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**Regulation of Gene Expression in the Basal
Ganglia**

By Carol. S. Simpson

Submitted for the Degree of Doctor of Philosophy

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SUMMARY

Using in situ hybridisation histochemistry we have demonstrated that :

1. Enkephalin gene expression is modulated by the dopaminergic system in the rat striatum. Intraperitoneal administration of the typical neuroleptic drugs haloperidol (1mg/kg) and fluphenazine (3mg/kg) significantly increase levels of proenkephalin mRNA in the caudal striatum compared to saline treated controls, although the atypical neuroleptic drug clozapine (3mg/kg) has no effect on the levels of proenkephalin mRNA in the rat. Methiothepin (10mg/kg), an antipsychotic with the ability to antagonise both serotonin and dopamine receptors can also induce proenkephalin mRNA levels in rat striatum. This induction is observed twenty-four hours after the drugs are given.

2. Tachykinin gene expression is modulated by the dopaminergic system in the rat striatum. Intraperitoneal administration of the typical neuroleptic drugs haloperidol (1mg/kg) and fluphenazine (3mg/kg) significantly reduce the levels of preprotachykinin mRNA in the rat striatum, compared to saline treated controls. This effect is observed twenty-four hours after drug administration. The atypical neuroleptic drug clozapine (3mg/kg) and the antipsychotic methiothepin (10mg/kg) have no significant effect on the levels of preprotachykinin mRNA in the rat striatum.

3. Somatostatin gene expression in the rat striatum is significantly increased by low doses of the atypical neuroleptic drug clozapine (3mg/kg) and by the antipsychotic drug methiothepin (10mg/kg), after acute administration. Acute treatment with typical neuroleptics have no effect in the rat striatum. Similarly, acute treatment

with clozapine (3mg/kg) and methiothepin (10mg/kg) significantly increase the levels of preprosomatostatin mRNA in the rat nucleus accumbens. Chronic treatment with clozapine (6mg/kg) also significantly increases preprosomatostatin mRNA levels in the rat striatum and in the olfactory tubercle.

4. Neuropeptide Y gene expression in the rat striatum is not changed significantly following acute neuroleptic drug treatment.

5. Chronic treatment with the neuroleptics fluphenazine (6mg/kg) and clozapine (6mg/kg) significantly increases the levels of D1 dopamine receptor mRNA in the rat rostral striatum. Chronic treatment with the above has no effect on the levels of 5-HT₆ receptor mRNA in the rat striatum.

6. Typical neuroleptics including haloperidol (1mg/kg) and fluphenazine (3mg/kg) can significantly induce the immediate early genes *c-fos*, *zif/268*, and *jun-B* in the rat striatum after 30-45 minutes. Haloperidol (1mg/kg), but not fluphenazine (3mg/kg), significantly induces *jun-D* mRNA in the rat striatum. Haloperidol (1mg/kg) can significantly increase *c-fos* and *jun-B* mRNA expression in the rat nucleus accumbens, while fluphenazine (3mg/kg) can significantly induce *zif/268* and *jun-B* mRNA expression in the rat nucleus accumbens. The atypical neuroleptic, clozapine (3mg/kg) had no significant effects on the expression of *c-fos*, *zif/268*, *jun-B*, and *jun-D* in either the rat striatum or accumbens regions. However, the immediate early, gene brain factor-1 (BF-1), was significantly induced in the rat caudal striatum twenty-four hours after clozapine (3mg/kg) treatment. *RHS2* gene expression was not affected by clozapine (3mg/kg) after twenty-four hours.

7. Striatal neurones in culture have a similar phenotype to striatal neurones in vivo. Cultured striatal neurones express the genes that encode enkephalins, substance P, somatostatin, and neuropeptide Y proteins similar to the in vivo situation, but not dynorphins. A small percentage of striatal neurones in culture are NADPH-positive and acetylcholinesterase-positive similar to the in vivo situation.

8. Dopamine ($5\mu\text{M}$) and SKF38393 ($1\mu\text{M}$), a D1/D5 receptor agonist, can significantly induce c-fos and zif/268 mRNA in cultured striatal neurones, compared to vehicle treated cultures. Maximal induction occurs after 45 minutes. This induction is blocked by SCH23390 ($1\mu\text{M}$), but not by eticlopride ($1\mu\text{M}$). The SKF38393-induced expression of both c-fos and zif/268 mRNA can be significantly reduced by pretreatment with the non-selective kinase inhibitor H7 ($50\mu\text{M}$) and with the selective protein kinase A inhibitor, KT5720 ($2\mu\text{M}$). The phorbol ester, 12-myristate-13-acetate (10^{-7}M) can significantly induce both c-fos and zif/268 mRNA after 45 minutes. The protein kinase C inhibitor, calphostin C ($1\mu\text{M}$), can significantly block the phorbol ester- induced expression of both c-fos mRNA and zif/268 mRNA. Calphostin C ($1\mu\text{M}$) can also significantly reduce the SKF38393- induced expression of c-fos mRNA, but is unable to block the SKF38393- induced expression of zif/268 mRNA. Quinpirole ($1\mu\text{M}$), a D2-like receptor agonist, can significantly induce c-fos mRNA in striatal cultures compared to control. The SKF38393- induced expression of both c-fos and zif/268 mRNA can be significantly reduced by coadministration of SKF38393 with quinpirole ($1\mu\text{M}$).

9. SKF38393 (1nM - $5\mu\text{M}$) can induce c-fos mRNA and zif/268 mRNA in a dose-dependent manner in striatal neurone cultures. Pretreatment with antisense oligonucleotides directed towards the D1 dopamine receptor and the D5 dopamine

receptor at a concentration of $1\mu\text{M}$ can significantly shift the dose response curves representing *c-fos* induction and *zif/268* mRNA induction by SKF38393, in a downward direction. However, pretreatment with the D1 sense oligonucleotide ($1\mu\text{M}$) has no effect on the dose response curve representing *c-fos* induction by SKF38393 (1nM - $5\mu\text{M}$). Although the D1 sense oligonucleotide caused a significant downward shift of the curve representing *zif/268* induction by SKF38393 (1nM - $5\mu\text{M}$).

10. Serotonin (5-HT) (1nM - $5\mu\text{M}$) can significantly induce both *zif/268* and *c-fos* mRNA in a dose-dependent manner in striatal neurone cultures. Pretreatment with ondansetron (200nM), a 5-HT₃ receptor antagonist, can block the 5-HT-induced expression of *zif/268* mRNA, and to a lesser extent *c-fos* mRNA. Pretreatment with the 5-HT_{1/2} receptor antagonists cyproheptadine (500nM), methiothepin (200nM), mesulergine (30 - 100nM) and spiperone (30 - 100nM) can block the 5-HT-induced expression of both *zif/268* and *c-fos* mRNA. The 5-HT_{2C} receptor agonist m-CPP (1nM - $5\mu\text{M}$) had no effect on the expression of *zif/268* or *c-fos* mRNA. Similarly 8-OH-DPAT (1nM - $5\mu\text{M}$), the 5-HT_{1A} receptor agonist, had no significant effects on either *zif/268* or *c-fos* expression in striatal cultures. Administration of the 5-HT_{2A} receptor agonist \pm DOI (1nM - $5\mu\text{M}$) had no significant effect on the expression of *zif/268* and *c-fos* mRNA in cultured striatal neurons.

Abbreviations

AchE	Acetylcholinesterase
GABA	γ -amino butyric acid
NPY	Neuropeptide Y
-ir	Immunoreactive
met-Enk	methionine enkephalin
cAMP	cyclic adenosine monophosphate
MPTP	1-methyl-4-phenyl-1,2,3,6,-tetrahydropyridine
NMDA	N-methyl-D-aspartate
PKA	protein kinase A
PKC	protein kinase C
DAG	Diacylglycerol
IEG	Immediate early gene
IP3	Inositol triphosphate
CREB	cAMP-response element binding protein
mRNA	messenger ribonucleic acid
EMEM	Earle's modified eagles medium
HBSS	Hank's balanced salts solution
NSE	neurone specific enolase
GFAP	glial fibrillary acidic protein
NGS	normal goat serum
PBS	phosphate buffered saline

Chapter 1

INTRODUCTION

1.1 Anatomical Organisation of the Basal Ganglia

The basic circuitry of the basal ganglia is classically known as being composed of multiple intrinsic loops through which most of the key structures of the system are reciprocally linked. As early as the 1970's it was proposed that the structures within the basal ganglia functioned as integrative centres, channelling information from different origins. The basal ganglia form a forebrain system that affects signals from a large part of the neocortex and focusses the output of this integrated information to target regions of the frontal lobe and brainstem involved in aspects of motor planning and motor memory.

The basal ganglia comprise a number of structures including the striatum, which is the largest structure in the basal ganglia, the substantia nigra, the globus pallidus, the subthalamic nucleus and the entopeduncular nucleus. The striatum, which itself comprises caudate nucleus, putamen and ventral striatum, receives an excitatory innervation from the substantia nigra pars compacta and dorsal raphe nucleus that is primarily dopaminergic and serotonergic respectively (Graybiel and Ragsdale., 1978), and also an excitatory glutamatergic innervation from specific cortical areas (Spencer., 1976; Divac *et al.*, 1977). In contrast, the striatum projects massively only to the globus pallidus, the substantia nigra pars reticulata and to the entopeduncular nucleus.

Immunocytochemical studies have revealed an inner complexity to the structure of the striatum. Biochemically specialised compartments were first detected in sections stained for acetylcholinesterase (AChE) activity (Graybiel and Ragsdale., 1978). Small AChE-poor regions referred to as striosomes or "patches" were embedded in an AChE-rich tissue of the striatum, which was

later referred to as the extrastriosomal matrix. In time, it was found that there were differences in the connections of striosomes and matrix compartments to the internal circuit within the basal ganglia, thus making functional differences between the striatal compartments likely. The matrix compartment receives the striatal inputs most directly related to sensorimotor processing. By contrast, striosomal compartments tend to receive inputs from neural structures, affiliated with the limbic system (Donoghue and Herkenham, 1986). Similarly the efferent projections from different striatal compartments show functional organisation. This mosaic pattern is presumably important for efficient and selective information processing by allowing a specific chemical environment to exist.

It has previously been suggested that the rat basal ganglia have a similar structural make-up to the human basal ganglia, and the organisation of connections within the rat basal ganglia have been shown to be remarkably similar to those in the human basal ganglia. Therefore the rat has been used in recent years as an adequate model for experimental research on basal ganglia function (Parent *et al.*, 1991).

1.2 Neurotransmitters and Neuropeptides in the Basal Ganglia

The basal ganglia contain a remarkable diversity of neuroactive substances, organised into functional subsystems that have unique developmental histories and vulnerabilities in neurodegenerative diseases.

Approximately 95% of striatal neurons *in vivo* are thought to be GABA-ergic projection neurons. Coexistence of GABA with neuropeptides in

various combinations in the principle circuits and interneurons of the basal ganglia, allows different combinations of cortical inputs, when projected into the basal ganglia, to be channeled not only into different output pathways, but also into different chemical environments.

In vivo, GABA-ergic projection neurons have been demonstrated to contain either enkephalin peptides or dynorphin and tachykinin peptides (Bolam *et al.*, 1985; Graybiel, 1990; Besson *et al.*, 1990). The remainder comprise possibly three populations of interneurons releasing acetylcholine, or GABA and neuropeptide Y (NPY), or somatostatin, NPY and nitric oxide (Bolam *et al.*, 1984; Phelps *et al.*, 1985; Smith and Parent, 1986; Graybiel, 1990; Vuillet *et al.*, 1990). Different neuronal populations have been demonstrated to project to different areas of the basal ganglia. It has been shown previously, with the use of retrograde tracing techniques, that the majority of neurons expressing enkephalins project to the globus pallidus and few to the substantia nigra. In contrast, neurons expressing dynorphin and substance P primarily project to the substantia nigra, with only small projections to the globus pallidus (Graybiel, 1986; Besson *et al.*, 1990). It has also been reported that there are differences in the abundance of mRNAs encoding neuropeptides contained in the same population of cells. For example, protachykinin mRNA appears to be considerably more abundant than prodynorphin mRNA (Young *et al.*, 1986), although both are located in striatoentopeduncular and striatonigral projection neurons. Similarly, somatostatin mRNA appears to be much more abundant than neuropeptide Y mRNA (Morris, 1989), although they are coexisting in the same cells. Protachykinin mRNA and prodynorphin mRNA show some indication of being preferentially located in the "patch" compartment of the striatum (Gerfen and Young, 1988; Morris *et al.*, 1989).

In vitro, there is only limited evidence to suggest what classes of neurons exist in the striatum. However, extracts of primary striatal cultures have been shown to contain immunoreactive (ir) somatostatin and substance P

(Kessler, 1986) and proenkephalin mRNA (Schwartz and Simantov, 1988; Vilijn et al., 1988), while analysis at the cellular level has shown the presence of GABA (Messer, 1981; Weiss et al., 1986; Surmeier et al., 1988), along with ir-leu-enkephalin (Surmeier et al., 1988) which could arise from the expression of either the prodynorphin gene or the proenkephalin gene and ir-somatostatin (Williams et al., 1991). Of these, only GABA, ir-leu-enkephalin and ir-somatostatin have been localised to neuronal cells rather than glia (Surmeier et al., 1988; Williams et al., 1991). Therefore, it remains unclear to what extent the other neuroactive substances detected in culture extracts are derived from non-neuronal cells.

1.3 Characterisation of Dopamine receptors

1.3.1 Molecular Biology

The dopaminergic system relies on the interaction of dopamine with several receptors. Dopamine can either inhibit or excite neurons (Bloom et al., 1965; McLennan and York., 1976). In 1979, Kebabian and Calne characterised two classes of dopamine receptor namely the D1 and D2 receptors. The concept of two distinct types of dopamine receptors mediating these two electrical effects (Cools and Van Rossum., 1976) was strengthened when it was found that the inhibition or excitation by dopamine could be selectively blocked (Struyker Boudier et al., 1974; Heiss and Hoyer., 1974). These two receptors were found to exert their biological actions by coupling to, and activating

different G-protein complexes. The D1 dopamine receptor interacts with G_s complexes resulting in the activation of adenylyl cyclase and an increase in intracellular cyclic AMP levels (Kebabian and Greengard., 1971). The D2 dopamine receptor interacts with G_i complexes to inhibit cyclic AMP production and thus decrease intracellular ^{CAMP} levels (De Camilli ^{et al.,} 1979).

Over the past decade the introduction of new molecular biological techniques have enabled the dopaminergic receptors to be cloned. The D2 receptor was originally cloned using a low stringency screening approach of rat complementary DNA ^{libraries} (Bunzow et al., 1988). However the polymerase chain reaction approach was used originally to clone the D1 receptor from rat striatum (Zhou et al., 1990; Sunahara et al., 1990) and from mouse neuroblastoma cells (Monsma et al., 1990). The low stringency approach to cloning the D1 receptor also proved to be successful (Deary et al., 1990).

Molecular cloning of G-protein linked receptors advanced rapidly with three further members of the dopamine receptor family being cloned, namely the D3, the D4 and the D5 dopamine receptors. The D3 receptor was originally identified using a DNA fragment of the D2 receptor as a probe under low stringency hybridisation conditions (Sokoloff et al., 1990; Giros., 1990). When expressed in eukaryotic cells this receptor was shown to bind D2, but not D1 ligands. By analysing mRNAs of a human cell line with D2 receptor complementary DNA (cDNA) probes, another D2-related mRNA was detected. The corresponding cDNA and gene analysis lead to the characterisation of the D4 receptor (Van Tol et al., 1991). The D4 receptor when expressed in COS-7 cells, binds D2 antagonists with a pharmacological profile distinct from, but reminiscent of that of the D2 receptor. The G-protein coupled D4 receptor has also been demonstrated to be coupled to adenylyl cyclase in the mouse retina (Cohen et al., 1992).

Finally the D1 dopamine receptor clone was used as a hybridisation probe to identify D1-related genes. A human D5 and a rat D1B receptor were

subsequently characterised (Zhou *et al.*, 1990; Sunahara *et al.*, 1991; Grandy *et al.*, 1991; Tiberi *et al.*, 1991; Weinshank *et al.*, 1991). The genes encoding for D5 and D1B receptors are intronless, unlike the genes encoding for the D2, D3 and D4 receptors, which are known to contain introns. The D5 and D1B display the same pharmacological profile, similar to that of the D1 receptor and are able to stimulate adenylyl cyclase activity.

1.3.2 Dopamine Receptor Ontogenesis

The recent development of molecular biology techniques such as *in situ* hybridisation (Le Moine *et al.*, 1991) and Northern blot analysis (Jaber *et al.*, 1992) have been used to study the localisation and regulation of dopamine receptor gene expression in rat and human brain. Using *in situ* hybridisation, it has recently been shown that the mRNA for the two isoforms of the dopamine D2 receptor, are detectable from gestational day 14 (G14) in the rat striatal primordium. At day 16, D2 receptor mRNA was present in the lateral part of the striatum and in the germinal ventricular zone lining the lateral ventricle. At day 18, D2 receptor mRNA was found in neurons of the caudate-putamen, the nucleus accumbens, the olfactory tubercle and the subependymal zone lining the lateral ventricle. The labelled cells were shown to have a neuroblastic and immature aspect before birth. After birth the topography and aspect of labelled cells was similar to the one observed in the adult animals (Guennoun and Bloch, 1991). A more sensitive method of detection known as reverse transcriptase polymerase chain reaction has recently been used to show D2 mRNA prenatally (Mack *et al.*, 1991). This method has been used to detect D1,

D1B and the D3 mRNA. The D1 receptor mRNA was detectable at gestational day 11 (G11) in the rat forebrain. The hybridisation signal remained relatively faint until G18 when it took on a more intense signal (Cadoret *et al.*, 1993). By gestational day 19, the D1 receptor mRNA was detectable by in situ hybridisation methodology (Guennoun and Bloch, 1992). The presence of D3 receptor mRNA was clearly detectable from G14 to G20 (with a maximum at G15), even though weak signals were observed as early as G11 (Cadoret *et al.*, 1993). The rat D1B receptor mRNA was detectable from gestational day 12, but the signal remained weak until G18-20 (Cadoret *et al.*, 1993). At present, the ontogenesis of the D4 receptor is not fully characterised.

The presence of dopaminergic markers has been reported at very early stages of brain development. Indeed, tyrosine hydroxylase and dopamine have been detected in the rat striatum at G14 (Foster *et al.*, 1987) before the full maturation of the dopaminergic system. It has also been suggested that the development of dopaminergic receptors is dependent on the development of the presynaptic nerve terminals containing the transmitters (Deskin *et al.*, 1981). Increasing evidence has shown that the dopamine receptors can be functional at very early stages of development. In avian retina embryonic cells in vitro, it has been demonstrated that stimulation of early dopamine receptors greatly inhibits the motility of the neuronal growth cone via a functional D1-type system (Lankford *et al.*, 1988). The D2 receptor has previously been shown to be modulated by guanylnucleotides as early as gestational day 15 (G15), suggesting that these receptors were already functionally coupled to a regulatory G-protein (Sales *et al.*, 1989). Furthermore, presynaptic D2 receptors appear to be functional as soon as dopamine release from striatal slices can be detected (De Vries *et al.*, 1992). Therefore the evidence to date suggests that the chemical environment (i.e. neurotransmitters) acts as a regulator of neuronal growth during development (Lankford *et al.*, 1988; Mattson, 1988).

1.3.3 Tissue Distribution of Dopamine Receptors

The distribution of the different dopamine receptor subtypes have revealed some important differences that may underlie functional differences. Since tissue distribution of the different dopamine receptors overlap in the central nervous system some selectivity may be obtained quantitatively rather than qualitatively. It has been shown using blotting analysis that the D3, D4 and D5 mRNAs are one to two orders of magnitude lower in abundance than D1 or D2 mRNAs (Sokoloff *et al.*, 1990; Van Tol *et al.*, 1991; Tiberi *et al.*, 1991). The anatomical distribution of both D1 and D2 subtypes is such that mRNA for both subtypes exists in high levels in the caudate-putamen, nucleus accumbens and the olfactory tubercle (Meador-Woodruff *et al.*, 1991). However, high levels of D2 mRNA, but not D1 mRNA are present in the substantia nigra and ventral tegmental area, whereas conversely the amygdala contains predominantly D1 mRNA. The distribution of D5 receptor mRNA was first reported to overlap the D1 mRNA (Sunahara *et al.*, 1991), but recent studies have subsequently conflicted with this view. It has since been found that D5 mRNA is present only in the hippocampus, hypothalamus and the parafascicular nucleus of the thalamus (Tiberi *et al.*, 1991; Meador-Woodruff *et al.*, 1992).

The newly cloned D2-like receptors, the D3 and the D4 subtypes have also been shown to have some unique tissue distribution. The distribution of D3 receptor gene transcripts in rat brain has been shown to be markedly different from that of the D2 receptor gene transcripts. A weak hybridisation signal has been observed in restricted parts of the striatum, although D3 receptor mRNA was found to be highly expressed in the olfactory tubercle-island of Calleja complex, the bed nucleus of stria terminalis and in the nucleus accumbens (Sokoloff *et al.*, 1990; Levesque *et al.*, 1992; Sokoloff *et al.*, 1992).

The expression of the D4 mRNA was shown to be relatively high in the monkey frontal cortex, midbrain area, amygdala and medulla, but lower levels were detected in the basal ganglia (Van Tol et al., 1991). In the rat, significant levels of expression were found in the hypothalamus, thalamus, olfactory bulb and frontal cortex. However, 20-fold higher levels of D4 mRNA were observed in the cardiovascular system (O' Malley et al., 1991). The D4 dopamine receptor transcripts have also been observed in the mouse retina (Cohen et al., 1992).

1.3.4 Cellular Localisation of Dopamine Receptors

The cellular localisation of dopamine receptors is currently a matter of much controversy. At present there are two schools of thought concerning this issue as some research groups support segregation of D1 and D2 dopamine receptors in the striatum (Gerfen et al., 1990; Le Moine et al., 1990; Gerfen, 1992; Le Moine et al., 1992), whilst other groups hold to the idea that D1 and D2 dopamine receptors occur on the same cell (Seeman et al., 1989; Bertorello et al., 1990; Piomelli et al., 1991; Surmeier et al., 1993).

Research supporting segregation of D1 and D2 dopamine receptors in the striatum is based on studies performed in intact rats, and thus the segregation is thought not to be a consequence of denervation of dopaminergic nerves. Stimulation of D1 and D2 receptors has been shown to preferentially affect the function of striatonigral and striatopallidal neurons, respectively, in the intact striatum. Treatment with D2 antagonists rapidly induces expression

of the immediate early gene *c-fos* specifically in striatopallidal neurons (Dragunow et al., 1990; Robertson et al., 1992). Conversely, indirect dopamine receptor agonists result in D1-mediated changes in the mRNA and protein levels of immediate early genes (Cole et al., 1992; Beretta et al., 1992; Cenci et al., 1992; Steiner et al., 1993) predominantly in striatonigral neurons. The effects of dopamine depletion on striatal neurons are also thought to be consistent with differential localisation of D1 and D2 receptors on striatonigral and striatopallidal neurons. Enkephalin mRNA and protein, which are predominantly localised to striatopallidal neurons (Gerfen and Young, 1988), are elevated by this depletion (Young et al., 1986; Gerfen et al., 1990; Li et al., 1990; Gerfen et al., 1991; Engber et al., 1992). Conversely in striatonigral neurons, mRNA and peptide levels of substance P and dynorphin are decreased (Young et al., 1986; Gerfen et al., 1990; Li et al., 1990; Gerfen et al., 1991). The levels of D1 and D2 receptor mRNA, together with receptor binding, are decreased and increased respectively (LaHoste et al., 1991., Joyce, 1991; Blunt et al., 1992). The lesion-induced alterations of peptide and D1 receptor mRNAs in striatonigral neurons are reversed by treatment with D1-selective agonists, whereas the alterations in peptide and D2 receptor mRNAs in striatopallidal neurons are reversed by D2 receptor agonists (Gerfen et al., 1990; Engber et al., 1992). In situ hybridisation studies have also demonstrated that D1 and D2 receptors are expressed preferentially in one of two populations of striatal efferent neurons, substance P-containing and enkephalin-containing neurons respectively, with little or no overlap of expression (Le Moine et al., 1990; Gerfen et al., 1990; Le Moine et al., 1991).

Alternatively the hypothesis for colocalisation of dopamine receptors is also strongly supported. On the basis of sensitive RNA detection techniques, current recordings obtained from single cells in vitro (Surmeier et al., 1992) and immunohistochemical data (Ariano et al., 1992), Surmeier and colleagues have suggested that D1, D2 and D3 receptors are largely colocalised in most

striatal efferent neurons, especially striatonigral neurons. Using immunohistochemical techniques, the D2 receptor staining pattern in both striatonigral and striatopallidal neurons has been observed by several research groups (Ariano et al., 1993; Boundy et al., 1993). Other immunocytochemical work relevant to the issue includes localisation of DARPP-32 which is strongly correlated with D1 receptor binding in both striatonigral and striatopallidal neurons. Recently, other signalling elements in the D1 receptor pathway have been found in both striatonigral and striatopallidal neurons. It has been shown that most medium spiny neurons are immunoreactive for a particular G-protein α -subunit that mediates D1 receptor activation of adenylate cyclase in the neostriatum (Herve et al., 1993). Some of the in vitro research supporting colocalisation has been argued against, because of the possible alterations in cellular environment affecting mRNA levels. However there has been much research performed in vivo supporting the convergence theory, including peptide expression assays (Haverstick et al., 1989; Morris and Hunt., 1991; Angulo, 1992) and electrophysiological studies (Hu et al., 1990). Ultimately, further research and more sensitive techniques are necessary in order to answer the question of dopamine receptor localisation.

1.4 Drug Pharmacology of Dopamine Receptors

1.4.1 D1-Like Receptors

The sensitivities of the cloned D1 and D5 receptors to antagonist drugs are very similar and are also generally the same as those D1 receptors in native tissue (Seeman and Van Tol, 1993). It has been previously shown, with the use of binding studies in COS-7 cells, that the compound SCH23390 is a very potent antagonist, producing K_i values of 0.35nM and 0.30nM at D1 and D5

receptors respectively. Fluphenazine was shown to have some antagonistic properties at the D1 and D5 receptors, although much less than SCH23390 (Sunahara *et al.*, 1991).

For dopamine receptor agonists, however, it has proved difficult to obtain accurate comparisons of the agonist sensitivities of the cloned receptors with those in native tissues. This is because dopamine receptors in native tissues can readily adopt either a high-affinity state or low-affinity state for an agonist, while cultured cells vary in their ability to reveal the high affinity state. COS-7 cells, for example, do not have sufficient or appropriate G-protein subunits to allow the high affinity state of a receptor to exist. Agonist drugs such as SKF38393, R-fenoldopam, and (-) apomorphine have a very high affinity for the D1-like receptors. However, an important consistent difference between the D1 and the D5 receptors is that dopamine itself is about ten times more potent at the D5 receptor (Sunahara *et al.*, 1991; Seeman and Van Tol, 1994).

1.4.2 D2-Like Receptors

Accurate values for many of the agonist potencies at D2, D3, and D4 receptors are not currently available. Nevertheless, some important selectivities are clearly emerging. For example, bromocriptine is about two orders of magnitude more potent at D2 and D3 receptors than at D4 receptors (Seeman and Van Tol, 1993). Moreover, 7-hydroxy-DPAT is approximately tenfold more potent at D3 receptors than at D2 receptors (Seeman and Van Tol, 1993). Quinpirole also has a relatively high affinity for all D2-like receptors (Van Tol *et al.*, 1991; Seeman and Van Tol, 1994).

In the case of antagonists acting at D2-like receptors, haloperidol and fluphenazine have relatively high affinity for all D2-like receptors. Clozapine has been shown to be one to ten orders of magnitude more potent at D4 receptors than at D2 or D3 receptors (Van Tol et al., 1991; Seeman and Van Tol, 1994). With the use of comparative binding studies, the density of D4 receptors in the human striatum has been shown to be approximately 10% of the total population of the D2-like receptors (Seeman et al., 1993). Methiothepin has also been shown to block dopaminergic receptors (Enjalbert et al., 1978; Lloyd and Bartholini, 1974). A small displacement of ³H-methiothepin was observed following treatment with high concentrations of haloperidol, a dopaminergic receptor antagonist, suggesting that part of bound ³H-methiothepin is also associated with dopaminergic receptors (Nelson et al., 1979).

1.5 Basal Ganglia Dysfunction- The Clinical Implications

1.5.1 Parkinson's Disease

Parkinson's disease was first characterised by James Parkinson in 1817 as a late onset disease manifested as tremor, rigidity, postural abnormalities and bradykinesia. By 1960, Ehringer and Hornykiewicz reported that these abnormalities were accompanied by a neurochemical change taking the form of a loss of dopamine in the corpus striatum. In the early eighties Parkinson's disease was referred to as a neurodegenerative disease characterised by the loss of dopaminergic neurons from the pigmented substantia nigra (Forno, 1982).

A considerable amount of research has been focussed on basal ganglia function. Animal studies, together with human postmortem analysis have provided a much greater understanding of the pathophysiological basis for Parkinson's disease. The basic observation of decreased striatal dopamine in patients with Parkinson's disease was mimicked in the laboratory with chemicals such as reserpine, the neurotoxins 6-hydroxydopamine, which causes a permanent lesion of dopaminergic projections to the striatum (Ungerstedt , 1971) resulting in severe neurological impairment in rats reminiscent of several symptoms of Parkinson's disease (Hornykiewicz and Kish, 1987), and 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) administered in primates which produces classic Parkinsonian-like symptoms thought to be due to the build up of a toxic metabolite (Langston, 1985). Further work revealed that dopamine depletion in rat brain via lesions increased dopamine turnover in the remaining terminals as a result of neuroplastic changes in the striatum (Zigmond et al., 1984). Moreover, recent evidence indicates that this neurochemical response is associated with the maintenance of the extracellular dopamine concentration at near normal levels despite the loss of dopamine terminals (Robinson and Wishaw, 1988; Zhang et al., 1988). Disruption of dopaminergic transmission with the use of dopamine antagonist drugs such as spiroperidol in rats previously lesioned, has been shown to result in significantly impaired behaviour patterns, but no effects were observed on intact rats given the same dose (Heffner et al., 1977). An increase in the spontaneous firing rates of striatal cells have also been associated with dopamine depletion, suggesting that the dopamine neurons exert an inhibitory influence on striatal activity (Orr et al., 1986).

A standard treatment for Parkinson's disease sufferers involves administration of levodopa, which is taken up by nerve terminals where it is decarboxylated to dopamine. The dopamine then activates receptors in the region of the depleted striatum and substantia nigra (Kandel and Schwartz,

1985; Gilman et al., 1985). Several studies have also demonstrated that dopaminergic D2 receptor agonists such as apomorphine can produce locomotion and stereotyped behaviour in lesioned animals when administered at dose levels which were shown to be ineffective in normal rats (Hollister et al., 1974; Kelly et al., 1975). This lead to the suggestion that dopamine receptors became "supersensitive" after lesioning (Hollister et al., 1975). The indirect dopamine receptor agonist amphetamine is also known to enhance locomotor activity at low doses, whilst producing repetitive stereotyped behaviour at high doses. These two aspects of animal behaviour have been attributed to separate neural systems, such that dopamine in the caudate is essential for stereotyped behaviour and dopamine in the nucleus accumbens septi is linked to an enhanced locomotor activity following amphetamine administration, after a 6-hydroxydopamine lesion (Kelly et al., 1975). Amphetamine has been shown to cause the release of dopamine from the caudate-putamen (Ungerstedt, 1971; Voigtlander and Moore, 1973).

Dopamine-containing neurons of the substantia nigra pars compacta are known to release dopamine from both the terminals in the striatum and from the dendrites in the substantia nigra pars reticulata (Cheramy et al., 1981). When dopamine neurons degenerate as in Parkinson's disease, or are destroyed by neurotoxins, the D2 dopamine receptors in the substantia nigra pars compacta disappear because the D2 receptors are almost entirely confined to dopamine neurons (Quik et al., 1979; Murrin et al., 1979). However, D1 receptor density in the substantia nigra pars reticulata, which is as high as the D1 receptor density in the striatum (Boysen et al., 1986), is unaffected by such dopamine neurotoxin lesions because the D1 receptors are presynaptic and are located on the terminals of descending striatonigral fibres (Savasta et al., 1986). Supersensitivity of dopamine D1 receptors has been demonstrated in the absence of dopamine stimulation (Savasta et al., 1986). Electrophysiological studies have also suggested that D1 dopamine receptor stimulation decreases

cell firing rates in the substantia nigra pars reticulata, presumably by increasing release of the inhibitory transmitter GABA from the terminals of the striatonigral fibres (Weick and Walters, 1987). This has also been demonstrated in nigral slice studies *in vitro* (Aceves *et al.*, 1991). It has since been suggested that a D1-D2 synergy exists, in which dopamine plays a central role as both a regulator and effector (Robertson, 1992).

A reduction in peptide levels in Parkinsonian basal ganglia has also been reported (Agid and Javoy-Agid, 1985). A decrease in methionine-enkephalin (met-Enk) and leucine-enkephalin has been observed in the putamen and in both segments of the globus pallidus (Taquet *et al.*, 1983, 1985) indicating that these neuronal systems are altered in the disease. The level of the neuropeptide substance P has also been shown to decrease in the lateral globus pallidus in Parkinson's disease patients (Mauborgne *et al.*, 1983).

The interaction between dopamine and other neurotransmitters in the basal ganglia, in particular the dopamine-glutamate interaction, has been implicated in neurotoxicity at the nigrostriatal dopamine terminals. Stimulation of excitatory amino acid receptors of the N-methyl-D-aspartate (NMDA) subtype has been shown to enhance dopamine release (Snell and Johnson, 1986; Clow and Jharnandas, 1989) and synthesis (Arias-Montano *et al.*, 1992) within the striatum. In the dopamine-depleted striatum, an elevation of NMDA receptor binding was shown and was found to be greatest in the dorsal caudate and putamen, regions that were found to have extensive loss of dopamine transporter sites (Weihmuller *et al.*, 1992). Similar findings of a relationship between reduced striatal dopamine content and increased binding to NMDA receptors have been reported in the 6-hydroxydopamine treated rat model of Parkinson's disease (Samuel *et al.*, 1990).

It has therefore been proposed that a decreased dopamine activity in the striatum of Parkinson's disease patients will, through direct and indirect basal ganglia pathways, release the substantia nigra pars reticulata and

entopeduncular neurons from their inhibitory controls, due to a lack of dopaminergic excitation from the substantia nigra pars compacta. Consequently thalamic targets of the substantia nigra pars reticulata and entopeduncular nucleus will be inhibited, leading to a diminished thalamocortical activation. This reduced cortical activation is thought to result in a decreased activity of striatocortical neurons and thus reduced striatal glutamate release. Thus the overall effect of dopamine loss in the striatum is inhibition of cortically initiated movement, to cause akinesia, hypokinesia, and bradykinesia. As a consequence, upregulation of NMDA receptors within striatal target neurons may occur as a compensatory mechanism (Weihmuller *et al.*, 1992). In coordination with this result, chronic NMDA receptor antagonism can lead to increased receptor sensitivity (Norman *et al.*, 1990).

1.5.2 Schizophrenia

The discovery of neuroleptics in 1952 (Delay *et al.*, 1952) initiated research seeking a possible biological basis for schizophrenia. Various hypotheses have been proposed and are currently under investigation.

The dopamine hypothesis of schizophrenia proposes that dopamine pathways are overactive in the disease state (Van Rossum, 1967; Matthysse, 1973). Much of the evidence for this hypothesis is based on the fact that neuroleptic drugs block dopaminergic D2 receptors (Seeman *et al.*, 1975; Creese *et al.*, 1976). This action of neuroleptics is thought to account for neuroleptic-induced Parkinsonism and hyperprolactinaemia. Additional evidence for a dopaminergic basis for schizophrenia comes from the exacerbating effects of dopamine-mimetic drugs (Seeman, 1987; Lieberman, 1987) and the elevation of dopamine D2 receptors in the post-mortem brains taken from schizophrenic patients (Cross *et al.*, 1981; Joyce *et al.*, 1988;

Reynolds *et al.*, 1988; Seeman and Niznik, 1990). The density of D1 dopamine receptors remains normal in schizophrenia (Seeman *et al.*, 1987; Seeman and Niznik, 1990).

A different approach to understanding the pathophysiology of schizophrenia is provided by use of phencyclidine, which can be used to induce a syndrome of both negative and positive schizophreniform symptoms. The behavioural effects of phencyclidine are due at least in part, to its ability to block glutamate function via the NMDA receptor (Javitt and Zukin, 1991). Reported alterations in glutamatergic synaptic markers, particularly of frontal cortex in schizophrenic patients (Nishikawa *et al.*, 1983; Deakin *et al.*, 1989) may provide neurochemical evidence for a dysfunction of this transmitter system. Hence it has been suggested that drugs effective in alleviating the symptoms of schizophrenia that do not respond to classical antipsychotics may do so by interaction with glutamate systems. Recent evidence has emerged showing human cortical pyramidal neurons to be enriched with 5-HT_{1A} receptors (Bowen *et al.*, 1992). This provides a potential receptor mechanism for the modulation of cortical glutamatergic neurons, raising the possibility that an interaction with 5-HT_{1A} receptors may contribute to the restoration of normal glutamatergic function after neuroleptic drug administration.

1.6 Second Messenger Pathways induced by Dopamine Receptor Stimulation

Cori and co-workers made the initial discovery that protein phosphorylation was essential for all forms of signal transduction (Cori and Cori., 1945). In the mid 1950's Sutherland and his associates contributed to this finding by discovering that cyclic AMP was the "second messenger" involved in the transmission of effects of noradrenaline in the liver, and later demonstrated

that cyclic AMP caused the transfer of phosphate groups between key proteins (Robison and Sutherland., 1971).

The ability of dopamine receptors to induce different second messenger pathways has thus far been studied in D1 and D2 receptors (Vallar *et al.*, 1990). The D1 and D2 dopamine receptors induce two types of signal transduction pathway, one obligatory and several cell specific. In D1 or D2 receptors the obligatory pathway is detected in every cellular environment and represents stimulation or inhibition of adenylyl cyclase, respectively. However, dopamine can also induce additional and sometimes different signal transduction pathways depending on the cell type.

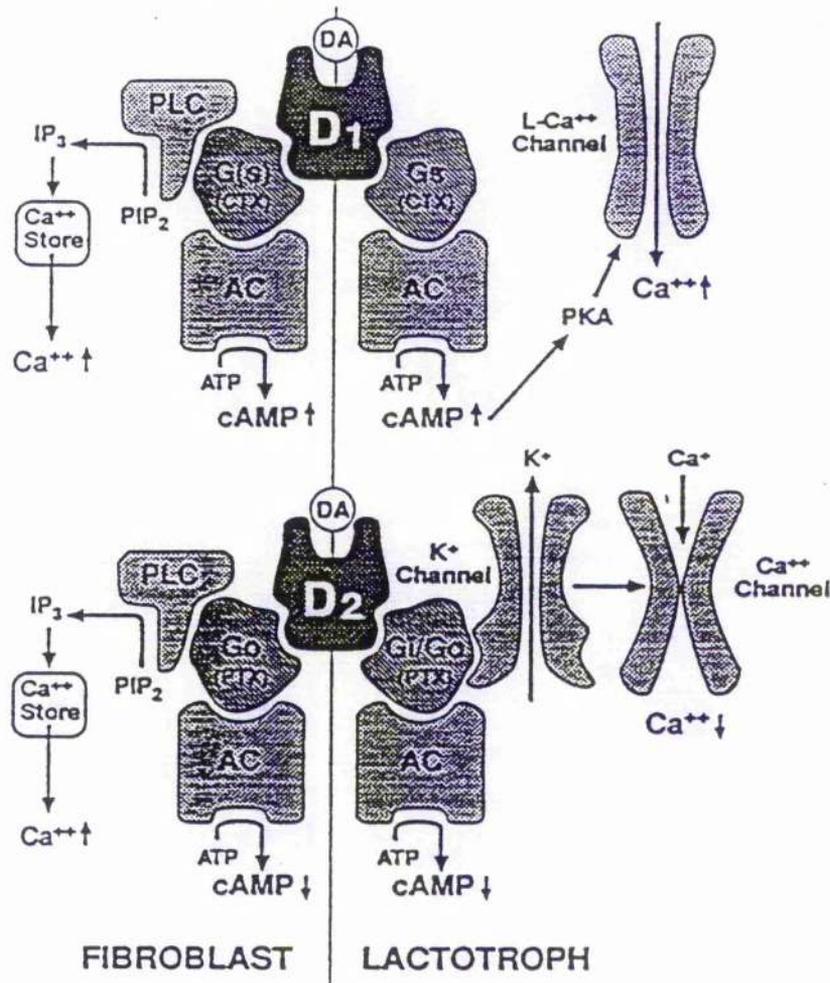


Figure 1 : Signalling pathways of the D1 and D2 dopamine receptors in the mouse fibroblast Ltk- and the rat somatomammotroph GH4Cl cells. DA represents dopamine, CTX and PTX means cholera toxin- and pertussis toxin-sensitive, respectively. G = G protein; PLC = phospholipase C; AC = adenylyl cyclase; PIP2 = phosphoinositol bisphosphate; IP3 = inositol triphosphate. The direction of arrows indicates whether the second messenger increases or decreases.

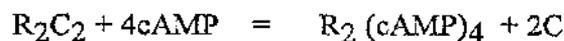
In every cell studied, stimulation of the D1 dopamine receptor increases cyclic AMP production. In addition, in GH4Cl cells, the D1 receptor potentiates activation of L-type voltage-dependent calcium channels in a cyclic dependent manner (Civelli et al., 1993). The D1 receptor has also been associated with an increased hydrolysis of phospholipids and calcium mobilization in xenopus oocytes (Mahan et al., 1990). The D2 receptor, while inhibiting adenylyl cyclase activity, does not affect phosphoinositide hydrolysis in GH4Cl cells and causes a decrease in intracellular free calcium mediated by a hyperpolarizing effect, mostly due to activation of potassium channels. ^(figure 1) However, in Ltk-cells, _A activation of the D2 receptor leads to an increase in intracellular free calcium partly due to release of calcium ions from intracellular stores following the rapid stimulation of phosphoinositide hydrolysis and, in part, to influx from extracellular medium (Gatti et al., 1991). In addition, in chinese hamster ovary (CHO) cells, D2 receptors have been shown to mediate the potentiation of arachidonic acid release by a mechanism that involves protein kinase C, and is independent of adenylyl cyclase inhibition (Kanterman et al., 1990).

Activation of kinase enzymes prior to protein phosphorylation and the resultant physiological effect has proved to be an important target for the understanding of receptor mediated events. It has become clear that a great many different protein kinases exist (Hunter, 1987). These kinases can be broadly divided into two large groups, the protein seryl or threonyl kinases and the protein tyrosyl kinases, based on their specificities towards the different amino acids in the protein (Krebs, 1985). A given kinase may phosphorylate a number of different proteins, however it usually recognises a short sequence of amino acids referred to as a 'consensus sequence'. This was first demonstrated by work on the cyclic AMP-dependent protein kinase (PKA) (Kemp et al., 1975; Humble et al., 1975; Kemp et al., 1976). Using synthetic peptides the consensus substrate sequence for PKA was defined (Kemp et al., 1977). It was concluded that PKA preferentially phosphorylated a

serine located at the second position after a pair of arginines as in the following sequence X-Arg-Arg-X-Ser-X (X= neutral amino acid).

1.7 Cyclic AMP-dependent Protein Kinase (PKA)

Cyclic AMP-dependent protein kinase (PKA) is an enzymatically inactive tetramer composed of two regulatory (R) and two catalytic (C) subunits. PKA undergoes dissociation when the enzyme is activated by cyclic AMP (Brostrom *et al.*, 1971; Gill and Garren, 1972; Kumon *et al.*, 1972; Tao *et al.*, 1970). The following equation for the activation reaction was composed after the quaternary structure of the enzyme had been determined (Beavo *et al.*, 1975; Rosen *et al.*, 1975) and the extent of cyclic AMP binding clarified (Corbin *et al.*, 1978).



The regulatory subunits exist in dimeric form which is retained after dissociation, whereas the catalytic subunits exist as monomers. Each R subunit binds two molecules of cyclic AMP.

In the brain PKA is broadly distributed in specific areas (Cadd and McC Knight., 1989). It plays an important role in neuronal regulation over different time frames. Protein kinase A acutely modulates the activity or function of neuron -specific enzymes (Zigmond *et al.*, 1989) and/or channel proteins (Li *et al.*, 1992; Sculptoreanu *et al.*, 1993) by catalysing the phosphorylation of these proteins. In such instances, evoked PKA activity produces rapid changes in neuronal function via direct phosphorylation of existing protein molecules. Alternatively, translocation of the active catalytic subunit of PKA into the nucleus and subsequent phosphorylation of nuclear proteins, including transcriptional activators, gives rise to slower long term changes in neuronal

function. In such instances PKA regulates de novo synthesis of mRNA leading to new protein synthesis.

In many cyclic AMP-responsive genes an octamer sequence motif known as the cyclic AMP (cAMP) response element (CRE) resides in the 5' upstream region and mediates cAMP responsiveness (Roesler *et al.*, 1989; Goodman *et al.*, 1990). PKA regulates the transcriptional activity of the CRE by catalysing the phosphorylation of transcription factors. One such transcription factor is the CRE-binding protein CREB (Hoeffler *et al.*, 1988; Gonzalez *et al.*, 1989). After phosphorylation by PKA CREB can activate transcription of CRE-containing genes (Gonzalez and Montminy., 1989; Lee *et al.*, 1990).

1.8 Protein Kinase C (PKC)

In the nervous system, activation of protein kinase C (PKC) has been related to enhancement of neurotransmitter release (Malenka *et al.*, 1986; Malenka *et al.*, 1987), control of growth differentiation (Burgess *et al.*, 1986; Spinelli and Ishii, 1983) and modification of neuronal plasticity (Routenberg, 1985).

PKC consists of a family of closely related proteins highly concentrated in the central nervous system. This protein kinase is present in a wide variety of tissues. Recent reports indicate that PKC's are heterogenously distributed in the brain and that neurons contain the highest levels of PKC (Huang, 1987; Brandt *et al.*, 1987). Molecular cloning has established the existence of several PKC isoforms, illustrating the diversity of this enzyme family. PKC's can be divided into two major groups, group A and group B, based on their sequence similarity at several coding regions. Group A PKC-catalysed phosphorylation

has been shown to be dependent on calcium and phosphatidylserine for its activation (Takai et al., 1979; Kaibuchi et al., 1981).

Protein kinase C can be activated by a naturally occurring product diacylglycerol (DAG), an intermediate in the biosynthesis and degradation of glycerolipids. DAG is thought to increase the affinity of the enzyme for calcium and phospholipid (Takai et al., 1979; Kishimoto et al., 1980). Tumour-promoting phorbol esters have also been shown to stimulate activity of PKC's in vitro (Blumberg, 1980; Chiang et al., 1981). It is currently the idea that DAG may be derived from the hydrolysis of a small pool of membrane phosphatidylinositol 4, 5-bisphosphate (PIP₂) by phosphatidylinositol-specific phospholipase C, resulting in a rise in inositol triphosphate (IP₃) and consequently intracellular calcium and diacylglycerol (Berridge, 1987). This signalling pathway has been linked to the actions of many neurotransmitters. Another potential source of DAG is derived through hydrolysis of phosphatidylcholine (Chan et al., 1989).

1.9 Immediate Early Genes

Immediate early genes (IEGs) were initially characterised in non-neuronal cells while studying growth factor responsive-genes. This research led to the discovery of a unique class of genes whose transcription can be activated within minutes. ^{independent of protein synthesis} The c-fos and c-myc protooncogenes were among the first IEGs to be identified (Kelly et al., 1983; Greenberg and Ziff, 1984). Many IEGs have since been identified in the brain and can be divided into distinct families based on their structural similarities. Among these families is the fos family whose members are closely related to c-fos (e.g. fos-B, Zerial et al., 1989) or the jun family, whose members are closely related to c-jun (e.g. jun-B, Ryder et al.,

1988) and jun-D (Ryder *et al.*, 1988) or the homeobox POU-domain containing proteins (e.g. RHS2, Le Moine and Scott Young, 1992) or the homeobox fork head gene, BF-1 (Mathis *et al.*, 1992). However, some IEGs are structurally unique, as for example zif/268, a zinc finger containing protein (Millbrandt, 1987). IEGs have a number of characteristics that makes them different from late response genes. Their expression tends to be low in quiescent cells, but is rapidly induced at the transcriptional level within minutes of extracellular stimulation. This transcriptional induction is transient and independent of new protein synthesis. Although, the subsequent shut-off of transcription requires new protein synthesis.

The IEG is currently thought to function as a transcription regulatory factor, that may act as a molecular switch coordinating changes in gene expression (Sheng and Greenberg, 1990). Direct evidence for a regulatory function came from the findings that many IEGs were found to encode transcription factors. This has been best established for c-fos and c-jun, both of which were initially identified as protooncogenes (Van Beveren *et al.*, 1983; Maki *et al.*, 1987). Recently it has been demonstrated that the products of these IEGs can interact with each other to form a heterodimeric transcription factor complex (Halzonetis *et al.*, 1988; Kouzarides and Ziff, 1988; Nakabeppu *et al.*, 1988; Rauscher *et al.*, 1988). This interaction was later shown to occur through a conserved dimerisation domain referred to as a 'leucine zipper' that is present in both fos and jun proteins (Landschultz *et al.*, 1988; Kouzarides and Ziff, 1988; Bos *et al.*, 1989; Gentz *et al.*, 1989; O'Shea *et al.*, 1989). The leucine zipper domain was shown to have an α - helical structure (Landschultz *et al.*, 1988). Adjacent to the leucine zipper, a highly basic stretch of amino acids were identified and shown to be required for the interaction of the fos/jun heterodimer with DNA (Kouzarides and Ziff, 1988; Gentz *et al.*, 1989; Neuberger *et al.*, 1989; Turner and Tjian, 1989). The heterodimer was demonstrated to bind with high affinity to the DNA consensus sequence -

TGACTCA- (reviewed by Curran and Franza, 1988). The sequence -TGACTCA- was initially identified as a phorbol ester-inducible promoter element and the binding site for a transcription factor activity termed AP-1 (Angel et al., 1987; Lee et al., 1987). Fos and jun have since been shown to be major components of the AP-1 site (Bohmann et al., 1987; Rauscher et al., 1988). Different combinations of fos/jun and jun/jun dimers have been demonstrated to bind to the AP-1 site in vitro (Nakabeppu et al., 1988; Cohen et al., 1989). There is also some evidence of functional specificity for some IEGs. It has been shown in vitro that c-fos/c-jun complexes activate transcription of genes containing AP-1 sites, while c-fos/jun-B complexes appear, under some circumstances, to repress transcription (Chiu et al., 1989; Schutte et al., 1989).

The mRNA encoding the immediate early genes Jun-D and c-jun is present constitutively in striatal neurons, and the corresponding proteins are likely to be available to form complexes with c-fos once it is synthesised. The mRNA encoding jun-B is not normally detectable in striatal neurons. Zif/268, the zinc finger containing immediate early gene protein, was shown to bind to a different DNA consensus sequence, -GCGGGGGGCG-. The DNA binding ^{domain encoded} by the homeobox genes, RHS2 and BF-1 is relatively large in comparison to other gene families studied. The POU-domain for RHS2 spans 145 amino acids (Le Moine and Scott Young, 1992) and for BF-1, the DNA binding sequence spans 56 amino acids (Tao and Lai, 1992).

1.10 Mechanisms of Induction of Immediate Early Genes

IEGs have been shown to be induced by a wide variety of physiological stimuli, including electrical excitation, neurotransmitters and growth factors (Greenberg et al., 1985, 1986; Morgan and Curran, 1986; Bartel et al., 1989). Studies have shown that many of the stimuli used to induce IEGs have a common feature in that they lead to or substitute for, calcium influx through voltage sensitive calcium channels (Greenberg et al., 1986; Morgan and Curran, 1986).

The molecular basis of IEG regulation by trans-synaptic stimulation has been studied primarily in PC12 cells. Transcriptional activation of *c-fos* as a result of calcium influx involves a specific site on the gene termed a calcium response element (CaRE) (Sheng et al., 1988). This site was later shown to be indistinguishable from the cAMP response element (CRE) and that cAMP and calcium signals converge on this element in vivo (Sheng et al., 1990). It has been demonstrated that both calcium and cAMP stimulation of PC12 cells result in the rapid phosphorylation of transcription factor CREB, the nuclear protein that binds specifically to the calcium/CRE (Gonzales and Montminy, 1989; Sheng et al., 1990). In response to cAMP, CREB is phosphorylated at a specific residue (Ser-133) known to be recognised by cyclic AMP dependent protein kinase (PKA) and to be important for activation of CREB as a transcription factor in vitro and in vivo (Yamamoto et al., 1988; Gonzales and Montminy, 1989). This phosphorylation is thought to activate CREB via an allosteric mechanism (Yamamoto et al., 1990), although it is still incompletely understood. CREB is phosphorylated at the identical Ser-133 position in response to calcium stimulation (Sheng et al., 1990). However evidence suggests that this phosphorylation is mediated by a calcium regulated protein kinase (Sheng et al.,

1990). The phosphorylated CREB then is thought to interact with the basic transcriptional machinery to stimulate gene transcription.

Genes containing calcium/CRE's include not only IEGs like c-fos and zif/268 (Christy *et al.*, 1988; Changelian *et al.*, 1989), but also a group of neuropeptide genes including somatostatin (Montminy *et al.*, 1986; Montminy and Bilezikjian, 1987), and proenkephalin (Comb *et al.*, 1986).

While calcium and cAMP signalling pathways are likely to mediate gene activation by many different neurotransmitters in the central nervous system, another intracellular signalling mechanism has been shown to be involved in the regulation of neuronal IEG expression. The activation of the IEG, c-fos, by the protein kinase C pathway has been demonstrated to be mediated by a distinct upstream regulatory element known as the serum response element (Gilman *et al.*, 1986; Prywes and Roeder, 1986; Treisman, 1986; Greenberg *et al.*, 1987; Sheng *et al.*, 1988; Gilman, 1988). The serum response element is a specific binding site for a nuclear protein termed the serum response factor (Gilman *et al.*, 1986; Prywes and Roeder, 1986; Treisman, 1986; Greenberg *et al.*, 1987). This interaction plays a crucial role in c-fos activation by a diverse set of stimuli, including phorbol esters (Sheng *et al.*, 1988).

1.11 Neuroleptics

Neuroleptics of various chemical classes have been used for more than 35 years for the treatment of psychiatric disorders (Leysen and Niemegeers, 1985). Neuroleptic drugs have been divided into two major classes, namely 'typical' neuroleptics, such as haloperidol and fluphenazine, and 'atypical' neuroleptics such as clozapine and fluperlapine. It is now well accepted that blockade of dopamine D2 receptors plays a major role in their antipsychotic

action (Leysen and Niemegeers, 1985; Meltzer *et al.*, 1989; Richelson, 1984; Seeman, 1980). However treatment with higher dosages of 'typical' neuroleptics has been shown to provoke adverse reactions such as extrapyramidal side effects following acute treatment and tardive dyskinesia after prolonged chronic treatment. Mesolimbic and mesocortical dopamine D2 receptors were suggested to play a role in the antipsychotic effects of the drugs whereas extensive occupancy of striatal D2 receptors was thought to be responsible for extrapyramidal side effects (Crow *et al.*, 1977; Stevens, 1973; Wilk *et al.*, 1975). The atypical neuroleptic, clozapine, has been shown to produce much less acute extrapyramidal side effects than typical neuroleptic drugs (Matz *et al.*, 1974). There have been no reliable reports that clozapine produces or exacerbates tardive dyskinesia, although it can block the symptoms of tardive dyskinesia (Meltzer and Luchins, 1984). In two relatively recent clinical trials, clozapine was shown to be a more effective antipsychotic drug than the typical neuroleptic chlorpromazine (Claghorn *et al.*, 1987; Kane *et al.*, 1988). Meltzer (1989) suggested that the clinical advantages of the use of clozapine may be due to a biological effect of the drug on neurotransmitter systems. They later demonstrated that atypical antipsychotic drugs could be distinguished from typical antipsychotic drugs on the basis of lower D2 and higher 5-HT₂ pK_i values.

More recently it has been suggested that potent serotonergic 5-HT₂ receptor antagonism could alleviate negative symptoms in schizophrenia. Moreover, 5-HT₂ antagonists were shown to reduce neuroleptic-induced catalepsy in rats (Balsara *et al.*, 1979; Hicks, 1990), to increase neuroleptic-induced dopamine turnover (Saller *et al.*, 1990), to cause a slight, indirect activation of midbrain dopaminergic neurons (Ugedo *et al.*, 1989), restoring phasic rather than tonic activity (Svensson *et al.*, 1989), and to reduce neuroleptic-induced extrapyramidal side effects (Reyntjens *et al.*, 1986;

Bersani *et al.*, 1990). Therefore it has been suggested that a critical balance between 5-HT₂ and dopamine D₂ receptor antagonism is essential for the beneficial effects of antipsychotic drugs (Meltzer *et al.*, 1989).

Clozapine, the prototype for atypical neuroleptics, shows higher affinity for 5-HT₂ than for D₂ receptors. However, it also shows a potent histamine H₁ receptor antagonism, which results in severe sedation. In addition, clozapine binds to 5-HT_{1C} receptors, to α ₁-adrenergic and to muscarinic receptors with 5-10 times higher affinity than to D₂ receptors. Therefore these additional properties may play a role in both the therapeutic and resultant side effects of clozapine treatment (Grohmann *et al.*, 1989). The relatively higher affinity of clozapine for D₄ receptors than for D₂ receptors was proposed as a possible underlying mechanism for the particular clinical properties of this compound (Van Tol *et al.*, 1991). Interest has consequently focussed on the ability of neuroleptic drugs to cause long-lasting alterations in basal ganglia function, by changing the pattern of gene expression in striatal neurons.

1.12 Long term effects on late response genes in the basal ganglia

In 1978, Hong and colleagues demonstrated that either chronic blockade of dopamine receptors by neuroleptic drugs or lesioning the nigrostriatal pathway could increase levels of enkephalin peptides in the striatum. Consistent with this, *in situ* hybridisation studies later revealed changes in mRNA levels of proenkephalin after chronic treatment with dopamine receptor antagonists. Chronic treatment with haloperidol, the relatively non-selective dopaminergic receptor antagonist, dramatically increased the levels of proenkephalin mRNA in the rat striatum (Sabol *et al.*, 1983; Tang *et al.*, 1983; Angulo *et al.*, 1986; Sivam *et al.*, 1986; Romano *et al.*, 1987; Morris *et al.*, 1988). *In situ* hybridisation

studies have also shown that unilateral nigral lesions induce a dramatic ipsilateral increase in striatal proenkephalin mRNA levels (Angulo *et al.*, 1986; Young *et al.*, 1986; Morris *et al.*, 1989; Cadet *et al.*, 1991), suggesting that the nigrostriatal pathway exerts a tonic inhibition on the activity of the striatal enkephalin-containing cells. The cells of the ventral tegmental area were demonstrated to exert a similar inhibition on the activity of the enkephalin cells of the nucleus accumbens (Morris *et al.*, 1989). More recently, the elevation in striatal proenkephalin mRNA levels following a nigrostriatal lesion, has been positively correlated with an increase in rotation rate after subcutaneous administration of amphetamine (Cadet *et al.*, 1991).

Conversely, studies have shown that the repeated administration of dopamine receptor antagonist drugs can decrease the concentration of substance P in the substantia nigra (Hong *et al.*, 1978; Hanson *et al.*, 1981; Le Douarin *et al.*, 1983; Oblin *et al.*, 1984; Sudzak and Gianutsos, 1984). Further evidence has demonstrated that the levels of protachykinin mRNA, the precursor protein for substance P, are decreased in the rat striatum after chronic treatment with the antipsychotic drug, haloperidol (Bannon *et al.*, 1986). From in situ hybridisation studies, the levels of striatal protachykinin mRNA has been shown to be decreased following a nigral lesion, suggesting a tonic dopaminergic facilitation of the activity of the tachykinin cells (Young *et al.*, 1986). Recent evidence suggests that the release of substance P is decreased in the ipsilateral substantia nigra after dopaminergic denervation of the striatum (Lindfors *et al.*, 1989).

In animal models of Parkinson's disease, met-Enk levels are increased in the striatum (Engber *et al.*, 1991; Dacko and Schneider, 1993), while substance P has been shown to be either unaltered (Sivam *et al.*, 1987) or decreased (Lindfors *et al.*, 1989). Ceballos and colleagues have more recently suggested that the population of neurons co-localising met-enkephalin and substance P can adapt according to the extent of degeneration in the substantia nigra in Parkinson's disease patients. Patients with greater than 80% loss of dopamine in

the caudate show an increase in both met-enkephalin and substance P levels. However, in patients with approximately 50% loss of dopamine content in the caudate, levels of both peptides were markedly reduced by 80%.

A decrease in somatostatin mRNA has been observed in the striatum of rats treated with haloperidol, suggesting that dopamine may be involved in the regulation of somatostatin mRNA levels (Salin *et al.*, 1988). Haloperidol decreases preprosomatostatin mRNA levels not only in the caudoputamen, but also in the nucleus accumbens and in the frontal cortex. Therefore ^{this} suggests a widespread influence of dopamine on somatostatin neurons in target areas of the mesencephalon. This is compatible with previous findings showing a decrease in immunoreactive somatostatin measured by radioimmunoassay in the striatum and in the nucleus accumbens after chronic haloperidol treatment (Beal and Martin, 1984; Radke *et al.*, 1988). It was suggested that this effect may be related to the blockade of a stimulatory action of dopamine on somatostatinergic neurons mediated through dopamine D2 receptors, since haloperidol was shown to preferentially block D2 receptors (Saller and Salama, 1986). However, it has since been shown that the region within the striatum, in which haloperidol significantly affects the levels of preprosomatostatin mRNA corresponds to an area of relatively low D2 dopaminergic binding sites (Dubois and Scatton, 1985; Boysen *et al.*, 1986). This implies that other mechanisms may be involved in the effect of haloperidol on preprosomatostatin mRNA levels in this brain region. Evidence to support this further came from the observation that lesions of the nigrostriatal dopaminergic neurons, which markedly reduce dopamine levels in the striatum do not affect the number of neurons expressing somatostatin-like immunoreactivity in the striatum ipsilateral to the lesion (Salin *et al.*, 1990).

The mRNA levels of the neuropeptide prodynorphin were not greatly affected by chronic neuroleptic drug treatment (Morris *et al.*, 1988). A similar lack of effect on expression of prodynorphin mRNA was observed following nigral lesions (Young *et al.*, 1986; Morris *et al.*, 1989).

Saller and colleagues (1989) found that chronic treatment with the atypical neuroleptic clozapine increased the intensity of preprosomatostatin mRNA in neurons, but not the number of neurons expressing the gene, in the nucleus accumbens. No effect was observed with clozapine in the striatum. The hypothesis that dopamine exerts opposite effects on the levels of preprosomatostatin mRNA in the striatum through an action on D1 and D2 dopamine receptors was thought to be compatible with this lack of effect in the striatum. However, it has been suggested that the blockade of other neurotransmitter receptors such as muscarinic, 5-HT₂, and noradrenergic receptors by clozapine (Kelley and Miller, 1975; Peroutka and Snyder, 1980) may be responsible for the different effects of clozapine and haloperidol on somatostatinergic neurons in the striatum (Saller *et al.*, 1990). This evidence substantiates the present suggestion for a differential effect of some antipsychotic drugs in the nigrostriatal and mesolimbic system (Chiado, 1988). The idea of a different mechanism of action of atypical and typical antipsychotic drugs has gained further support from the finding that suggests that clozapine, but not haloperidol, increases the levels of glutamic acid decarboxylase mRNA, the enzyme involved in the synthesis of GABA, in neurons of the globus pallidus (external pallidum), implying a differential effect of these antipsychotic drugs on one of the major output systems of the basal ganglia (Mercugliano *et al.*, 1989).

Dopaminergic receptor antagonists of D1 and D2 receptors have also been demonstrated to affect in opposite directions, the levels of immunoreactive neuropeptide Y, a peptide colocalised with somatostatin in striatal neurons (Sandell *et al.*, 1986; Smith and Parent, 1986). Chronic treatment with SCH23390, which preferentially blocks D1 dopamine receptors, increases the number of neurons expressing neuropeptide Y-like immunoreactivity in the striatum. The dopaminergic D2 receptor antagonists, haloperidol and sulpiride were shown to have an opposite effect (Kerkerian *et al.*, 1988). Similarly at the cellular level it was shown that neuropeptide Y mRNA could be clearly increased

in the rat striatum following chronic administration of the D1/D5 receptor antagonist SCH23390 for three days (Morris, 1989).

1.13 Effect of Antipsychotic Drug Treatment on Immediate Early Gene Expression

The antipsychotic drugs are effective in the treatment of severe psychiatric disorders. Although many exert serious side effects and have limitations in efficacy which in turn has spurred a search for better compounds and a greater understanding of the pharmacology of antipsychotic drugs. The full therapeutic effect of antipsychotic drugs generally takes several weeks to emerge. Therefore it has been postulated that their interaction with neurotransmitter receptors are merely the initial step in their actions, and that the therapeutic effects result from adaptive processes that may occur in response to repetitive or chronic receptor occupancy by the drugs. A likely candidate mechanism for such drug induced neural plasticity is receptor -mediated regulation of neuronal gene expression. It is currently the idea that the induction of cellular IEGs may be a critical signal transduction step in neural plasticity induced by neuroleptic drugs, with the protein products of IEGs functioning to activate or repress genes that encode proteins involved in the differentiated functions of target neurons.

Recent evidence has shown that the dopaminergic system in brain is capable of responding in an adaptive manner to neuroleptic drug exposure, by inducing a number of dopamine D2 receptors (Seeman, 1980). This neurotransmitter system has also been shown to participate in the induction of certain IEGs. The dopamine precursor L-DOPA has previously been demonstrated to activate the IEG *c-fos* in the rat striatum ipsilateral to a 6-hydroxydopamine lesion of the substantia nigra via stimulation of dopamine D1 receptors (Robertson *et al.*,

1989). In agreement with this, systemic administration of D1 receptor agonists, in particular SKF38393, have been shown to induce c-fos and zif/268 mRNA in striatal neurones of rat brain after lesioning (Robertson *et al.*, 1990; Cole *et al.*, 1992). In contrast, haloperidol, a relatively non-selective dopamine receptor antagonist together with selective dopamine D2 receptor antagonists such as YM 09151, eticlopride and (+) butaclamol have been shown to induce c-fos protein in rat striatal neurons (Miller, 1990; Deutch *et al.*, 1992; Sirinathsinghji *et al.*, 1994) and in the nucleus accumbens (Dragunow *et al.*, 1990; Deutch *et al.*, 1992), but not in other brain regions. Similarly, Nguyen and colleagues (1992) have shown that both c-fos and zif/268 can be induced in the caudate-putamen at the mRNA level by systemic administration of haloperidol or amphetamine. However the atypical neuroleptic, clozapine, was unable to induce c-fos mRNA in the caudate-putamen, but was able to induce the expression of c-fos in the nucleus accumbens. A similar induction of fos immunoreactivity has been demonstrated in the shell but not the core region of the nucleus accumbens in the rat after acute systemic clozapine treatment (Deutch *et al.*, 1992). Whilst the selective D1 dopamine receptor antagonist, SCH23390 was demonstrated to have no effect on the levels of fos protein (Dragunow *et al.*, 1990). A similar lack of effect on the expression of c-fos mRNA in rat striatum has been observed following systemic administration of the selective D2 receptor agonist LY 171555, but this compound was able to reverse completely the haloperidol induced expression of c-fos mRNA (Miller, 1990). It was later shown that a 6-hydroxydopamine lesion of the substantia nigra alone could lead to a long lasting expression of fos related antigens in striatal neurons ipsilateral to the side of the lesion (Dragunow *et al.*, 1991). At present there is circumstantial evidence linking the haloperidol induction of c-fos to the subsequent increase in proenkephalin mRNA (Sonnenberg *et al.*, 1989; Donovan *et al.*, 1992), but this has been questioned by recent results suggesting that it is not the fos protein that interacts with the proenkephalin enhancer, but it is a cAMP-responsive element

binding protein that appears to play a major interactive role (Konradi et al., 1993).

A number of psychomimetic drugs including D-amphetamine and cocaine, with markedly different pharmacologic and clinical properties, have also been reported to activate striatal c-fos expression and a clear drug-specific pattern of gene activation has been shown in neurons of the sensory-motor and limbic striatum (Robertson et al., 1989; Graybiel et al., 1990; Nguyen et al., 1992). The mechanism of induction of c-fos is thought to involve the stimulation of D1 dopamine receptors (Robertson et al., 1989; Graybiel et al., 1990). Therefore, both PKA and calcium or PKC signalling may underlie the induction of c-fos, as D1 receptors are positively coupled to adenylate cyclase (Stoof and Keibian., 1981) and to inositol phospholipid (Mahan et al., 1990) signalling pathways. It has previously been shown in vitro that the IEG c-fos can be induced through interaction of nuclear proteins with cAMP-responsive regulatory elements (CREs) present in the gene (Greenberg et al., 1985) and some PKC effects have been reported as well (Goodman, 1990).

It is known that c-fos can form dimers with members of the jun family of IEGs in order to effect target gene transcription. The mRNAs encoding two members of this family, c-jun and jun-D, are present constitutively in striatal neurons and the corresponding proteins are likely to be available to form complexes with c-fos when it is synthesised. The mRNA encoding another IEG belonging to this family, jun-B, is not normally detectable in striatal neurons. However there is some evidence for the induction of jun-B after acute administration of cocaine (Hope et al., 1992), amphetamine and apomorphine (Cole et al., 1992). These drugs are known to have agonist properties at a number of receptors, including dopamine receptors.

As yet there is still no firm evidence linking the induction of any immediate early gene with any effect on a late-reponse gene. However, there has been some speculation on a potential link between the induction of fos and jun

proteins and enkephalin mRNA expression (Sonnenberg et al., 1989) Although, contrary to this suggestion, more recent research implies that it is not the fos protein that interacts with the enkephalin enhancer (Konradi et al., 1993).

1.14 Characterisation of Serotonin Receptor Subtypes

Serotonin (5-hydroxytryptamine, 5-HT) is a biogenic amine neurotransmitter that is known to be involved in the homeostatic regulation of the fundamental physiological functions such as sleep, temperature regulation, learning and memory and pain. Disruptions in the serotonergic system may be a critical factor in many pathological disorders affecting society today, including anxiety, depression, schizophrenia, eating disorders and obsessive compulsive disorder (Lopez-Ibor, 1988). A centralisation of the serotonergic system in the raphe area of the central nervous system, together with its diverse projections to all brain areas confirms the potential of serotonin to be a homeostatic regulator.

The transduction of serotonergic signals across the neuronal membrane is mediated by a diversity of receptor subtypes that in mammals appear to fall into seven distinct classes, designated 5-HT₁ to 5-HT₇, on the basis of their affinities for serotonin or sequence similarities. With the exception of the 5-HT₃ receptor which is a ligand gated ion channel, all of the other members of the serotonin receptor family belong to the large class of receptors that are linked to their effector function via guanine-nucleotide binding proteins with the putative seven transmembrane domain structure characteristic of many receptors (Hoyer and Schoeffter, 1991). Within receptor classes, a number of serotonin receptor subtypes have been identified. The functional significance, if any, in terms of the physiological role of many of these newly cloned receptors is as yet to be identified and therefore is of great importance in understanding the functions of serotonin in the brain.

The 5-HT₁-like subcategory has been further subdivided into four different subtypes (Hartig *et al.*, 1992), which have recently been cloned, namely the 5-HT_{1A} (Kobilka *et al.*, 1987; Fargin *et al.*, 1988), the 5-HT_{1B} (Voigt *et al.*, 1991; Branchek *et al.*, 1991), the 5-HT_{1D} (Hamblin and Metcalf, 1991) in rat,

the 5-HT_{1D} in the canine (Zgombick *et al.*, 1991), the 5-HT_{1Db} (Demchyshyn *et al.*, 1992) in human and the 5-HT_{1E} (Hartig *et al.*, 1992) in the rat. All members of the 5-HT₁ subgroup are characterised by their high affinities for serotonin ($K_i < 100\text{nM}$) and their ability to inhibit cellular cAMP production (De Vivo and Mayaani, 1990). Distinction between the receptors in this group has been made on the basis of the affinities for a series of drugs showing some subtype selectivity e.g. 8-hydroxy DPAT is a selective agonist for the 5-HT_{1A} receptor (Gozlan *et al.*, 1983) and rawolscine a selective agonist for the 5-HT_{1D} receptor (Frazer *et al.*, 1990). 5-HT_{1A} receptors have been widely studied and agonist compounds acting at this receptor site have been shown to possess anxiolytic (Traber and Glasser, 1987; Gleeson *et al.*, 1989; Brocco *et al.*, 1990), antidepressive (Cervo *et al.*, 1988; Glasser, 1988; Glennon, 1990), antihypertensive (Hartog and Wouters, 1988; Mandal *et al.*, 1989; Stubbs *et al.*, 1991) and possibly antipsychotic (Ahlenius, 1989) and neuroprotective (Bielenberg and Burkhardt, 1990; Bode-Greuel *et al.*, 1990) properties. Indeed the 5-HT_{1A} partial agonists buspirone, ipsapirone and gepirone have been shown to exert anxiolytic and antidepressive effects in humans (Glennon, 1990; Robinson *et al.*, 1989; Traber and Glasser, 1987). The 5-HT_{1B} receptor was distinguished from the 5-HT_{1A} subtype with the use of the drug spiperone, as 5-HT_{1A} sites have high affinity for spiperone, while conversely the 5-HT_{1B} site has a low affinity for spiperone (Pedigo *et al.*, 1981). A similar distribution of 5-HT_{1A} receptors has been observed in many different animal species (Marcinkiewicz *et al.*, 1984; Waeber *et al.*, 1989). The highest density of 5-HT_{1A} sites was observed in the limbic system including the hippocampus, septum, amygdala and cortical limbic areas. This high concentration in the limbic system is thought to be consistent with their involvement in mood and anxiety. The recently cloned 5-HT_{1B} receptor subtype has been shown to have high affinity binding of ³H-5-HT with a different pharmacological profile to the 5-HT_{1A} subtype. In situ hybridisation has revealed that the 5-HT_{1B} mRNA is

located within cells of the dorsal and median raphe nuclei suggesting an autoreceptor function, in the CA1 region of the hippocampus, in the striatum, cortex and in the cerebellum. The human mRNA encoding the 5-HT_{1B} receptor has been shown to be most abundant in the striatum (Jin *et al.*, 1992). Both the 5-HT_{1B} and the 5-HT_{1D} receptor subtypes are defined on both pharmacological and biochemical grounds by their ability to inhibit adenylate cyclase activity (Schoeffter *et al.*, 1990). The 5-HT_{1B} receptor has been suggested to be the evolutionary antecedent of the 5-HT_{1D} receptor which is present in bovine and human brain in high concentrations in the substantia nigra and basal ganglia (Heuring and Peroutka, 1987). The 5-HT_{1E} receptor mRNA has been located on the terminal fields of some serotonergic neurons. Characterisation of this receptor subtype is still necessary.

Another subclass of 5-HT receptors is the 5-HT₂ subtype. At present there are three structurally similar subtypes referred to as 5-HT_{2A}, 5-HT_{2B} (formerly the 5-HT_{2F}), and 5-HT_{2C} (formerly the 5-HT_{1C}) (Pritchett *et al.*, 1988) whose activation results in phospholipase C-mediated phosphatidylinositol lipid hydrolysis (Conn and Sanders-Bush, 1986; Conn *et al.*, 1986) which liberates the second messengers diacylglycerol and inositol triphosphate. Diacylglycerol activates the enzyme protein kinase C, whereas inositol triphosphate increases the intracellular free calcium concentration (Jaken, 1989; Irvine, 1989). Both increases in intracellular calcium and activation of protein kinase C have been observed in response to activation of 5-HT_{2A} (Pritchett *et al.*, 1988; Berg *et al.*, 1994) and 5-HT_{2C} receptors (Lutz *et al.*, 1993; Boddeke *et al.*, 1993). In addition to coupling to phospholipase C-mediated phosphatidylinositol lipid hydrolysis, 5-HT_{2A} and 5-HT_{2C} receptors have been shown to mediate phospholipase A₂-mediated arachidonic acid release (Felder *et al.*, 1990). Long term exposure to agonists (Dillon-Carter and Chuang, 1989; Sanders-Bush and Breeding, 1990; Ivins and Molinoff, 1991) or antagonists (Blackshear *et al.*, 1983; Sanders-Bush and Breeding, 1988) have also been demonstrated to down

regulate these receptors. The newly cloned 5-HT_{2B} (formerly 5-HT_{2F}) receptor (Foguet *et al.*, 1992; Kursar *et al.*, 1992; Loric *et al.*, 1992) has recently been characterised. The 5-HT_{2B} receptor subtype was shown to stimulate production of inositol 1,4,5-trisphosphate in transformed cells and has a unique pharmacological profile consistent with the evidence that this receptor subtype mediates contractions in the rat stomach fundus (Wainscott *et al.*, 1993).

The distribution of 5-HT₂ receptor subtypes shows some overlap, although there are some differences. Both the 5-HT_{2A} and 5-HT_{2C} subtype mRNA have been shown to be present at intermediate levels in the caudate-putamen, accumbens nucleus and substantia nigra pars compacta. However, their distribution in the caudate-putamen is quite different. The 5-HT_{2C} receptor mRNA is in addition present in the basal nucleus of Meynert, stria terminalis, entopeduncular nucleus and subthalamic nucleus and was found to be most predominant in the amygdala. While the 5-HT_{2A} transcripts were restricted to only a few nuclei in these areas (Pompeiano *et al.*, 1993). It has very recently been shown that there are important fundamental differences in the signal transduction systems for 5-HT_{2A} and 5-HT_{2C} subtypes. In transfected CHO cells activation of 5-HT_{2C}, but not 5-HT_{2A} receptors reduces receptor mediated inhibition of adenylate cyclase activity. Furthermore, this effect of the 5-HT_{2C} subtype is independent of activation of PKC and an increase in intracellular calcium concentration (Berg *et al.*, 1994). Attempts to visualise transcripts for the third member of the 5-HT₂ receptor family, the 5-HT_{2B} (5-HT_{2F}) with probes and PCR analysis has so far failed in rat brain (Foguet *et al.*, 1992).

The 5-HT₃ receptor is a ligand-gated ion channel which when activated causes a fast depolarising response. Kilpatrick and colleagues (1987) were the first to identify 5-HT₃-binding sites in the brain. More recently, Maricq and colleagues have isolated a cDNA clone encoding a 5-HT₃ receptor from NCB-20 cells (Maricq *et al.*, 1991). The mRNA encoding this receptor is identified in

brain, spinal cord and in the heart. The receptor protein is located mainly in the dorsal hindbrain, nuclei of the cranial nerves, and the area postrema (Palacios *et al.*, 1990).

A 5-HT₄ receptor subtype is also known to exist, however it is the least well characterised of all the 5-HT receptors. The 5-HT₄ receptor has been detected in the CNS by measuring 5-HT stimulation of cyclic AMP formation either in intact neurons in culture (Dumuis *et al.*, 1988, 1989) or in isolated neuronal membranes (Bockaert *et al.*, 1989, 1990), or in the gastrointestinal tract by monitoring 5-HT-mediated changes in contractile responses of the guinea pig ileum (Clarke *et al.*, 1989; Craig and Clarke, 1990). This receptor has also been characterised in radioligand binding studies (Clarke *et al.*, 1989), but has not yet been cloned.

A fifth member of the 5-HT receptor family has been isolated from mouse brain (Plassat *et al.*, 1992). This 5-HT receptor is referred to as the 5-HT_{5A}, as a 5-HT_{5B} subtype has recently been sequenced (Matthes *et al.*, 1993). 5-HT₅ receptors are known to have very high affinity for ergot compounds such as lysergic acid diethylamide and methysergide, but have relatively low affinity for 5-HT. The 5-HT_{5A} subtype was found to be expressed predominantly in the cerebellum, cerebral cortex, hippocampus and olfactory bulb. While the 5-HT_{5B} receptor mRNA was expressed predominantly in the medial habenula and in the CA1 field of the hippocampus and raphe nuclei.

A recent report has described the cloning of another structurally and pharmacologically distinct 5-HT receptor from rat brain, namely the 5-HT₆ receptor. This receptor subtype has been shown to exhibit high affinity for a number of antipsychotic and tricyclic antidepressant drugs, including clozapine, amoxapine and amitriptyline (Monsma *et al.*, 1993). In cells transfected with this receptor, serotonin elicits a potent stimulation of adenylyl cyclase activity which is blocked by antipsychotic and antidepressant drugs (Sebben *et al.*, 1994). This receptor subtype has been localised in motor limbic and cortical regions of the

brain, with the highest levels present in the striatum, a potentially important site of action of antipsychotic drugs with regard to both extrapyramidal symptoms and perhaps cognitive functioning. (Monsma *et al.*, 1993). A similar 5-HT receptor subtype has recently been isolated in mouse neuroblastoma cells (Unsworth and Molinoff, 1993).

Most recently a 5-HT₇ receptor subtype has been proposed (Shen *et al.*, 1993). The serotonergic 5-HT₇ receptor has since been cloned and shown to be positively coupled to adenylate cyclase when transfected into a chinese hamster ovary cell line (Tsou *et al.*, 1994). Similar to the 5-HT₆ receptor, this receptor subtype has high affinity for antipsychotic drugs such as clozapine. The 5-HT₇ receptor has not yet been fully characterised, however, both the 5-HT₆ and 5-HT₇ receptor subtypes are known to pharmacologically resemble the 5-HT₂ receptor (Monsma *et al.*, 1993; Shen *et al.*, 1993).

1.15 Drug Pharmacology of Serotonin receptors

Early on in the studies on neuroleptic drugs, evidence was obtained demonstrating that the serotonergic system could influence the action of antipsychotic drugs (Kostowski *et al.*, 1972) and some neuroleptic drugs could increase 5-HT turnover, perhaps by blockade of 5-HT receptors (Leysen *et al.*, 1978). With the discovery of specific agonists and antagonists for serotonin receptor subtypes, a potential role of particular clinical importance was evident, since the binding affinities of many atypical antipsychotics such as clozapine, melperone and amperozide (Meltzer *et al.*, 1989; Christensson *et al.*, 1990) are particularly high for certain serotonin receptor subtypes.

In vitro, the pK_i values of typical neuroleptic drugs for 5-HT₂-binding sites tends to vary over a wide range, with a number being quite potent, such as

methiothepin and spiperone ($pK_i = 9.4$) (Meltzer *et al.*, 1989). Both fluphenazine ($pK_i = 8.6$) and haloperidol ($pK_i = 7.7$) have some affinity for the 5-HT_{2A} receptor subtype (Meltzer *et al.*, 1989; Audinot *et al.*, 1993). Haloperidol has a much lower affinity for the 5-HT_{2C} serotonin receptor subtype (Audinot *et al.*, 1993). Previous studies have also demonstrated that haloperidol has only a weak affinity for the 5-HT_{1A}, 5-HT_{1B}, 5-HT_{1D} and 5-HT₃ receptors (Meltzer *et al.*, 1989). Therefore it has been suggested that haloperidol would not be expected to achieve effective blockade at clinically relevant doses at 5-HT_{1D}, 5-HT_{2C}, or 5-HT₃- binding sites in humans with the possible exception of weak 5-HT_{2A} receptor blockade, because the clinical dose is based on its potent D₂ receptor antagonism ($pK_i = 9.0$) (Meltzer and Nash, 1991).

Many atypical antipsychotic drugs have been found to have a relatively high affinity for 5-HT₂-binding sites *in vitro* as compared to the typical drugs relative to their respective affinities for D₂ dopamine receptors (Meltzer *et al.*, 1989). Clozapine has been shown to have high affinity for both the 5-HT_{2A} ($pK_i = 8.3$, Meltzer *et al.*, 1989; $pK_i = 8.3$, Roth *et al.*, 1991; $pK_i = 7.6$, Audinot *et al.*, 1993) and the 5-HT_{2C} receptor ($pK_i = 8.14$, Roth *et al.*, 1991; $pK_i = 8.1$, Audinot *et al.*, 1993). As yet, it is not clear whether atypical drugs, like clozapine, are superior in treating schizophrenia compared with typical antipsychotic drugs, and whether they also do not produce tardive dyskinesia. It is likely that at clinically effective doses of clozapine, which correlate with its affinity for the D₂ receptor (Seeman, 1981), clozapine produces effective occupancy of the 5-HT_{2A} and 5-HT_{2C} receptors because it has a significantly higher affinity for these receptors than for the D₂ receptor. Clozapine has also been reported to have submicromolar affinity for 5-HT_{1A} receptors in human brain tissue (Mason and Reynolds, 1992) and a moderate affinity for the 5-HT₃ receptor (Watling *et al.*, 1990).

There is a possibility that, because of pharmacokinetic and pharmacodynamic reasons, in vitro affinities may not predict in vivo occupancies of specific drugs. Active metabolites, ^{may} rapidly dissociate from a receptor site ^{there may be} or local differences in drug metabolism due to differences in brain cytochrome P450 enzymes may cause significant differences between in vivo occupancy and in vitro affinities (Meltzer and Nash, 1991). Meltzer *et al.* (1989) have previously demonstrated that some drugs such as chlopromazine, are significantly more effective in blocking D2 receptors in vivo than would be predicted from their in vitro affinities.

The highly selective D1-like receptor antagonist, SCH23390, has also been reported to interact with the central 5-HT system (Bischoff *et al.*, 1988). Similar to selective 5-HT2 receptor blockers, SCH23390 was shown to inhibit [³H]-spiperone binding in the rat frontal cortex, in vivo. In vitro, SCH23390 was shown to inhibit [³H]-ketanserin binding to 5-HT2 sites. Biochemical parameters linked to dopamine and 5-HT were not changed, except in the striatum where SCH23390 increased homovanilic acid levels. SCH23390 was shown to have only a weak affinity for 5-HT1B and 5-HT1A receptors.

In general terms, the following research work was performed in order to understand further the role of the immediate early gene in striatal functioning, which at present is unclear. We have attempted to investigate how major transmitter systems, in particular the dopaminergic and serotonergic systems, have the ability to activate certain immediate early genes known to be present in the striatum. With the use of a number of selective drugs, novel molecular biological approaches and an in vitro 'model' of the striatum, a wide spectrum of possible receptor mechanisms will be investigated. The ultimate aim of which will be to isolate a receptor mechanism and second messenger pathway prior to the induction of an immediate early gene under investigation.

Since a number of disease states including Parkinson's disease and schizophrenia have been identified, and in turn linked to a dysfunction of the basal ganglia, we were particularly interested in the neuropeptide systems in the striatum which are known to be vulnerable in many neurological diseases. Neuroleptic drugs are the conventional treatment for schizophrenia, therefore to understand how neuroleptics effect transmitter / neuropeptide systems in the striatum may provide an important insight into certain diseases. Recently it has been proposed that the activation of immediate early genes may be connected to a change in the expression of a late response gene (for example, a gene encoding a neuropeptide). Therefore, to aid in the understanding of this potential relationship, we have also investigated how neuroleptics effect a number of immediate early genes. Hopefully this research will provide useful information on the complex functioning of the basal ganglia.

Chapter 2

METHODS

2.1 Oligonucleotide Probes

Oligonucleotide probes of unique sequence were synthesised on an Applied Biosystems DNA synthesiser. The probe sequences were as follows: c-fos (45-mer), complementary to nucleotides spanning amino acids 1-15 (Curran et al., 1987) ; Jun-B (60-mer), complementary to nucleotides spanning the last 20 amino acids of the predicted protein (Ryder et al., 1988) ; Jun-D (60-mer), complementary to the last 20 amino acids of the predicted protein (Ryder et al., 1989) ; zif/268 (45-mer), complementary to nucleotides spanning amino acids 2-16 (Millbrandt, 1987) ; preproenkephalin (45-mer), complementary to bases 388-432 of the predicted gene sequence (Howells et al., 1984; Yoshikawa et al., 1984) ; prodynorphin (45-mer), complementary to bases 862-906 of the predicted gene sequence (Civelli et al., 1985; Morris et al., 1986) ; proneuropeptide Y (45-mer), complementary to bases 1624-1668 of the predicted gene sequence (Larhammar et al., 1987) ; preprosomatostatin (45-mer) , complementary to the sequence encoding for amino acids 78-92 of the predicted protein (Montminy et al., 1984) ; preprotachykinin (36-mer), complementary to bases 172-207 of the predicted gene sequence (Krause et al., 1987) ; BF-1 (45-mer), complementary to bases 445-489 (Tao and Lai, 1992) ; RHS2 (45-mer), complementary to bases 35-79 (Le Moine et al., 1992) ; D1 dopamine receptor (45-mer), complementary to bases 142-186 (Monsma et al., 1990) ; D2 dopamine receptor (45-mer), complementary to bases 640-684 (Bunzow et al. , 1988), and the 5-HT6 receptor (45-mer), complementary to bases 702-746 (Monsma et al., 1992); Tubulin T26 (45-mer) complementary to part of the 3' non-coding region:

GGGAAACAGCATAGAAGCATCGATGCCTGC AGCTAGTGCT GGAGC (Miller et al., 1987); D1 dopamine receptor antisense complementary to bases 1-15 (Sunahara et al., 1990) ; D5 dopamine receptor antisense complementary

to bases 1-15 (Sunahara *et al.*, 1991) and a D1 sense. A sense "message" probe for preprosomatostatin and Microtubule-associated protein (MAP2) were also synthesised.

2.2 In situ Hybridisation on cryostat sections

In situ hybridisation was performed according to the method described previously (Morris *et al.*, 1986, Wisden *et al.*, 1991 ^{Wisden and Morris, 1994}). Frozen coronal brain sections (20 μ m) were cut using a cryostat (Reichert-Jung) at the level of the rostral and medial striatum (taken 1.6mm anterior and 0.4mm posterior to bregma). The sections were mounted onto sterile fridge-cooled slides (BDH) previously baked in aluminium foil at 180°C and coated in a 0.01% solution of poly-L-lysine hydrobromide made up in Diethylpyrocarbonate (DEPC)-treated water. The sections were then dried at room temperature. Following this, the sections were fixed in an ice cold 4% (weight/volume) solution of depolymerized paraformaldehyde freshly prepared in 1x phosphate-buffered saline (1xPBS) [10 x PBS is 1.3M sodium chloride, 70mM di-sodium hydrogen orthophosphate, 30mM sodium di-hydrogen orthophosphate made up in sterile water.] for 5 minutes. The sections were then transferred into 1xPBS for 1-2 minutes and then dehydrated in graded ethanol solutions (i.e. 70%, 95%, and 100%) for several minutes.

0.6pmol oligonucleotide probes (45-mers) were 3' -end labelled with 10pmol of the isotope 5- $[\alpha$ - ³⁵S]dATP (NEN, specific activity 1300 Ci/mmol), i.e. using a 33:1 molar ratio of isotope to oligonucleotide and with terminal deoxynucleotidyl transferase (TdT, Pharmacia). The specific activity of the probes was in the region of 1.0×10^9 dpm/ μ g.

For a 12.5 μ l reaction volume, in a sterile eppendorff tube mix together:
4.0 μ l DEPC-water

2.5 μ l 5 x reaction buffer. Reaction buffer (5x) contains 1M potassium cacodylate, 125mM Tris/HCl, 1.25mg ml⁻¹ bovine serum albumin, pH 6.6, at 25°C.

2.0 μ l of oligonucleotide at concentration of 0.3pmol μ l⁻¹ for a 45-mer, 0.3pmol is equivalent to 5 ng μ l⁻¹)

2.8 μ l [α -³⁵S]dATP (1300 Ci/mmol)

1.2 μ l TdT .

The reagents were mixed very carefully by pipette. Then ^{the mixture was} incubated for 1-3 hours in a water bath at 30-35°C. The reaction was ^{then} inhibited by adding 40 μ l DEPC-treated water. A Sephadex G-25 spin column was then prepared using a 1ml syringe plugged at the tip with glass wool. The syringe was filled with Sephadex G-25 solution (i.e. a 1:1 solution of sephadex G-25 powder and Tris/EDTA buffer[10mM Tris, 1mM EDTA] and spun for 1 minute to obtain a column. Then 50 μ l of the reaction mixture was added to the Sephadex G-25 spin column and spun at 2000 rpm for 1 minute in a MSE MINOR centrifuge, to remove unincorporated nucleotides. After collecting the eluate in a sterile eppendorff tube, 2 μ l of the eluate was analysed by liquid scintillation counting. The counts would be expected to be in the range 300 000 to 800 000 d.p.m. 2 μ l⁻¹. Then 2 μ l of a 1M solution of dithiothreitol (DDT) made up in sterile water was added to the tube. This is known to preserve the probe from oxidation. The probe was stored at -20°C until use.

Sections were removed from 100% ethanol and allowed to air dry. Following this, the sections were hybridised with the selected probe at 42°C overnight. In a sterile eppendorff tube, the radiolabelled probe was diluted 1:100/1:200 in a hybridisation buffer containing 50% deionised formamide, 0.02% poly adenylic acid, 10% 4x standard saline citrate (4x SSC) [1x SSC= 0.15M sodium chloride, 0.015 M sodium citrate, pH= 7.0], 5% 0.5M sodium phosphate, pH 7.0, 1% 0.1M sodium pyrophosphate and 10% dextran sulphate made up in sterile water. After adding 2 μ l of DTT to the mixture and vortexing

thoroughly, 120 μ l of probe/hybridisation buffer was applied to each slide. The slides were gently covered with a parafilm coverslip. Any air bubbles present were removed with blunt ended forceps. A small piece of tissue paper was saturated with 4 x SSC and placed alongside the slides in a sealed petri dish to maintain humidity. The slides were then incubated overnight at 42°C.

After overnight incubation slides were transferred into coplin jars containing 250ml of prewarmed 1 x SSC at 55°C. The parafilm was gently peeled off and discarded. The slides were washed twice in 1 x SSC at 55°C for approximately 30 minutes per wash with constant agitation. Following this, the slides were transferred into 0.1 x SSC at room temperature for 15 minutes, and then briefly dehydrated through 70% ethanol, 95% ethanol and 100% ethanol, before being allowed to air dry.

The sections were then exposed to autoradiographic film (Kodak XAR-5) at room temperature. Dry slides were attached to paper with scotch tape and exposed to autoradiographic film for 3 days to 2 weeks depending on the probes used and the efficiency of their tailing. Under safe light conditions films were developed for 1 minute in D19 (Kodak), rinsed in water for 30 seconds, fixed in Unifix fixative (Kodak) for 4 minutes and finally rinsed in water for 20 minutes. Once the films had been developed, the slides were dipped in Ilford K5 liquid photographic emulsion. Slides were dipped under safe light in a filtered 1:1 solution of emulsion : distilled water containing 0.5% glycerol prewarmed to 45°C. Slides were allowed to dry before being transferred to light-tight slide boxes containing fresh silica gel. The boxes were stored in the fridge at 4°C until time of development. As a general rule slides were stored for 5 X the exposure time on film.

Exposed boxes were allowed to come to room temperature. Under safe light conditions the slides were then transferred into glass racks prior to immersion in 250ml D19 developer (Kodak) at 19°C for 2 minutes. After developing, slides were immersed in water for 30 seconds and then transferred

into 250ml of a freshly prepared solution of 30% sodium thiosulphate for 4 minutes. The slides were then transferred into water for at least 30 minutes in order to wash off excess fixative. Sections were then counterstained in either a 0.2% solution of neutral red for 2-5 minutes or a 0.25% solution of cresyl violet for 5 minutes and dehydrated in serial solutions of ethanol for several minutes. Finally sections were transferred into histoclear for several minutes. Excess histoclear was allowed to drain off, following which the slides were mounted with glass coverslips (Chance) and DPX mounting medium.

2.3 In situ hybridisation of cell cultures

The method used for in situ hybridisation of cell cultures was essentially identical to that described for in situ hybridisation of cryostat sections. However we found it necessary to dilute 'hot' probes to 1:400 with hybridisation buffer to reduce the risk of background labelling to a minimum.

Autoradiographic films and slides were developed as previously described. Cultures were counterstained with a 0.2% solution of neutral red for 1 minute or with a 0.25% solution of cresyl violet for 1-5 minutes before dehydration and coverslipping.

2.4 Controls for in situ hybridisation

For in situ hybridisation using synthetic oligonucleotides, two types of control experiment were employed.

1. Competition hybridisations - where sections/cultures were hybridised with a 25 -fold excess of the unlabelled oligonucleotide probe.

2. Sections/cultures were hybridised with a 'sense' oligonucleotide probe having the same length, base composition and specific activity as a region on the messenger RNA (mRNA).

2.5 RNA Extraction

The brains from six wistar rats (250-300g) were removed and tissue from several brain regions including the cerebellum, hippocampus, striatum, cortex and olfactory bulb was carefully dissected out and rapidly frozen on dry ice. The brain tissue was weighed individually and stored at -80°C in sterile plastic tubes until extraction.

Total RNA was extracted using the guanidinium/ hot phenol method previously described by Chomzynski and Sacchi in 1987. Under sterile conditions, tissue of predetermined weight was put into a 50ml plastic tube containing 5ml/g guanidinium isothiocyanate solution (guanidinium isothiocyanate solution = 100g guanidinium thiocyanate powder, 5% 1M Tris, pH 7.6, and 2% 0.5M EDTA made up to 94.84mls in DEPC-treated water. Heat solution to 60°C, sterile filter, and add 21.2mls 20% sarcosyl and 2.1mls β - mercaptoethanol such that the total volume = 212mls) and a few drops of antifoam. Using a polytron homogeniser at setting 7, tissue was homogenised for a few minutes. An equal volume (5ml/g) of phenol preheated to 60°C, was then added and the tissue allowed to stand in a water bath at 60°C. Following this, 2.5ml/g preheated salt solution, solution III, was added at 60°C (solution III = 100mM sodium acetate, pH 5.2, 10mM Tris/ hydrochloric acid, pH 7.4, and 1mM EDTA). Then 5ml/g chloroform, maintained at room temperature, was added to the tissue and the tube shaken vigorously for 5-10 minutes whilst in the water bath at 60°C. The tissue was then cooled on ice and centrifuged at

4000rpm for 10 minutes at room temperature. The aqueous phase was removed to another tube and re-extracted with phenol/chloroform (volume ratio = 3:1). Following this, the aqueous layer was removed to another tube and re-extracted four times with phenol/chloroform (volume ratio = 1:1) and finally extracted with chloroform alone (volume ratio = 1:1). Then 2.5 volumes of ethanol was added to precipitate the RNA. At this stage strands of RNA were visibly forming in the ethanol. After overnight storage at -20°C an RNA pellet was formed. The tube was spun for 10 minutes at 4000rpm at room temperature, and then the pellet resuspended in 50µl DEPC-treated water and diluted 1:100. The optical density at 260/280 was measured, and 10µl aliquots stored at -80°C until use.

2.6 Northern Blot Analysis

The method for northern blotting was performed according to the method previously described by Wisden *et al.*, in 1988. Firstly a formaldehyde gel was prepared in a total volume of 150mls by heating together 1.64g agarose powder with 96.4mls DEPC-treated water in a conical flask previously baked to 180°C. Once melted, the gel was allowed to cool slightly. Following which, 23.6mls formaldehyde was added together with 30mls DEPC-treated 5 X Running buffer (5 X Running buffer = 200mM morpholinopropanesulphonic acid (MOPS), pH 7.0, 50mM sodium acetate, pH 7.0, and 5mM EDTA, the solution appears yellow after autoclaving) in a fume hood. Meanwhile the sides of an ethanol cleaned perspex gel holder were taped securely with scotch tape. The comb was attached to one end approximately 2.0mm from the base. The gel holder was then positioned in a horizontal gel electrophoresis apparatus (BIO-RAD) previously sterilised with ethanol. Warm gel was slowly

poured into the holder to ensure no bubble formation and allowed to set. Once the gel had set, the tape and the comb were removed. A 1 x MOPS solution was added to the apparatus and the gel was then pre-run at 65 volts for 1 hour. During this time brain tissue RNA samples were prepared by adding together:

10 μ l aliquots of p(A) + RNA

4.6 μ l 5 X Running buffer

3.5 μ l formaldehyde

10 μ l formamide.

The sample was vortexed and left for 15 minutes at 55°C in order to denature the RNA. The sample was then chilled on ice. 4 μ l loading buffer (loading buffer = 50% glycerol in water, 1mM EDTA, 0.4% bromophenol blue, and 0.4% xylene cyanole made up in DEPC-treated water) was added and then the samples were allowed to run for 4 hours at 120 volts or until the bromophenol dye had migrated three-fourths down the gel. A standard DNA sample was run alongside the tissue samples as a marker. Then the gel was removed from the gel holder and placed onto clingfilm. After cutting the standard DNA sample away from the gel, it was immersed in a ethidium bromide solution (1 μ g/ml) for 15 minutes and visualised under ultra violet light. The gel was then photographed on an ultraviolet transilluminator, before being blotted to record marker positions. The gel to be blotted was then transferred directly onto nylon membrane. After soaking approximately ten pieces of Whatman 3MM filter paper approximately 23 x 50cm, in 10 X SSC, the filter paper was placed onto a tray. Carefully the gel was then placed upside down (i.e. sample well openings down) on top of the filter papers. Following this, a piece of hybond filter paper (nylon membrane) was cut to fit the gel and placed on top and smoothed out carefully. Then, 2 x 3MM wet filters (Whatman) were packed on top followed by a box of tissues. Finally a weight was placed over the gel to ensure good transfer. The gel was then left overnight to allow the transfer process to take place. By capillary action, the 10 X SSC would be expected to

soak up the tissues and in the process, the single strand RNA in the gel would be transferred to the underside of the nylon membrane. The RNA was then crosslinked by ultraviolet irradiation onto the nylon membrane. The blot could then be stored in the freezer at -80°C until use.

Northern blots were hybridised with oligonucleotide probes previously 3'- end labelled with the enzyme TdT and the isotope ^{32}P -dATP (6 000 Ci/mmol, Dupont) at 42°C overnight. The probes were labelled to a specific activity of 1.0×10^{10} dpm/ μg . The method of probe labelling was identical to the method previously described under in situ hybridisation with ^{35}S - labelled probes, however the quantity of components within the reaction mixture were adjusted according to the radioisotope, such that the reaction mixture contained:

- 2.5 μl 5 X buffer
- 1.25 μl oligonucleotide
- 1.55 μl DEPC-treated water
- 6.0 μl ^{32}P - dATP
- 1.2 μl TdT.

When using the isotope ^{32}P -dATP, a high energy β - particle emitter, all procedures should be shielded by perspex to avoid radiation contamination. Blots were prehybridised with 10ml hybridisation buffer for 1 hour at 40°C . Following this, the blot was hybridised with a 3'- end labelled probe overnight at 42°C in a heat-sealed plastic bag. The following day, the blot was washed in 1 X SSC at 55°C for 1 hour, then in 1 X SSC for 15 minutes at room temperature, and finally briefly rinsed in 0.1 X SSC at room temperature. The blot was then covered with cling film and smoothed. Finally the northern blot was exposed to Kodak XAR-5 autoradiographic film at -80°C with an intensifying screen for approximately 2-5 days. Development of the film was as previously described. The blot was then washed in 0.5 X Denhardt's solution, 25mM Tris-HCl, pH 7.5, and 0.1% sodium dodecyl sulphate for 60 minutes at

90-95°C, followed by a wash in distilled water at 65°C for 15 minutes. The nylon membrane was then reusable for hybridisation with another ³²P-labelled probe.

2.7 Primary Neuronal Culture

Striatal neurons were cultured using a modification of the methods of Thayer et al (1986), Surmeier et al (1988) and Vilijn et al (1988). Timed pregnant Wistar rats (200-300g) were killed with an overdose of anaesthetic injected intraperitoneally (Sagatal) on gestational day 17 (E17). Embryos were removed from the uterus, decapitated and placed in ice cold calcium-free Earle's Modified Eagles Medium (EMEM). Following removal of the brains from the crania, the dorsal striata of each embryo was dissected out in cold EMEM using the optic chiasm as the posterior marker (Mísgeld and Dietzell, 1988). From each dissecting session the tissue was pooled from ten embryos (on average). The tissue was then minced and washed three times in cold EMEM. Cells were dissociated in 10ml of a 0.67% trypsin solution made up in sterile saline, at 37°C for 45 minutes. The trypsin was inactivated with 1.5mls heat-inactivated foetal calf serum (FCS). Following trypsinisation the cells were washed three times in cold EMEM and spun at 500g in a DAMON/IEC centrifuge for 2 minutes per wash. The cells were then resuspended in Dulbecco's Modified Eagles medium (DMEM) supplemented with 20% FCS (volume/volume), and penicillin-streptomycin solution (100 units penicillin, 100µg/ml streptomycin). The tissue was triturated in supplemented DMEM using a pasteur pipette and cells were seeded at a density of approximately 0.15×10^6 cells/cm² as monolayer cultures onto either sterile glass coverslips in 24-well plastic culture plates (Nunc) or onto 8-well multichamber glass slides

(Nunc) previously coated with poly-D-lysine (4 μ g/ml) for 24 hours, rinsed in EMEM and then coated with laminin (6 μ g/ml) for 6 hours. The cultures were grown in a humidified incubator supplied with 95% oxygen and 5% carbon dioxide. After 24 hours the medium was changed to DMEM supplemented with 10% horse serum, and on day 5 the medium was replaced with DMEM/10% horse serum containing 20 μ M cytosine arabinoside in order to control proliferation of glial and other non-neuronal cells. On day 9 the cytosine arabinoside was omitted. The culture medium was changed every 4 days thereafter in the 24-well culture plates used for immunocytochemistry. However for the cells seeded onto multichamber slides to be used for drug stimulation, the medium was changed to DMEM supplemented with 'Ultrosor G' (Gibco, BRL), and penicillin-streptomycin solution. This was necessary to obtain serum-free conditions. Cells were generally used 12-20 days after seeding. Prior to use, cells were fixed in a 4% buffered paraformaldehyde solution.

2.8 Immunohistochemistry

Prior to immunohistochemistry, cultures were fixed in an ice cold 4% solution of buffered formalin in 0.1M phosphate-buffered saline (PBS). The cell culture wells were rinsed three times in 0.1M PBS for 10 minutes and then incubated in 0.1M PBS containing 20% normal goat serum (NGS), at room temperature for 60 minutes. The cultures were then incubated overnight with specific antibodies against neurone specific enolase (NSE), met-enkephalin (met-ENK), and glial fibrillary acidic protein (GFAP), at dilutions of 1: 20 000, 1: 60 000, and 1: 40 000 respectively, in 50mM PBS with 1% NGS, 1.5% sodium chloride, and 0.5% Triton X-100 added. After rinsing in PBS three

times for 5 minutes per rinse, the cultures were incubated with biotinylated secondary antibody (Vector) for 60 minutes, rinsed in 0.1M PBS twice for 10 minutes, and then incubated with ABC-conjugated horse radish peroxidase (ABC Elite kit, Vector) for 60 minutes in accordance with manufacturers instructions. The peroxidase was visualised with 0.1% diaminobenzidine made up in 0.1M phosphate buffer, pH 7.2, containing 0.02% hydrogen peroxide and 0.05% nickel chloride. 50 μ l of filtered DAB chromogen was added to each well. A blue/grey reaction product was visible 2-15 minutes after the addition of DAB chromogen.

2.9 NADPH-Diaphorase Staining

NADPH-diaphorase histochemistry was performed according to the method of Valtschanoff et al (1992). Tissue culture slides or slide-mounted sections (50 μ m) of rat brain were fixed by immersion in 4% paraformaldehyde solution in 0.1M phosphate buffer, pH 7.2. Slides were preincubated in 0.1M phosphate buffer containing 0.25% Triton X-100 for 5-10 minutes and then transferred to a freshly prepared buffer-Triton solution containing 0.5 mg/ml β -NADPH and 0.2mg/ml nitro blue tetrazolium in a light-tight container. The slides were incubated in this solution at 37°C for 1-2 hours. After incubation the slides were rinsed in phosphate buffer, dehydrated in serial solutions of alcohol and finally coverslipped with DPX mounting medium.

2.10 Acetylcholinesterase Staining

Adult rat striatal sections and cultured striatal neurones were collected on slides as previously described. The slides were then washed four times in distilled water for 3 minutes per wash and incubated in a modified 'Koelle' medium containing 500 μ g/ml copper sulphate, 750 μ g/ml glycine, 1.15mg/ml acetylthiocholine iodide, and 0.05mg/ml ethopropazine, in sodium acetate buffer, pH 5.0. The slides were incubated in modified 'Koelle' medium for approximately three hours at 37°C. After incubation the slides were washed four times, 3 minutes per wash, in distilled water and then immersed in an ammonium sulphide solution made up in 0.5M acetic acid, for 60 seconds. Following this, the slides were washed four times in distilled water for 3 minutes each wash. The stain was then enhanced with a 0.25% silver nitrate solution for 30 seconds followed by four rinses of 3 minutes each in distilled water. Finally slides were dehydrated in 70%, 95%, and 100% ethanol respectively for 4 minutes each and coverslipped in DPX mounting medium.

2.11 Drugs and Compounds Used

Tris[hydroxymethyl] aminomethane, dimethyl sulphoxide, N,N-dimethyl formamide, ethylenediaminetetraacetic acid, (3-[N-Morpholino]propanesulfonic acid) MOPS, cyproheptadine hydrochloride, cresyl violet acetate, poly-L-lysine hydrobromide, poly-D-lysine hydrobromide, 3-hydroxytyramine (dopamine), 5-hydroxytryptamine, fluphenazine dihydrochloride, haloperidol, nickel chloride hexahydrate, nitro blue tetrazolium, hydrogen peroxide, cytosine β -D-arabinoside, DL- Dithiothreitol, diethylpyrocarbonate, acetylthiocholine iodide, ethopropazine, phorbol-12-myristate-13-acetate, polyadenylic acid potassium salt, β - nicotinamide adenine dinucleotide phosphate tetrasodium salt, glycerol, β - mercaptoethanol, diaminobenzidine, triton X-100, phorbol-12 myristate-13-acetate (Sigma). Methiothepin mesylate, mesulergine hydrochloride, (\pm) -1-(2,5-dimethoxy-4-iodophenyl)-2-aminopropane hydrochloride (DOI), S(-)-eticlopride hydrochloride, (\pm) SKF-38393, R(+) - SCH-23390 hydrochloride, spiperone hydrochloride, (\pm) -8-hydroxy-2- (di-n-propylamino)-tetralin hydrobromide, 1-(3-chlorophenyl)piperazine 2-hydrochloride, (-)-quinpirole hydrochloride (RBI). Sodium chloride, sodium dihydrogen orthophosphate, disodium hydrogen orthophosphate, sodium thiosulphate, sodium citrate, DPX mountant, sodium acetate, sodium hydroxide, copper sulphate (BDH). Silica gel 4-7 mesh (Fisons). Antifoam A, polyethylene glycol, guanidinium thiocyanate, sodium N-lauroylsarcosinate, formamide, formaldehyde (Fluka). Ammonium sulphide solution, glycine, paraformaldehyde (Aldrich Chemical Co. Ltd.). Ultra pure DNA Grade Agaroses, sodium dodecyl sulphate, xylene cyanole, bromophenol blue, ethidium bromide tablets (BIO-RAD). Clozapine (Sandoz). Ondansetron (Glaxo). Sephadex G-25 medium, dextran sulphate sodium salt, terminal deoxynucleotidyl transferase (Pharmacia). Terminal deoxynucleotidyl transferase buffer (Boehringer-Mannheim). [α - 35 S]dATP,

α ³²P-dATP (NEN, Dupont). PGEM marker (Promega) Acetic acid, hydrochloric acid, chloroform (Prolab). HistoClear (National Diagnostics). Ethyl alcohol (Hayman Ltd.). Silver nitrate (Johnson Matthey Materials Technology U.K.). Dulbecco's Modified Eagle's medium, Earle's Modified Eagle's medium, Dulbecco's Modified Eagle's medium nutrient mixture - F12, heat-inactivated foetal calf serum, Hank's Balanced salts solution without calcium and magnesium, Ultrosor G, trypsin solution (2.5%), trypsin-EDTA solution (0.25%), mouse laminin, penicillin-streptomycin solution, horse serum, normal goat serum (Gibco, BRL). Neurone specific enolase, met-enkephalin, glial fibrillary acidic protein (Affiniti). 8-well multichamber slides (Nunc).

All drug compounds used for in vivo experiments were dissolved in 0.9% physiological saline, with the exceptions of haloperidol which was dissolved in 0.2% acetic acid , fluphenazine dihydrochloride, and clozapine which were dissolved in 0.5% dimethylsulphoxide.

All drug compounds used for in vitro experiments were dissolved in DEPC-treated water, with the exceptions of haloperidol which was dissolved in 0.2% acetic acid, fluphenazine dihydrochloride and clozapine which were dissolved in 0.5% dimethylsulphoxide, KT5720, calphostin C, and phorbol-12-myristate-13 acetate which were dissolved in dimethylsulphoxide. The stock drug solutions were then diluted in Hank's Balanced Salts Solution, to obtain the final concentration.

2.12 'In vivo' Experimental Design

2.12.1 Acute Drug Treatment

Four groups of male Wistar rats (180-250g) were treated acutely with vehicle (isotonic saline, n=5), haloperidol (1mg/kg, n=3), fluphenazine (3mg/kg, n=3; 6mg/kg, n=3), clozapine (3mg/kg, n=5). Drugs were delivered by intraperitoneal injection in volumes of 1ml/kg of body weight at staggered 5 minute time intervals. After 30 minutes the animals were killed by stunning and decapitation, the brains rapidly removed and placed onto dry ice. The brains were then processed for the detection of mRNA encoding the transcription factors c-fos, zif/268, jun-B and jun-D using in situ hybridisation histochemistry.

Five groups of male wistar rats (180-250g) were treated for a 24 hour time period with vehicle (physiological saline, n= 5), haloperidol (1mg/kg, n= 3), fluphenazine (3mg/kg, n= 3), clozapine (3mg/kg, n= 6), and methiothepin (10mg/kg, n= 3). Drugs were injected intraperitoneally (1ml/kg of body weight) at staggered time intervals. After 24 hours, the animals were killed by stunning and decapitation and their brains processed for the detection of mRNA encoding a specific region of the preproenkephalin, preproneuropeptide Y, preprosomatostatin, preprotachykinin, RHS2 and BF-1 peptides, using in situ hybridisation histochemistry.

2.12.2 Chronic Drug Treatment

Three groups of male Wistar rats (250-280g) were chronically treated for 5 days with vehicle (osmotic mini pump containing vehicle, n= 4), clozapine (3mg/kg, n= 4), and fluphenazine (3mg/kg, n= 4). The drugs were administered

via osmotic mini pumps (Alzet). Freshly prepared drug solutions were carefully injected into the osmotic mini pumps according to the manufacturers instructions. The animals were anaesthetised with equethesin (4ml/kg) and the osmotic mini pumps inserted subcutaneously into the hind limb region. Drug release was moderated by the pump to 1 μ l/ hour. Animals were weighed on the second, third and fifth day after insertion of the osmotic mini pumps and checked regularly for any symptomatic effects of the drugs. After five days the animals were killed by stunning and decapitation, and their brains rapidly processed for the detection of mRNAs encoding preprosomatostatin, and also the dopaminergic D1 receptor and the serotonergic 5-HT₆ receptor.

2.12.3 Time Course Experiments

Male Wistar rats (180-250g) were treated with vehicle (n= 2 per time point) or either fluphenazine (6mg/kg) or clozapine (3mg/kg) for the following time periods: 15 minutes (n= 3), 30 minutes (n= 3), 45 minutes (n= 3), 90 minutes (n= 3) and 3 hours (n= 3). Both drugs were delivered by intraperitoneal injection in volumes of 1ml/kg. After the specified times, the animals were killed by stunning and decapitation, and their brains rapidly frozen on dry ice. The brains were then processed for the detection of mRNA encoding the transcription factors c-fos, jun-B, and zif/268, and the late response genes proenkephalin and preprosomatostatin.

2.13 'In vitro' Experimental Design

All striatal neurone cultures were treated with drugs for 45 minutes at 37°C. The drugs were delivered directly to the bathing medium in 20µl volumes, making the assumption that the total volume bathing the cells was 0.4ml. After 45 minutes the cells were fixed in a 4% buffered paraformaldehyde solution, rinsed in 1 X PBS, dehydrated in ethanol and processed for in situ hybridisation as previously described.

2.13.1 Time Course Experiment In Vitro

Striatal neurone cultures were treated with haloperidol (final bath concentration 2µM) administered to each well of an 8-well multichamber slide for the following time intervals: 90 minutes (n= 3), 60 minutes (n= 3), 45 minutes (n= 3), 30 minutes (n= 3), 20 minutes (n= 3), 10 minutes (n= 3), 5 minutes (n= 3), and at time zero (n= 3). The slides were then fixed and processed for the detection of mRNA encoding the transcription factors zif/268, jun-B and c-fos.

2.13.2 Treatment with Agonist Compounds

Striatal neurone cultures were treated with vehicle (Hank's Balanced Salts Solution), the dopaminergic agonists SKF 38393 (final bath concentration 1nM-5µM), dopamine (final bath concentration 1nM-5µM), and quinpirole (final bath concentration 1µM), the serotonergic agonists 8-hydroxy-DPAT (final bath concentration 1nM-5µM), mCPP (final bath concentration 1nM-5µM), ±-DOI (final bath concentration 1nM-5µM) and the protein kinase C

stimulating drug phorbol-12-myristate-13-acetate (final bath concentration 10^{-7} M) at 37°C. After 45 minutes the cells were fixed and processed for the detection of c-fos and zif/268 mRNA using in situ hybridisation.

2.13.3 Treatment with Antagonist Compounds

Striatal neurone cultures were pretreated for 30 minutes with the dopaminergic antagonist compounds SCH 23390 (final bath concentration 1 μ M), eticlopride (final bath concentration 1 μ M), the serotonergic antagonist compounds cyproheptadine (final bath concentration 500nM), ondansetron (final bath concentration 200nM), methiothepin (final bath concentration 200nM), mesulergine (final bath concentration 30, 100nM), clozapine (final bath concentration 100nM), and spiperone (final bath concentration 30,100nM) prior to the addition of SKF 38393 (final bath concentration 1nM-5 μ M) or 5-hydroxytryptamine (final bath concentration 1nM-5 μ M) respectively. Similarly cultures were pretreated for 30 minutes with the protein kinase A inhibitor, KT5720 (final bath concentration 2 μ M) and the protein kinase C inhibitor, calphostin C (final bath concentration 1 μ M) prior to the addition of SKF 38393 (final bath concentration 1 μ M). Cultures treated with calphostin C were left under conventional room lighting for the first five minutes of the incubation period in order to activate this compound (Bruns et al., 1991). After 45 minutes the cells were fixed and processed for the detection of c-fos and zif/268 mRNA using in situ hybridisation.

2.14 Dopamine Receptor Antisense Experiment

Striatal neurone cultures were treated for 2 days at 37°C with 15-mer antisense oligonucleotides complementary to the D1 dopamine receptor and to the D5 dopamine receptor. The antisense oligonucleotides were delivered in a volume of 0.5µl, such that the final bath concentration was 1µM. As an appropriate control for specificity of effects of these oligonucleotides, cultures were also treated with a D1 sense oligonucleotide, identified as having the same sequence as a region of the D1 receptor mRNA, administered at a concentration of 1µM. After 2 days, the cultures were acutely treated with SKF 38393 (1nM-5µM) for 45 minutes, at 37°C. Slides were fixed and then processed for the detection of both c-fos and zif/268 mRNA using in situ hybridisation.

2.15 Image Analysis and Statistics Used

To quantitate any drug induced changes in the levels of mRNA for the transcription factors c-fos, zif/268, jun-B, jun-D, RHS2, BF-1, for the neuropeptides preproenkephalin, preprotachykinin, preprosomatostatin, proneuropeptide Y, and for the 5-HT6, and dopamine D1 receptors, initially densitometry of film images was done using the MCID SCIR image analysis system. Bilateral measurements of mean relative optical density (R.O.D.) were taken from brain sections on autoradiographic film, in the regions of the striatum and nucleus accumbens from two sections per animal. For cell culture analysis, four measurements of mean R.O.D. were taken from each well of an 8-well multichamber slide exposed to film. For a higher cellular resolution quantification the density of silver grains were measured after dipping slides in liquid photographic emulsion using a Macintosh image analysis system ("image

v1.52" software [NIH]). For a single culture well, a total of fifteen measurements of mean number of pixels occupied by silver grains per unit area, were taken from three fields of view.

Following this, analysis of variance ANOVA, (one-way unstacked or two-way stacked) was performed on the data for each drug treatment (Minitab). The Dunnett's post-hoc test was then used to assess statistical significance of any drug induced changes between individual groups in all experiments unless stated otherwise. However, for the in vivo experimental work on immediate early genes it was necessary to use the more stringent statistical test, the Mann-Whitney U test. Two-way ANOVA, general linear model (Minitab), was used to analyse in vitro drug dose response curves in the presence and absence of antagonist compounds.

Chapter 3

Regulation of Late Response Gene Expression in vivo

INTRODUCTION

3.0 Neuroleptic Modulation of Late Response Genes In Vivo

It has been known for some time that neuroleptics act by blocking dopamine receptors and are clinically effective in the treatment of a number of diseases which are thought to originate from a dopaminergic dysfunction in the central nervous system. Disease states such as schizophrenia and Parkinson's disease are known to be manifestations arising from a dysfunction of the dopaminergic system within the brain. Blockade of dopamine receptors in the striatum by neuroleptic drugs such as haloperidol have since been shown to cause an elevation in the levels of preproenkephalin mRNA and its corresponding protein product (Hong *et al.*, 1978; Sabol *et al.*, 1983; Tang *et al.*, 1983; Angulo *et al.*, 1986; Sivam *et al.*, 1986; Romano *et al.*, 1987; Morris *et al.*, 1988). Conversely, neuroleptic drug treatment has been shown to reduce the levels of the brain neuropeptide substance P in the striatum (Bannon *et al.*, 1986).

However, most of this research has been performed under conditions of chronic drug administration, probably due to the fact that neuroleptic drug treatment is clinically effective only after prolonged administration. At present there is virtually no information on the acute effects of neuroleptic drugs on late response genes in the region of the striatum. With the rapid progression of research into immediate early genes and the recent findings that some typical neuroleptic drugs can elevate the levels of certain immediate early genes including c-fos, jun-B and zif/286 in the striatum (Nguyen *et al.*, 1992), the acute effects of neuroleptics may provide a greater understanding of the sequence of events following dopamine receptor blockade.

The atypical neuroleptics such as clozapine, which are known to produce less extrapyramidal side effects (Ayd, 1961; Claghorn *et al.*, 1983),

and are therefore an improved treatment for certain disease states, have been shown to exert different effects on certain late response genes in comparison to typical neuroleptics drugs. For example the typical neuroleptic, haloperidol, is known to decrease the levels of preprosomatostatin mRNA in the striatum and accumbens (Salin et al., 1989). Whereas the atypical neuroleptic, clozapine, has been shown to increase the levels of somatostatin mRNA in the nucleus accumbens (Saller et al., 1989). This differential regulation of gene expression by atypical compared to typical neuroleptics, may contribute to the different clinical profile of drugs such as clozapine. It is known that clozapine is an anomoloy, in that it has a much higher affinity for the dopamine D 4 receptor than the typical neuroleptic drugs (Van Tol et al., 1991). Therefore it is of great interest to understand further the complete profile of clozapine in terms of its effect on late response genes and the possible receptor mechanisms involved in the actions of this class of drugs.

RESULTS

Short-term Regulation of Proenkephalin Gene Expression in vivo

3.1 Early time course of Fluphenazine on Proenkephalin mRNA expression

Following intraperitoneal administration of fluphenazine (6mg/kg) to male wistar rats (n= 3), the levels of proenkephalin mRNA were not significantly changed 15, 30, 45, 90, and 180 minutes after drug administration, compared to saline treated control animals (n= 2 per treatment group) injected at the corresponding time points. It can be seen that the mRNA levels for the enkephalin gene in the rostral striatum [Figure 3.1.1] were not significantly affected by fluphenazine (6mg/kg) at time periods up to three hours. A similar lack of effect was observed in the caudal striatum after treatment with fluphenazine (6mg/kg) compared to saline treated control animals [Figure 3.1.2].

3.2 Effect of acute Antipsychotic drug treatment on Proenkephalin mRNA levels in the caudal striatum 24 hours later

Chronic treatment with neuroleptics is well known to cause an increase in the levels of the enkephalin gene in the region of the striatum (Sabol *et al.*, 1983; Tang *et al.*, 1983; Angulo *et al.*, 1986; Sivam *et al.*, 1986; Romano *et al.*, 1987; Morris *et al.*, 1988). Therefore we were interested in the acute effects of neuroleptic drug treatments on the expression of the enkephalin gene after 24 hours.

Figure 3.1.1 : Effect of fluphenazine on preproenkephalin mRNA levels in the rat rostral striatum. Fluphenazine (6mg/kg, n= 3, closed triangles) or vehicle (0.9% saline, n= 2, open triangles) were injected i.p. and the animals killed after the specified time intervals i.e.- 15, 30, 45, 90, and 180 minutes, and their brains processed for in situ hybridisation. Results, obtained from film autoradiographs, are expressed as mean relative optical density \pm s.e.m.

Figure 3.1.2 : Effect of fluphenazine (6mg/kg) on preproenkephalin mRNA levels in rat caudal striatum. Fluphenazine (6mg/kg; n= 3, closed triangles) or vehicle (0.9%; n= 2, open triangles) were injected i.p., the animals killed after the specified time intervals i.e- 15, 30, 45, 90 and 180 minutes, and their brains processed for in situ hybridisation. Results, obtained from film autoradiographs, are expressed as mean relative optical density \pm s.e.m.

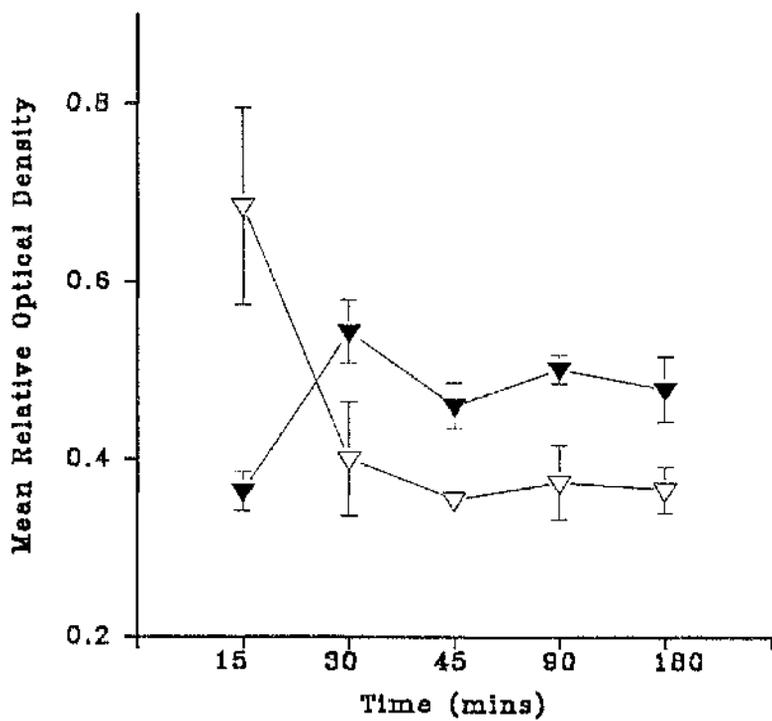
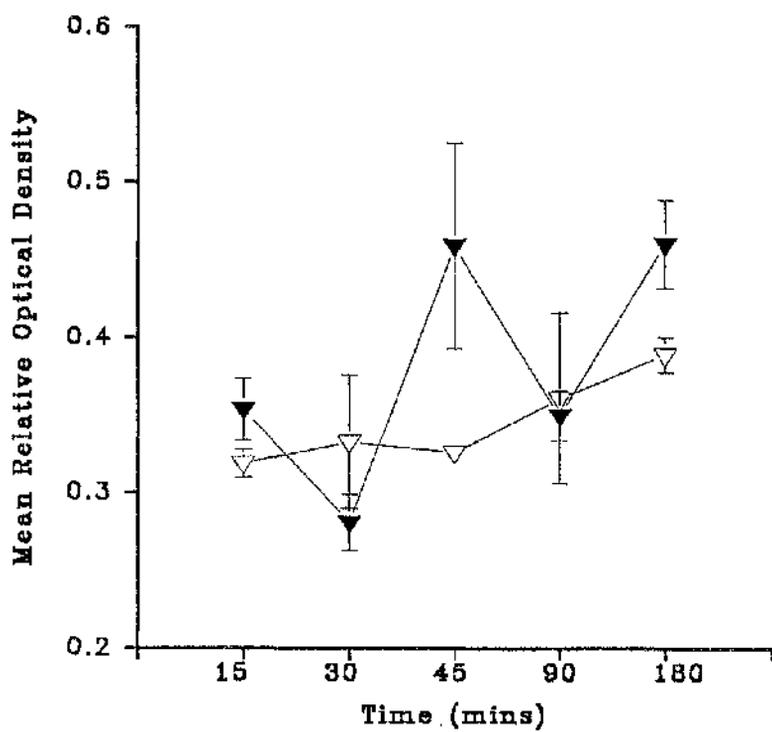
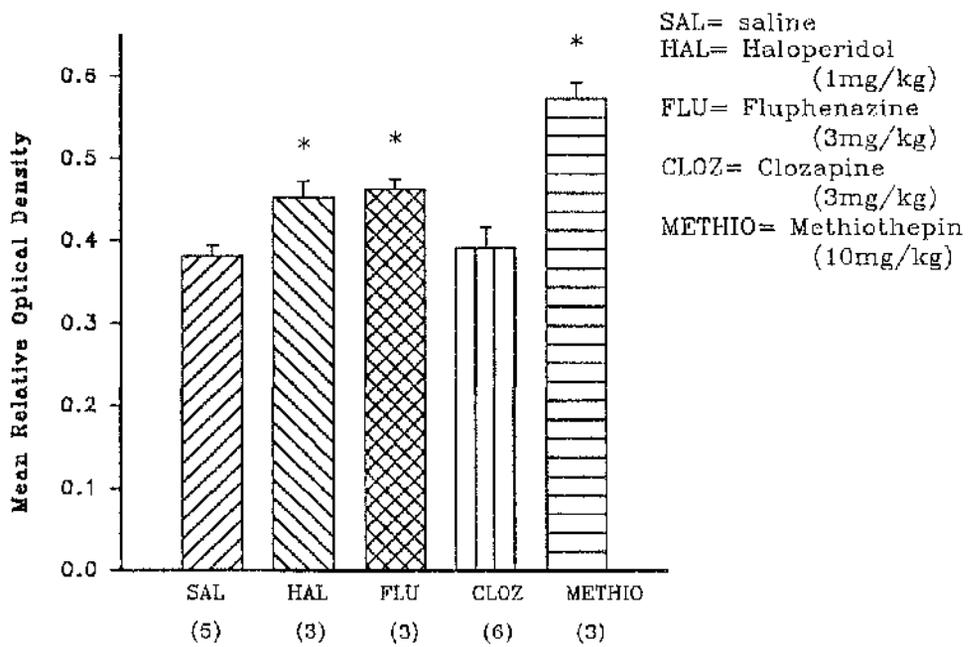


Figure 3.2.1 : Effect of acute neuroleptic drug treatment on the expression of proenkephalin mRNA in the rat caudal striatum. Animals were injected i.p. with vehicle (0.9% saline), haloperidol (1mg/kg), fluphenazine (3mg/kg), clozapine (3mg/kg), or with methiothepin (10mg/kg). After 24 hours the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m.; the number of animals per group is shown in parentheses. Statistical significance was determined by one way ANOVA followed by post-hoc Dunnett's test. $p < 0.05^*$ relative to saline group.



In saline treated animals a moderate hybridisation signal for preproenkephalin was observed in the caudal striatum. Typical neuroleptic drug treatments, together with methiothepin were demonstrated to significantly increase the levels of proenkephalin mRNA in the caudal striatum [$F(4, 15) = 14.40, p = 0.001$]. Following intraperitoneal administration of the typical neuroleptic haloperidol (1mg/kg, $n = 3$) a significant induction in the levels of preproenkephalin mRNA were seen ($p < 0.05$) [Figure 3.2.1]. A similar increase was observed following treatment with fluphenazine (3mg/kg, $n = 3$), and this induction was shown to be significantly different from saline treated control rats ($n = 5$), such that $p < 0.05$ [Figure 3.2.1]. The atypical neuroleptic clozapine ($n = 6$) had no effect on the expression of the enkephalin gene. However, following administration of the non-selective 5-HT_{1/2} receptor antagonist, methiothepin (10mg/kg, $n = 3$), also having a relatively high affinity for the dopamine D₂ receptor, the levels of preproenkephalin were significantly induced such that $p < 0.05$ [Figure 3.2.1].

Short-term regulation of preprotachykinin gene expression in the rat striatum

3.3 Effect of acute drug treatment on the expression of preprotachykinin mRNA levels in the rat striatum 24 hours later

It has previously been reported that chronic treatment with typical neuroleptics can decrease the levels of preprotachykinin mRNA in the rat striatum (Bannon *et al.*, 1986). In this study we have investigated the acute effects of typical and atypical neuroleptic drugs on preprotachykinin gene expression.

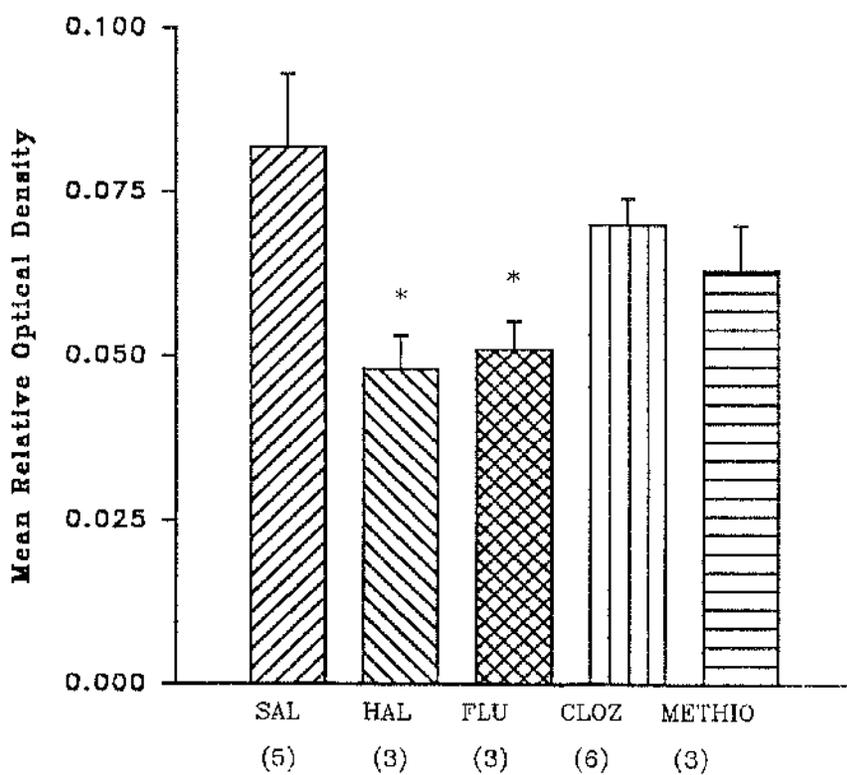
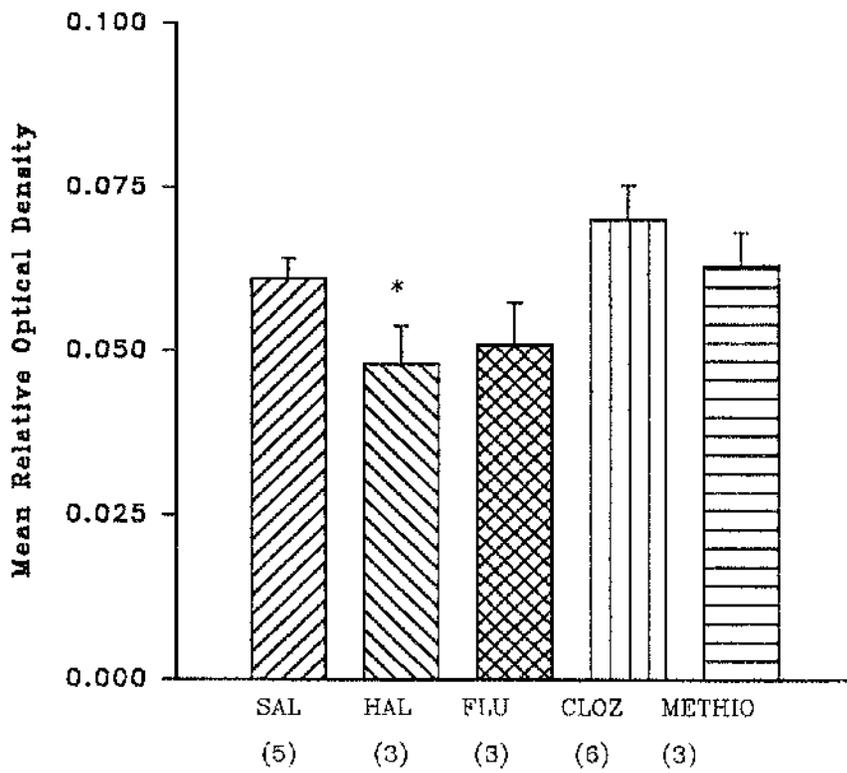
Following intraperitoneal injections of saline (n= 5), or haloperidol (1mg/kg; n= 3), or fluphenazine (3mg/kg; n= 3), or clozapine (3mg/kg; n= 6), or methiothepin (10mg/kg; n= 3), the levels of preprotachykinin mRNA were measured in the rat striatum 24 hours after drug administration.

Analysis of variance revealed that the typical neuroleptics had an effect on the level of preprotachykinin mRNA in both the rostral [$F(4, 15) = 6.92$, $p < 0.001$] and caudal [$F(4, 15) = 3.82$, $p < 0.05$] striatum. Haloperidol (1mg/kg, n= 3) caused a reduction in the levels of preprotachykinin mRNA in the rostral striatum, compared to saline treated control animals. This decrease was shown to be significantly different from the levels expressed in saline treated controls ($p < 0.05$) [Figure 3.3.1]. Fluphenazine (3mg/kg, n= 3) had a tendency to decrease preprotachykinin mRNA levels in the rostral striatum, although this decrease was not found to be significant. The atypical neuroleptic clozapine (3mg/kg, n= 6) or methiothepin (10mg/kg, n= 3) had no effect on preprotachykinin mRNA levels after 24 hours in the rostral striatum. [Figure 3.3.1]

A similar significant reduction in the levels of preprotachykinin mRNA compared to saline treated controls, was observed in the caudal striatum 24 hours after treatment with haloperidol (1mg/kg, n= 3) ($p < 0.05$) [Figure 3.3.2]. Fluphenazine (3mg/kg, n= 3), also caused a significant decrease in preprotachykinin mRNA levels in the caudal striatum, such that $p < 0.05$ [Figure 3.3.2]. However, clozapine (3mg/kg, n= 6) and methiothepin (10mg/kg, n= 3) had no effects on the levels of preprotachykinin mRNA expressed in the rat caudal striatum. [Figure 3.3.2].

Figure 3.3.1 : Effect of acute neuroleptic drug treatment on the expression of preprotachykinin mRNA in the rat rostral striatum. Animals were injected i.p. with vehicle (0.9% saline), or haloperidol (1mg/kg), or fluphenazine (3mg/kg), or clozapine (3mg/kg) or with methiothepin (10mg/kg). After 24 hours the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m.; the number of animals per group is shown in parentheses. Statistical significance was determined by one-way ANOVA, followed by post-hoc Dunnett's test. $p < 0.05^*$ relative to saline group.

Figure 3.3.2 : Effect of acute neuroleptic drug treatment on the expression of preprotachykinin mRNA in the rat caudal striatum. Animals were injected i.p. with vehicle (0.9% saline), or haloperidol (1mg/kg), or fluphenazine (3mg/kg), or clozapine (3mg/kg) or with methiothepin (10mg/kg). After 24 hours the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m.; the number of animals per group is shown in parentheses. Statistical significance was determined by one-way ANOVA, followed by post-hoc Dunnett's test. $p < 0.05^*$ relative to saline group.



Short-term regulation of preprosomatostatin gene expression in the rat striatum

3.4 Effect of acute drug treatment on the levels of preprosomatostatin mRNA in the caudal striatum and in the nucleus accumbens 24 hours later

Previous studies have suggested that high doses of clozapine can cause an induction of preprosomatostatin mRNA in the rat nucleus accumbens (Salin *et al.*, 1989). Therefore in this study we have investigated the effect of acute (24 hour) treatment with low doses of clozapine, together with two typical neuroleptics haloperidol and fluphenazine, on the expression of preprosomatostatin mRNA.

The levels of preprosomatostatin mRNA in the caudal striatum were not changed by the administration of haloperidol (1mg/kg, n= 3) or fluphenazine (3mg/kg, n= 3) compared to saline treated control animals (n= 5). However, following treatment with clozapine (3mg/kg, n= 6) or methiothepin (10mg/kg, n= 3), an increase in the level of preprosomatostatin mRNA was observed in the region of the caudal striatum. This induction was shown to be significantly different from saline treated control animals [$F(4, 15) = 9.46, p = 0.01, p < 0.05$] [Figures 3.4.1]. The effect of clozapine (3mg/kg, n= 6) on the expression of preprosomatostatin in the caudal striatum was significantly different from control animals, such that $p < 0.05$ [Figure 3.4.1]. This acute effect of clozapine can also be seen in figure 3.4.2, as an increase in the number of silver grains over cell bodies in the striatum expressing the somatostatin gene. Treatment with methiothepin (10mg/kg, n= 3), caused a greater increase in the levels of preprosomatostatin mRNA in the rat caudal striatum than did

clozapine, and this increase was significantly different from control animals ($p < 0.01$) [Figure 3.4.1].

A similar pattern of induction was observed in the rat nucleus accumbens. Analysis of variance revealed that the drugs under investigation caused an induction in somatostatin gene expression after 24 hours [$F(2, 9) = 7.97$, $p = 0.006$, $p < 0.01$]. Clozapine (3mg/kg, $n = 6$) caused an increase in the expression of preprosomatostatin mRNA in the core region of the accumbens. This increase was significantly different from the levels expressed in saline treated rats ($p < 0.01$). [Table 1]. Similarly administration of methiothepin (10mg/kg, $n = 3$) resulted in an induction of preprosomatostatin mRNA in the core region 24 hours after administration ($p < 0.01$) [Table 1].

<u>Drug Treatment</u>	<u>Nucleus Accumbens</u>
Saline	11.10 \pm 0.82
Clozapine	22.63 \pm 2.60 **
Methiothepin	18.15 \pm 1.50 **

Table 1: The expression of preprosomatostatin mRNA in the rat nucleus accumbens following a 24 hour treatment with vehicle (physiological saline), clozapine (3mg/kg, $n = 6$), or with methiothepin (10mg/kg, $n = 3$). Results, obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.c.m. Significance was determined by one-way ANOVA, followed by *post-hoc* Dunetts t-test. $p < 0.01$ **

Figure 3.4.1 : Effect of acute neuroleptic drug treatment on the expression of preprosomatostatin mRNA in the rat caudal striatum. Animals were injected i.p. with vehicle (0.9% saline), haloperidol (1mg/kg), fluphenazine (3mg/kg), clozapine (3mg/kg), or with methiothepin (10mg/kg). After 24 hours the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m.; the number of animals per group is shown in parentheses. Statistical significance was determined by one-way ANOVA, followed by post-hoc Dunnett's test. $p < 0.05$ * $p < 0.01$ ** relative to saline group.

Figure 3.4.2 : Effect of acute treatment with clozapine on the expression of preprosomatostatin mRNA in the rat striatum. Photomicrographs show the distribution of silver grains over cell bodies in the caudal striatum. On the left, the animal was injected i.p with vehicle (0.9% saline) (A) and on the right, with clozapine (3mg/kg) (B) Animals were killed after 24 hours, following which their brains were processed for in situ hybridisation. Scale bar represents 25 μ m.

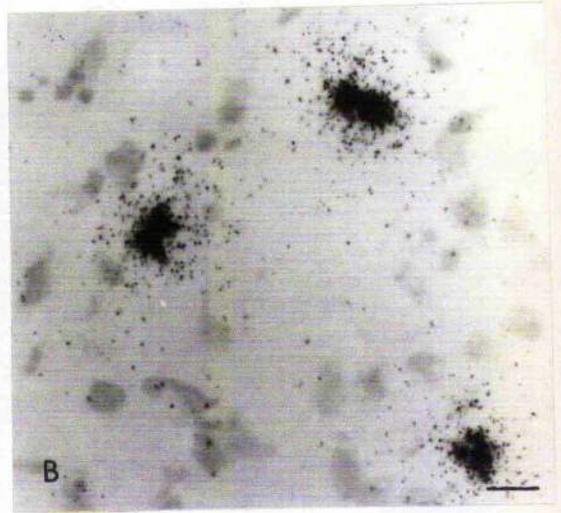
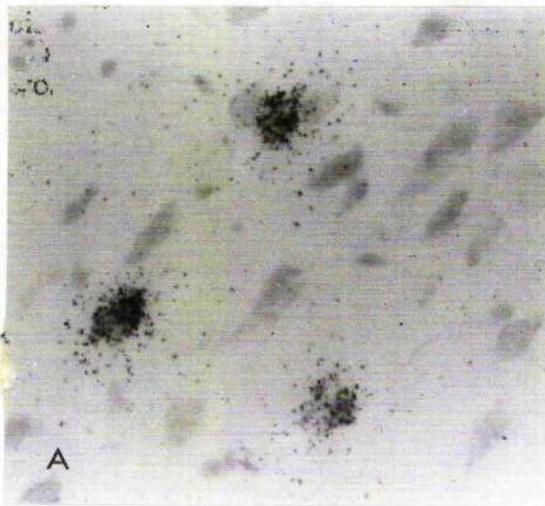
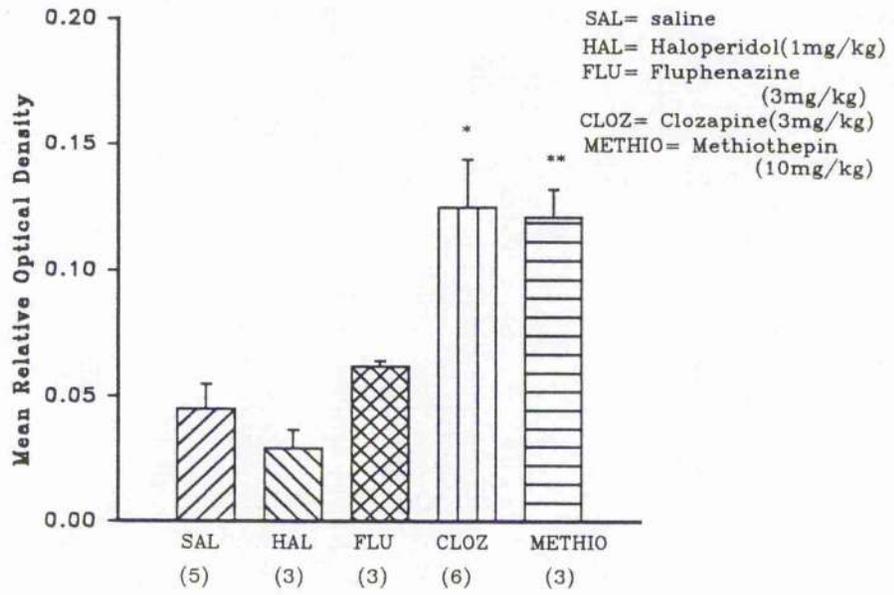
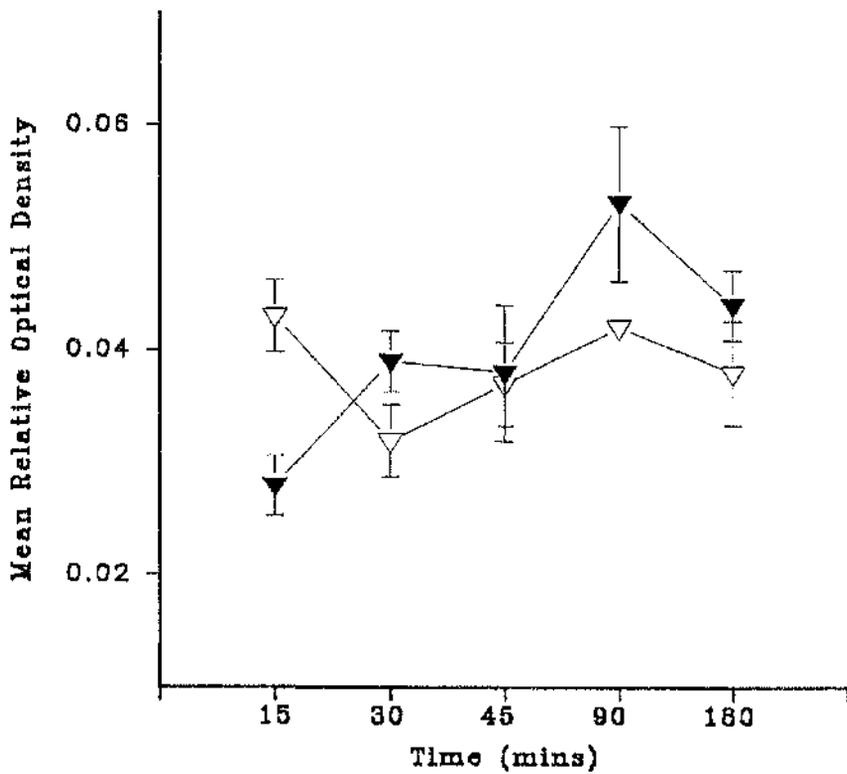
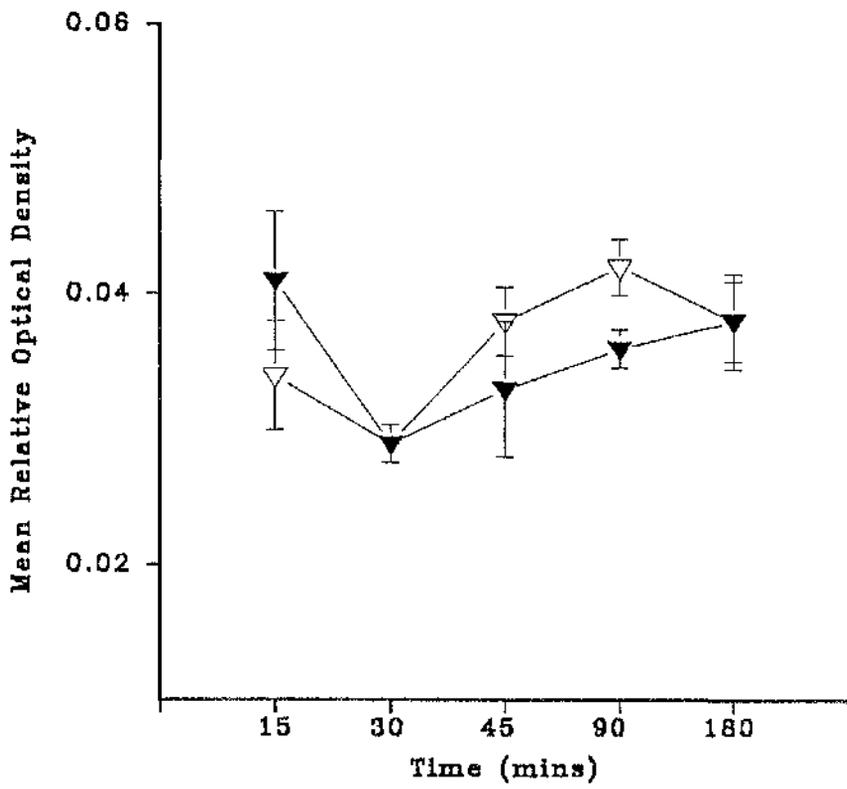


Figure 3.5.1 : Time course effect of clozapine on somatostatin mRNA levels in the rat rostral striatum. Clozapine (3mg/kg; n= 3, closed triangles) or vehicle (0.9% saline; n= 2, open triangles) were injected i.p., the animals killed after the specified time intervals i.e.- 15, 30, 45, 90, and 180 minutes, and their brains processed for in situ hybridisation. Results, obtained from film autoradiographs, are expressed as mean relative optical density \pm s.e.m.

Figure 3.5.2 : Time course effect of clozapine on somatostatin mRNA levels in the rat caudal striatum. Clozapine (3mg/kg; n= 3, closed triangles) or vehicle (0.9% saline; n= 2, open triangles) were injected i.p., the animals were killed after the specified time intervals i.e.- 15, 30, 45, 90, and 180 minutes, and their brains processed for in situ hybridisation. Results, obtained from film autoradiographs, are expressed as mean relative optical density \pm s.e.m.



3.5 The early time course of the clozapine effect on the expression of preprosomatostatin mRNA in the rat striatum

The levels of preprosomatostatin mRNA following treatment with saline (n= 2) or clozapine (3mg/kg; n= 3) were not significantly different after the following time points: 15, 30, 45, 90, 180 minutes, in either the rostral striatum [Figure 3.5.1] or caudal striatum [Figure 3.5.2] in the rat.

Long-term regulation of somatostatin gene expression in the rat striatum and nucleus accumbens

3.6 Effect of low doses of clozapine given chronically, on the expression of preprosomatostatin mRNA in the rat striatum and olfactory tubercle

Animals were treated with either saline (n= 4) or clozapine (6mg/kg /day, n= 4), administered via an osmotic mini pump for 5 consecutive days, following which the levels of preprosomatostatin mRNA were measured by in situ hybridisation. Moderate levels of preprosomatostatin mRNA were expressed in saline treated animals, in a small population of cells known to express the somatostatin gene in both the rostral and caudal striatum. Following chronic treatment with clozapine (6mg/kg, n= 4), an increase in the level of preprosomatostatin mRNA was observed in both the rostral and caudal striatum. The induction of preprosomatostatin mRNA in the rostral region was shown to be significantly different from the levels expressed in saline treated rats [$F(1, 6) = 7.92, p=0.022, p < 0.05$] [Figure 3.6.1]. A similar, and highly

Figure 3.6.1 : Effect of chronic treatment with clozapine on the expression of preprosomatostatin mRNA in the rat rostral striatum. Clozapine (6mg/kg) or vehicle (0.9% saline) was administered via an osmotic mini pump for 5 days. Following this, the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m. The number of animals per group are shown in parentheses. Significance was determined by one-way ANOVA, followed by post-hoc Dunnett's test. $p < 0.05^*$ relative to saline group.

Figure 3.6.2 : Effect of chronic treatment with clozapine on the expression of preprosomatostatin mRNA in the rat caudal striatum. Clozapine (6mg/kg) or vehicle (0.9% saline) was administered via an osmotic mini pump for 5 days. Following this, the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m. The number of animals per group are shown in parentheses. Significance was determined by one-way ANOVA, followed by post-hoc Dunnett's test. $p < 0.01^{**}$ relative to saline group.

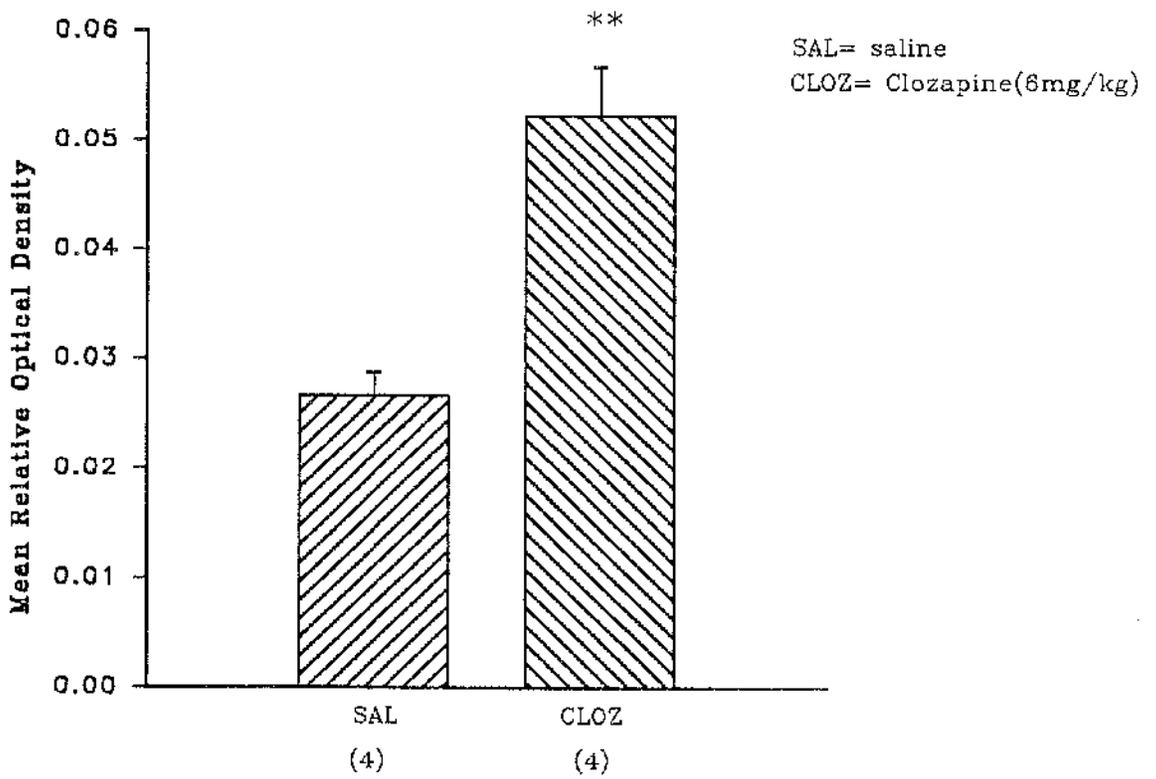
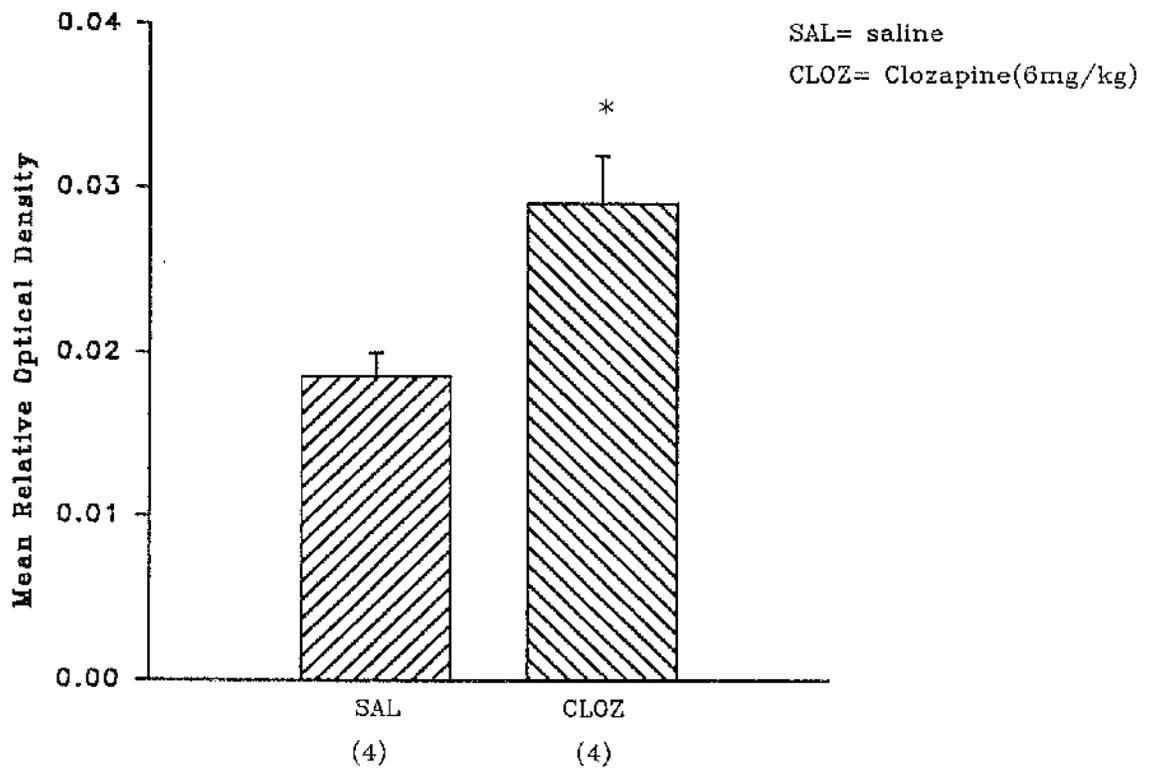
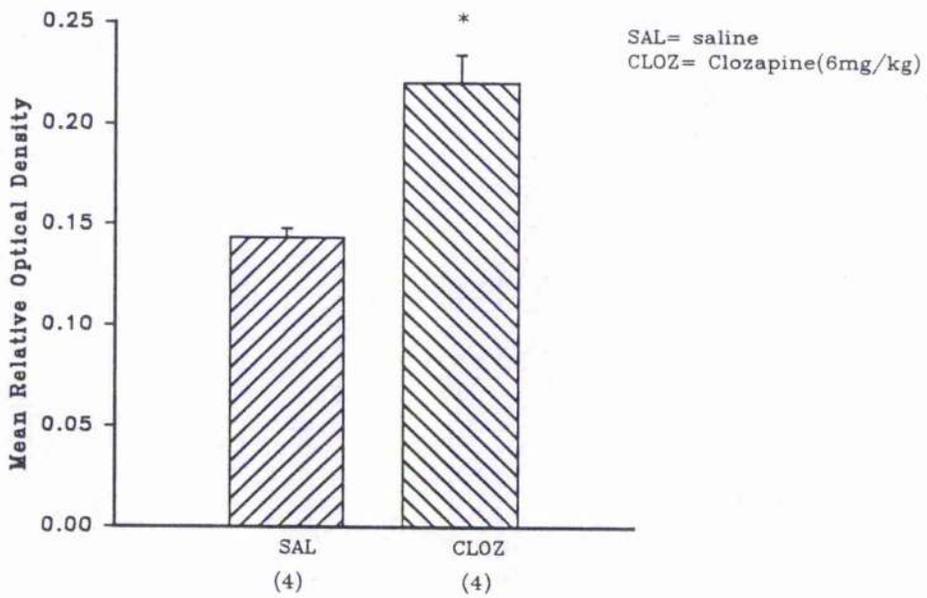
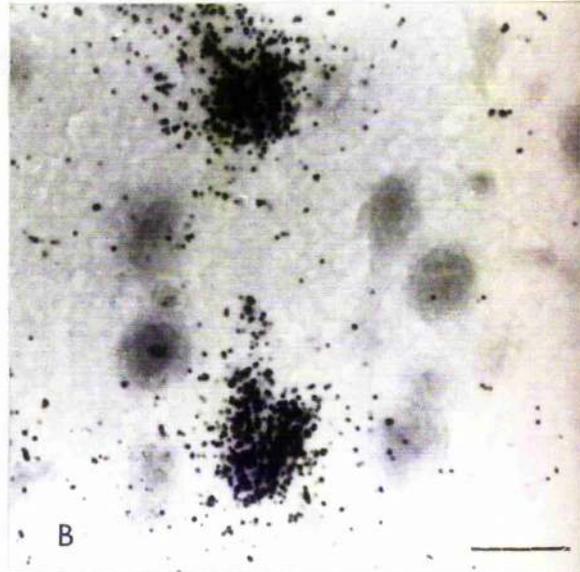
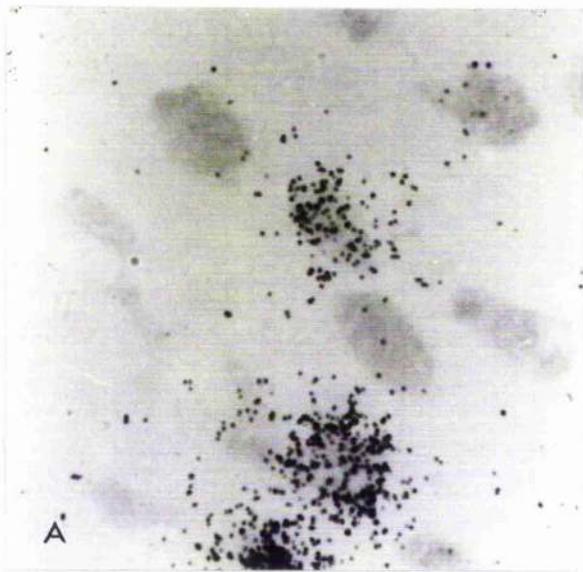


Figure 3.6.3 : Effect of chronic treatment with clozapine on the expression of preprosomatostatin mRNA in the caudal striatum of the rat. Photomicrographs show the distribution of silver grains over striatal cell bodies that express the somatostatin gene. On the left, is an animal chronically treated with saline (A), and on the right , an animal chronically treated with clozapine (6mg/kg), via osmotic mini pump for 5 days (B). $p < 0.01^{**}$. Scale bar represents $25\mu\text{m}$.

Figure 3.6.4 : Effect of chronic treatment with clozapine on the expression of preprosomatostatin mRNA in the rat olfactory tubercle. Clozapine (6mg/kg) or vehicle (0.9% saline) were administered via osmotic mini pumps for 5 days. Following this, the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m. The number of animals per group are shown in parentheses. Significance was determined by one-way ANOVA, followed by post-hoc Dunnett's test. $p < 0.05^*$ relative to saline group.



significant increase in the levels of preprosomatostatin mRNA compared to the saline treated controls, was also observed in the caudal regions of the rat striatum [$F(1,6) = 20.24, p = 0.005, p < 0.01$] [Figures 3.6.2, 3.6.3], following low doses of clozapine.

In the olfactory tubercle, a similar induction of the somatostatin gene was observed after chronic treatment with clozapine (6mg/kg, n= 4), and this induction was shown to be significantly different from saline treated control animals [$F(1,6) = 19.28, p = 0.019, p < 0.05$] [Figure 3.6.4].

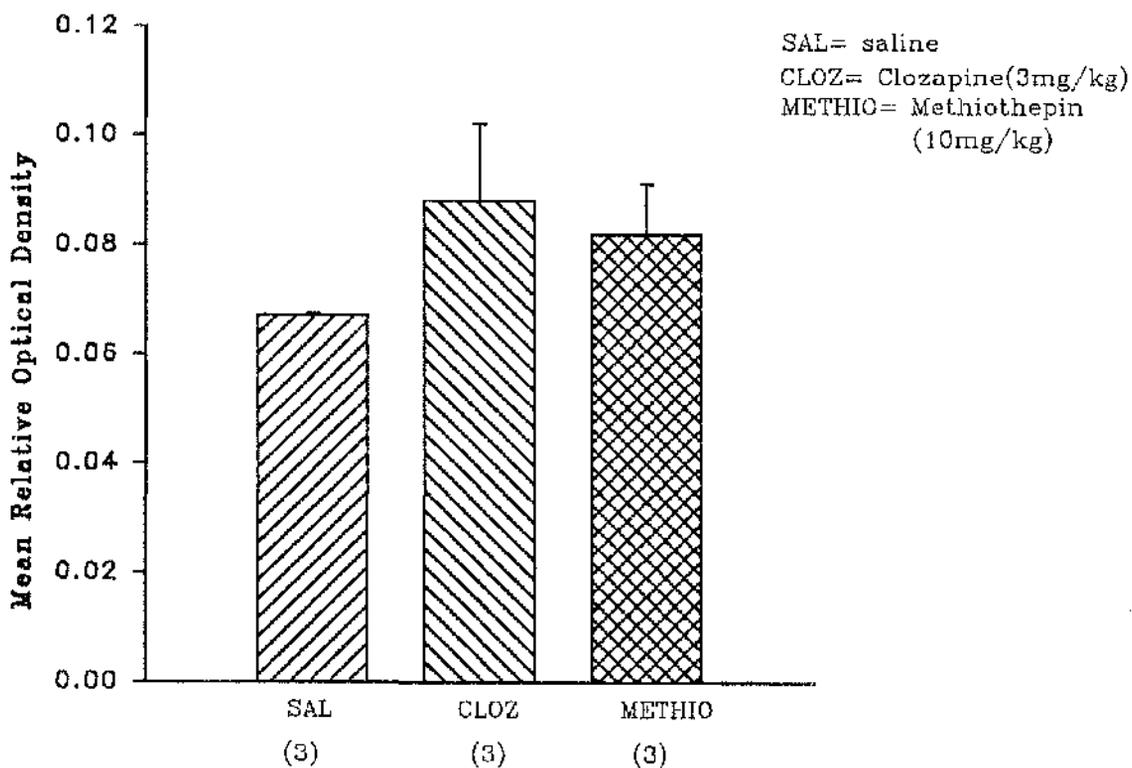
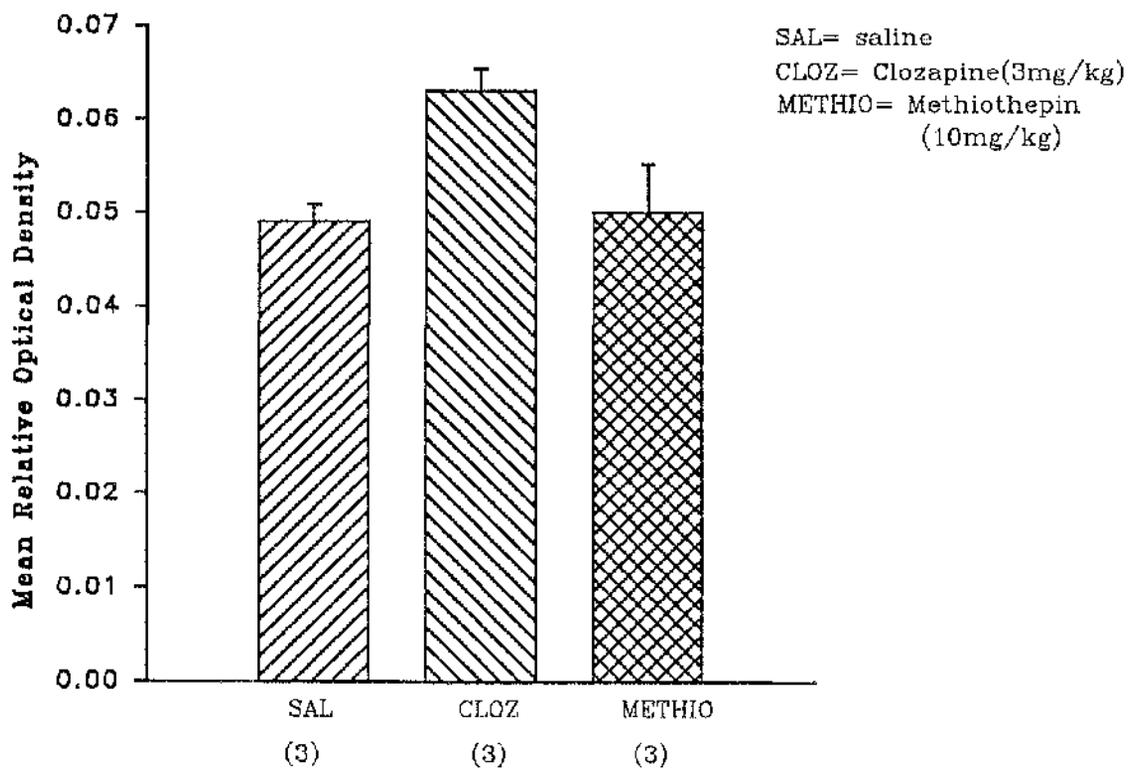
Short-term regulation of proneuropeptide Y gene expression in the rat striatum

3.7 Effect of acute drug treatment on the expression of proneuropeptide Y mRNA levels in the rat striatum 24 hours later.

Following intraperitoneal injection of saline (n= 3), or clozapine (3mg/kg, n= 3), or methiothepin (10mg/kg, n= 3), proneuropeptide Y was measured in the regions of the rostral and caudal striatum 24 hours after drug administration. No significant change in the level of proneuropeptide Y mRNA was seen in either the rostral [Figure 3.7.1] or in the caudal [Figure 3.7.2] regions of the rat striatum, compared to saline treated control animals.

Figure 3.7.1 : Effect of acute drug treatment on the expression of proneuropeptide Y mRNA in the rat rostral striatum. Clozapine (3mg/kg, n= 3) , methiothepin (10mg/kg, n= 3) or vehicle (0.9% saline, n= 3) were injected i.p. After 24 hours, the animals were killed and their brains processed for in situ hybridisation. Results, obtained from film autoradiographs, are expressed as mean relative optical density + s.e.m.

Figure 3.7.2 : Effect of acute drug treatment on the expression of proneuropeptide Y mRNA in the rat caudal striatum. Clozapine (3mg/kg, n= 3), methiothepin (10mg/kg,n= 3) or vehicle (0.9% saline, n= 3) were injected i.p. After 24 hours the animals were killed and their brains processed for in situ hybridisation. Results, obtained from film autoradiographs, are expressed as mean relative optical density + s.e.m.



Long-term effects of neuroleptics on receptors

3.8 Effect of chronic neuroleptic drug administration on the expression of the dopamine D1 receptor mRNA in the rat striatum

The expression of the D1 dopamine receptor in the rat striatum is known to be homogenously distributed throughout the striatum.

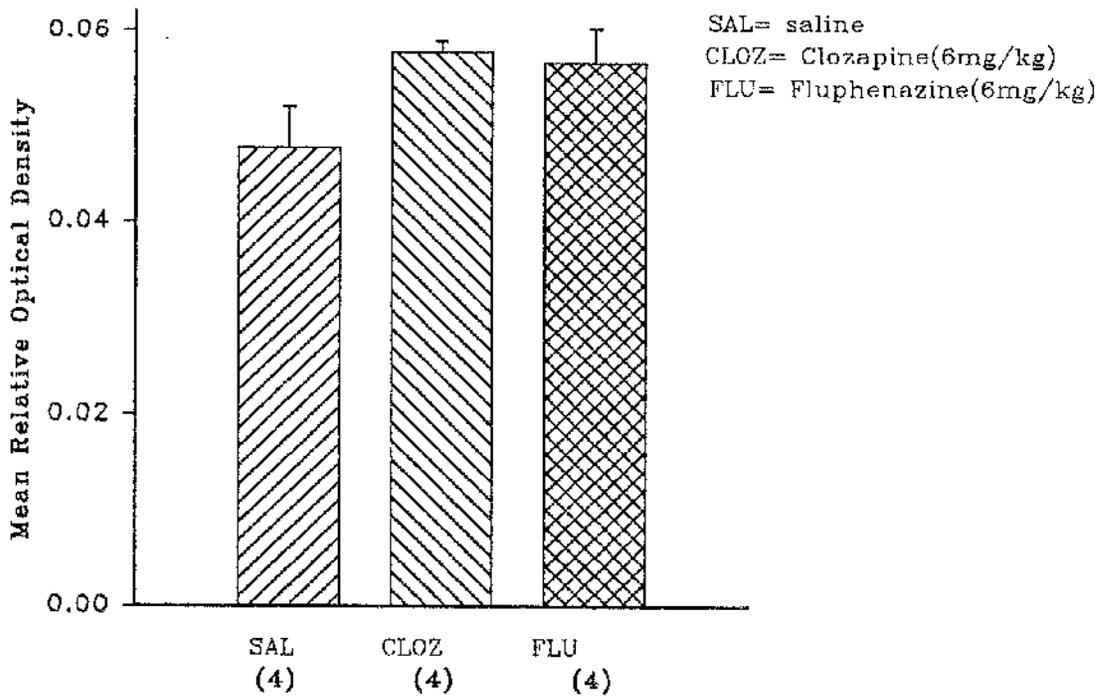
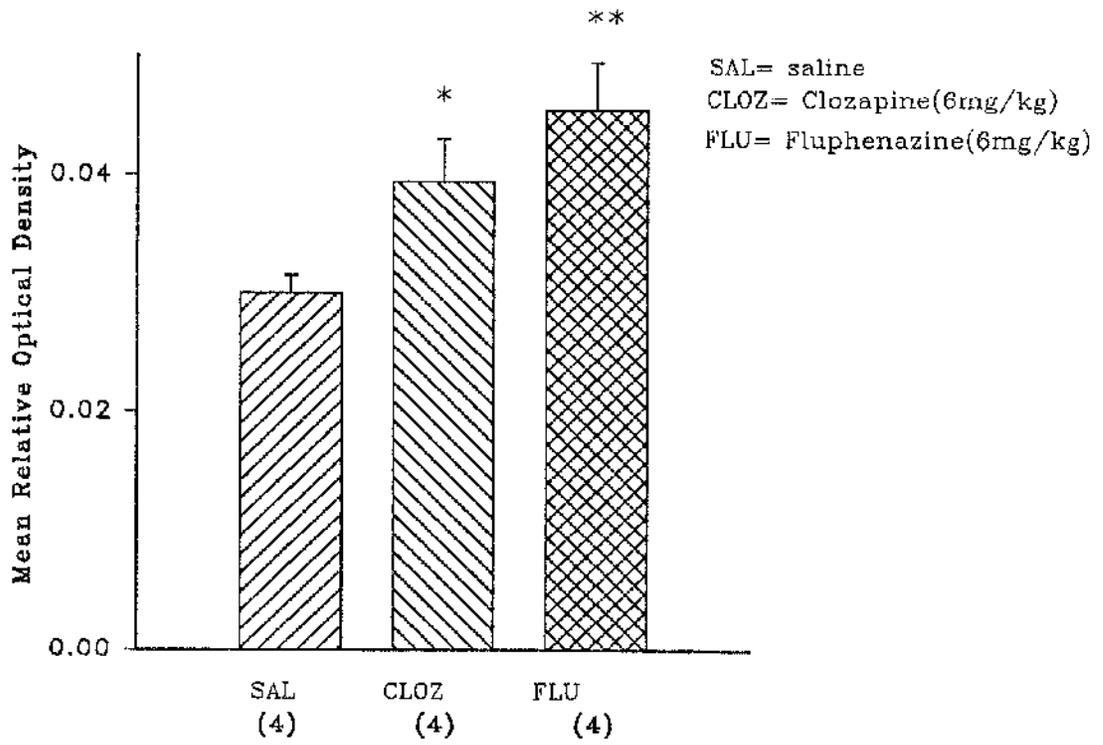
Animals were treated chronically with saline (n= 4), clozapine (6mg/kg, n= 4) or with fluphenazine (6mg/kg, n= 4) for 5 days, following which the levels of D1 receptor mRNA were measured as previously described.

It was found that both clozapine (6mg/kg, n= 4) and fluphenazine (6mg/kg, n= 4) when given chronically could increase the expression of D1 mRNA in the rostral striatum of the rat, $F(2, 9) = 5.84$, $p = 0.01$, $p < 0.01$ [Figure 3.8.1]. The increase in D1 mRNA after clozapine treatment was significantly different from control animals, such that $p < 0.05$. Similarly chronic treatment with fluphenazine (6mg/kg, n= 4) caused an increase in D1 mRNA in the rostral striatum. This increase was greater than the induction due to clozapine, such that $p < 0.01$ [Figure 3.8.1].

In caudal regions, both clozapine (6mg/kg, n= 4) and fluphenazine (6mg/kg, n= 4) had a tendency to increase the levels of D1 mRNA expressed in the striatum. Although statistical analysis revealed no significant differences when compared to the control animals [Figure 3.8.2].

Figure 3.8.1 : Effect of chronic neuroleptic drug treatment on the expression of dopamine D1 receptor mRNA in the rat rostral striatum. Clozapine (6mg/kg), or fluphenazine (6mg/kg), or vehicle (0.9% saline) were administered via osmotic mini pumps for 5 days. Following this, the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m. The number of animals per group are shown in parentheses. Significance was determined by one-way ANOVA, followed post-hoc Dunnett's test. * $p < 0.05$, ** $p < 0.01$ relative to saline group.

Figure 3.8.2 : Effect of chronic neuroleptic drug treatment on the expression of dopamine D1 receptor mRNA in the rat caudal striatum. Clozapine (6mg/kg), or fluphenazine (6mg/kg), or vehicle (0.9% saline) were administered via osmotic mini pumps for 5 days. Following this, the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m. The number of animals per group are shown in parentheses. Significance was determined by one-way ANOVA, followed by post-hoc Dunnett's test.



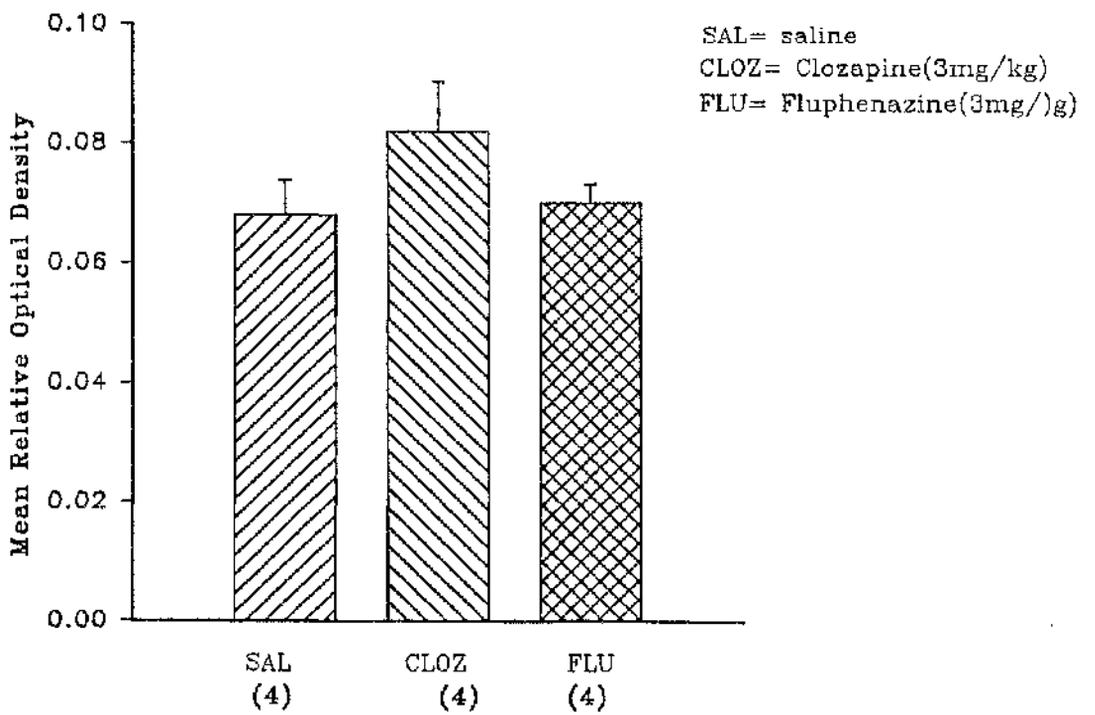
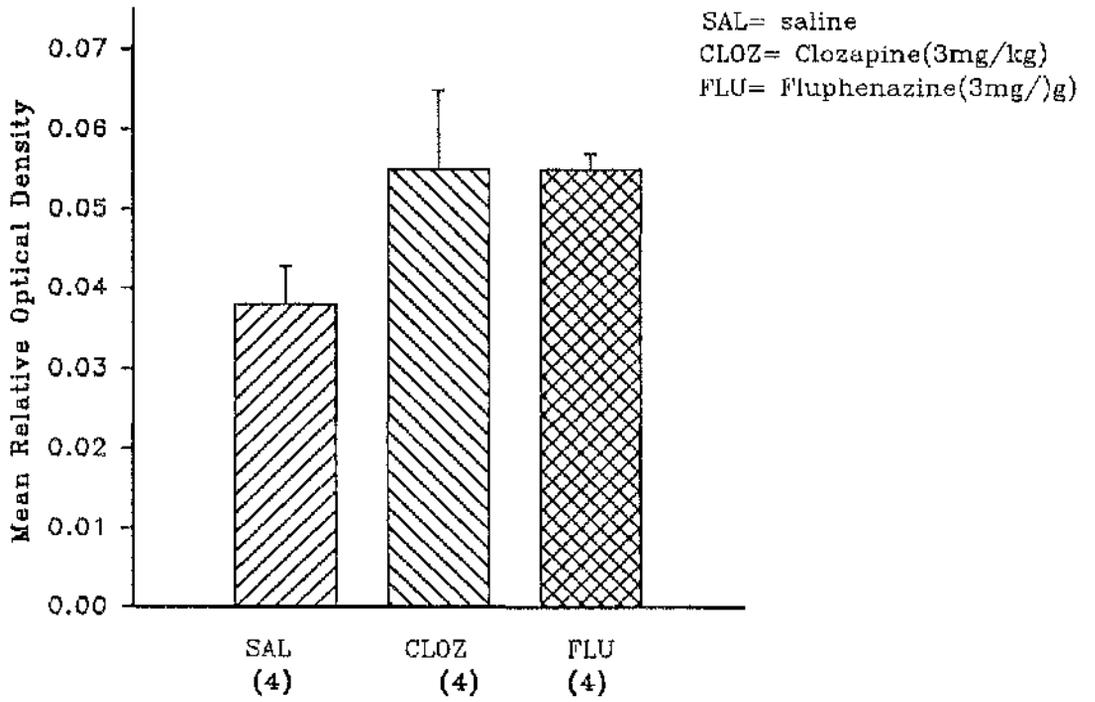
3.9 Effect of chronic neuroleptic drug administration on the levels of 5-HT₆ receptor mRNA in the rat striatum

The recently cloned 5-HT₆ receptor is known to be expressed in the rat striatum and have a high affinity for certain antipsychotic drugs. Therefore we have investigated if chronic administration of the neuroleptic drugs clozapine (6mg/kg, n= 4) or fluphenazine (6mg/kg, n= 4) have any effect on the levels of 5-HT₆ receptor mRNA compared to saline treated control animals.

However, it was found that low doses of clozapine (6mg/kg) or fluphenazine (6mg/kg) had no effect on the levels of 5-HT₆ mRNA in either the rostral [Figure 3.9.1] or the caudal regions in the rat striatum [Figure 3.9.2], such that the levels of 5-HT₆ mRNA were not significantly different from vehicle treated animals.

Figure 3.9.1 : Effect of chronic neuroleptic drug treatment on the expression of serotonergic 5-HT₆ receptor mRNA in the rat rostral striatum. Clozapine (6mg/kg), or fluphenazine (6mg/kg), or vehicle (0.9% saline) were administered via osmotic mini pumps for 5 days. Following this, the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m. The number of animals per group are shown in parentheses.

Figure 3.9.2 : Effect of chronic neuroleptic drug treatment on the expression of serotonergic 5-HT₆ receptor mRNA in the rat caudal striatum. Clozapine (6mg/kg), or fluphenazine (6mg/kg), or vehicle (0.9% saline) were administered via osmotic mini pumps for 5 days. Following this, the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m. The number of animals per group are shown in parentheses.



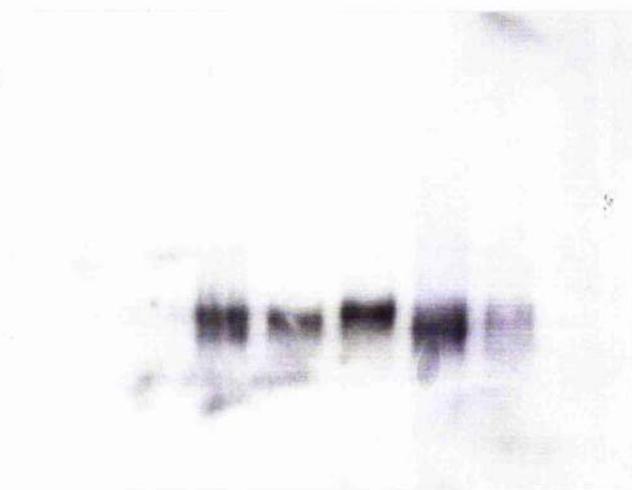
3.10 Northern blot hybridisation analysis of rat RNA

After total poly (A) + RNA (8 μ g/ sample) from each brain region was denatured, electrophoresed on a 1% agarose/ formaldehyde gel and transferred to a nylon membrane, the blot was hybridised with a 32 P- labelled tubulin T26 probe, as described under methods. The tubulin isoform T26 is a cytoskeletal protein present in all brain tissue and therefore was used as an appropriate control for RNA content in brain tissue samples. The blot was exposed to film for forty-eight hours at -80°C. As shown in the autoradiograms of figure 3.20, the rat T26 probe hybridised with a single major band of total poly (A) + RNA [Figure 3.10.1, panel a] of length 1800 nucleotides from all brain regions tested (i.e.- striatum, hippocampus, cortex, olfactory bulb and cerebellum), but not from rat liver. As can be seen from the autoradiogram, the intensity of labelling was very similar in all regions after incubation with a tubulin T26 mRNA probe. In accordance with previous findings, the size of band is consistent with that described for rat tubulin T26 in the brain (Miller et al., 1987). Any other bands appearing on the blot were considered to represent degraded material.

To assess the sensitivity and specificity of the rat preproenkephalin probe, the northern blot of total poly (A) + RNA was hybridised with a 32 P- labelled rat preproenkephalin mRNA probe (45-mer, previously described). After a thirty-six hour exposure to film at -80°C, a single band of length approximately 1500 nucleotides was observed in the region of the striatum, but not in the liver, a tissue essentially devoid of proenkephalin peptides [Figure 3.10.1, panel b]. This size is consistent with that previously determined for bovine and rat striatal preproenkephalin mRNA (Dandekar and Sabol, 1982; Tang et al., 1983). The pattern of expression of preproenkephalin mRNA on

Figure 3.10.1 : Northern blot analysis of (A) tubulin T26 mRNA and (B) preproenkephalin mRNA from various regions of the rat brain. Total poly(A)+ RNA (8 μ g/sample) taken from five brain regions [striatum (ST), hippocampus (H), cortex (CTX), olfactory bulb (OB), and cerebellum (CB) and also from the liver (L)] was denatured and electrophoresed on a 1% agarose, formaldehyde gel, transferred to a nylon membrane and hybridised with either (A) a ³²P-labelled tubulin T26 probe (previously described) or with (B) a ³²P-labelled preproenkephalin probe (previously described) as described under methods.

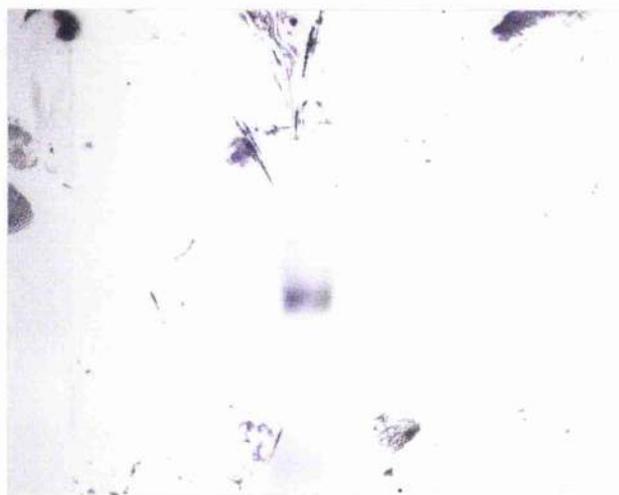
A



1.8 Kb

K CB H ST CTX OB

B



1.5 Kb

the northern blot appeared relatively free from non-specific hybridisation, and showed greatest abundance in the rat striatum, with virtually no signal in other brain regions. However, with a longer time of exposure, it may be expected that a signal representing preproenkephalin mRNA may appear in the region of the cortex and cerebellum, but with a much lower intensity (Yoshikawa *et al.*, 1984).

DISCUSSION

Short-term Regulation of Proenkephalin Gene Expression in vivo

These studies were undertaken to initially determine whether enkephalin gene expression is induced by certain antipsychotic drugs after their acute administration. It is well known information that antipsychotic drugs can induce proenkephalin mRNA levels when given chronically (Sabol *et al.*, 1983; Tang *et al.*, 1983; Angulo *et al.*, 1986; Sivam *et al.*, 1986; Romano *et al.*, 1987; Morris *et al.*, 1988). We have looked at short-term regulation of a number of late response genes, in an attempt to gain further understanding into the functional significance of the class of genes referred to as immediate early genes. By gaining information on the possible receptor mechanism involved in the induction of both late response genes and immediate early genes we have been able to contribute indirect information that may aid in understanding the relationship between late response genes and immediate early gene induction, a theory that currently remains controversial and not well understood.

3.11 Early time course of Fluphenazine on Proenkephalin mRNA expression

The drug fluphenazine was used in this study because of its profile as a typical neuroleptic. Clinical use of typical neuroleptic drugs is frequently associated with the appearance of undesirable side effects including extrapyramidal movement disorders, which further divide into a parkinsonian-like syndrome and tardive dyskinesia, dependent on the length of drug exposure (Baldessarini, 1985; Tarsy and Baldessarini, 1986; Meltzer, 1991). These unwanted extrapyramidal behavioural side effects are thought to be a

consequence of extensive blockade of dopaminergic D2-like receptors in the nigrostriatal (A9) dopamine system (Baldessarini, 1979; Klawans et al., 1980). Fluphenazine is reported to have a marginal selectivity for the D2-like dopamine receptors (Saller et al., 1987), over the D4 dopamine receptor (Van Tol et al., 1991) with a moderate potency at dopamine D1-like receptors (Christensen et al., 1984), which was an important reason for selecting this drug for the following studies.

The results obtained in this study show that acute injection of the typical neuroleptic drug fluphenazine has no effect on the levels of proenkephalin mRNA in the rat striatum up to three hours after the initial injection, when compared to control animals. As in other studies, changes in the mRNA level have been correlated to changes in protein levels (Shiomi et al., 1986; O'Connell et al., 1987; Adler et al., 1988). It may be assumed that fluphenazine does not affect enkephalin protein levels in either the rostral or caudal region of the rat striatum up to three hours after administration. The dosage used would be expected to bind to greater than 50% of D2 dopamine receptors in vivo according to the reported affinity of fluphenazine for D2 dopamine receptors (Van Tol et al., 1991). As the number of animals per group were particularly small, this may have introduced a greater error, and thus masked any small changes in proenkephalin gene expression that may have occurred, at any one time point. In a previous report it has been suggested that a single acute injection of haloperidol can induce proenkephalin mRNA in some animals by fifteen minutes, and in all animals tested by forty-five minutes (Konradi et al., 1993). Similarly the mRNA levels of other peptidergic transmitters, for example, neurotensin, can be affected by acute treatment with dopaminergic D2 receptor antagonists (Augood et al., 1991). The lack of an induction of proenkephalin mRNA after acute treatment with fluphenazine may reflect a slower drug action of this typical neuroleptic. A longer time period

may be necessary for fluphenazine to penetrate the blood-brain barrier and thus have an effect on the levels of proenkephalin mRNA.

3.12 The effect of Antipsychotic drug treatment on Proenkephalin mRNA levels twenty-four hours after administration

In this study the effect of the typical neuroleptics haloperidol and fluphenazine, which have relatively high affinities for D2-like dopamine receptors and moderate potency at D1-like dopamine receptors (Christensen *et al.*, 1984), were compared with the atypical neuroleptic clozapine, which has been demonstrated to show some selectivity for D4 dopamine receptors (Van Tol *et al.*, 1991) and with the 5-HT_{1/2} receptor antagonist drug methiothepin, also known to bind to dopamine receptors, on the expression of proenkephalin mRNA in the rat striatum twenty-four hours after drug administration.

From the results obtained it was observed that the typical neuroleptics haloperidol and fluphenazine increased the expression of proenkephalin mRNA in the rat striatum after twenty four hours. These antipsychotic drugs are known to induce enkephalin mRNA and enkephalin protein after chronic administration (Hong *et al.*, 1978; Sabol *et al.*, 1983; Tang *et al.*, 1983; Angulo *et al.*, 1986; Sivam *et al.*, 1986; Romano *et al.*, 1987; Morris *et al.*, 1988). At present it is generally accepted that the clinical therapeutic effects of these antipsychotic drugs are mediated through blockade of dopamine D2-like receptors (Creese *et al.*, 1976; Seeman *et al.*, 1976; Baldessarini, 1985). The primary evidence for this hypothesis is two-fold. Firstly, most of the available antipsychotic drugs are dopamine D2-like receptor antagonists (Carlsson, 1978; Peroutka and Snyder, 1980; Baldessarini, 1985). Evidence for this comes from an increased dopamine turnover and antagonism of dopamine D2-like

agonist effects, during acute administration, and an increase in D2-like dopamine receptor density following chronic administration (Carlsson, 1978; Muller and Seeman, 1978; Baldessarini, 1985; Wilmot and Szczepanik, 1989). Secondly it has been shown that there is a strong correlation between the relative potencies of antipsychotic agents in competing for dopamine D2 sites *in vitro* and the doses of antipsychotic drugs used in the clinical treatment of psychoses (Peroutka and Snyder, 1980; Seeman, 1980; Cohen and Lipinski, 1981). However, it should be noted that the drugs and doses used would distinguish poorly between D2, D3 and D4 dopamine receptor subtypes. Furthermore, the antipsychotic activity of neuroleptic drugs may, in fact, be related to the interaction of neuroleptics with other transmitter systems in the brain. At clinically effective doses, neuroleptics interact with a high degree of affinity with the α -adrenergic, histamine, muscarinic acetylcholine and serotonin receptors (Peroutka and Snyder, 1980).

A similar induction of proenkephalin mRNA was observed twenty-four hours after treatment with the non-selective 5-HT_{1/2} receptor antagonist methiothepin. Recently, it has been suggested that potent serotonin 5-HT₂ receptor antagonism could alleviate negative symptoms in schizophrenia. Moreover, 5-HT₂ antagonists have been shown to reduce neuroleptic-induced catalepsy in rats (Balsara *et al.*, 1979; Hicks, 1990), to increase neuroleptic-induced dopamine turnover (Saller *et al.*, 1990), to cause a slight, indirect activation of midbrain dopaminergic neurons (Ugedo *et al.*, 1989), restoring phasic rather than tonic activity, and to reduce neuroleptic-induced extrapyramidal side effects (Reyntjens *et al.*, 1986; Bersani *et al.*, 1990). As methiothepin has previously been shown to possess the ability to block dopaminergic receptors in binding studies (Lloyd and Bartholini, 1974; Enjalbert *et al.*, 1978; Nelson *et al.*, 1979; Christensen, 1985), it is likely that the methiothepin-induced increase in proenkephalin mRNA levels in the rat striatum may be due its actions at dopamine D2-like receptors *in vivo*.

Although the possibility still remains that a 5-HT receptor blockade may also be contributing to this effect. Both haloperidol and fluphenazine are known to possess 5-HT₂ receptor binding properties, but with much lower affinity than methiothepin (Meltzer *et al.*, 1989).

The atypical neuroleptic drug clozapine, which is known to be associated with a low incidence of extrapyramidal side effects, a property that as yet has not been attributed to any one physiological effect of the drug, was shown to have no effect on proenkephalin mRNA levels twenty-four hours after administration. Previous studies have revealed that chronic treatment with clozapine does not increase the density of dopamine D₁ or dopamine D₂ receptors (Seeger *et al.*, 1982; Severson *et al.*, 1984; Lee and Tang, 1984; Rupniak *et al.*, 1984; Cohen and Lipinski, 1986). It has been suggested that the atypical antipsychotic agents exert preferential effects on the mesolimbic (A10) dopamine system *in vivo* (Borison *et al.*, 1983; Borison and Diamond, 1983).

These results are the first observations indicating that striatal proenkephalin mRNA is induced after acute treatment, not only with certain typical neuroleptics, but also with the 5-HT_{1/2} receptor antagonist compound, methiothepin, and not with the atypical neuroleptic drug clozapine. It may be suggested that the extrapyramidal side-effects associated with typical neuroleptic drug treatment could be partially attributed to the rapid induction of the enkephalin gene in the striatum shortly after drug administration and the subsequent sustained elevation with chronic use. This would be consistent with evidence that, in rodents with dopamine lesions, the degree of extrapyramidal motor deficit is related to the magnitude of the change in proenkephalin gene expression (Cadet *et al.*, 1992). The induction of proenkephalin mRNA following acute typical neuroleptic drug treatment is most likely due to blockade of D₂ or D₃ dopamine receptor subtypes *in vivo*, rather than the D₄ dopamine subtype, as clozapine, having some selectivity for the D₄ receptor over the D₂ or D₃ receptor subtypes (Van Tol *et al.*, 1991), was ineffective.

Although in contrast to this, a previous study has demonstrated a decrease in proenkephalin gene expression after chronic treatment with high doses of clozapine (Mercugliano and Cheeselet, 1991). It must also be considered that 5-HT receptor blockade may contribute to this effect, since the non-selective 5-HT receptor antagonist methiothepin, together with both the typical neuroleptics fluphenazine and haloperidol, share the ability to block 5-HT₂ receptors.

Short-term Regulation of Preprotachykinin mRNA Expression in vivo

3.13 The effect of Antipsychotic drug treatments on the levels of Preprotachykinin mRNA in the rat striatum twenty-four hours after administration

In this study the typical neuroleptic haloperidol clearly reduced the level of preprotachykinin mRNA in the rat rostral striatum after twenty-four hours, whilst both haloperidol and fluphenazine were shown to decrease preprotachykinin mRNA levels in the rat caudal striatum after acute treatment.

These results are in agreement with previous studies demonstrating that the level of protachykinin mRNA, the mRNA encoding for substance P, is reduced in the rat striatum after chronic treatment with the antipsychotic drug haloperidol (Hong *et al.*, 1983; Bannon *et al.*, 1986). This is the first observation for an acute effect of these neuroleptics on preprotachykinin mRNA levels in vivo. There was no significant change in preprotachykinin mRNA levels in the striatum, twenty-four hours after administration of the atypical neuroleptic, clozapine. This is in agreement with previous reports

indicating that the atypical neuroleptic clozapine does not alter preprotachykinin mRNA in the rat striatum following chronic treatment (Bannon *et al.*, 1987; Shibata *et al.*, 1990). The dose of clozapine used here would be expected to be blocking 80% of α -1 receptors, approximately 65% 5-HT₂ receptors, with relatively little blockade of D₂ dopamine receptors (Schotte *et al.*, 1993). Since clozapine has a decreased propensity for inducing extrapyramidal side-effects, it is conceivable that decreased tachykinin gene expression in the basal ganglia is somehow related to the production of these adverse effects. It is known that chronic administration of most antipsychotic drugs causes a depolarisation inactivation of dopamine neurons innervating the rat striatum. This results in a decreased number of spontaneously active dopamine cells and a diminished release of striatal dopamine (Chiodo, 1988). It is the decrease in dopamine neurotransmission that apparently causes decreased tachykinin biosynthesis similar to that seen after experimental dopamine cell lesioning or in post-mortem material from Parkinson's disease patients (Mauborgne *et al.*, 1983; Tenovno *et al.*, 1984; Young *et al.*, 1986; Clevens and Beal, 1989). In contrast, atypical antipsychotic drugs elicit neither depolarisation inactivation of dopamine cells innervating the striatum nor decreased tachykinin biosynthesis. However, both typical and atypical antipsychotic drugs induce depolarisation inactivation of those dopamine neurons innervating primarily limbic targets (Chiodo, 1988), leading to speculation that decreased limbic dopamine influence is the common mechanism of antipsychotic drug action. Although various typical and atypical antipsychotic agents may differ somewhat in their affinities for various dopamine receptor subtypes, some other property or properties of atypical drugs (such as antiadrenergic or antiserotonergic effects) may be responsible not only for the differing propensities for induction of adverse side effects, but also for additional therapeutic effects (Chiodo, 1988; Meltzer, 1988). The dose of clozapine used in this study was chosen with the knowledge that clozapine

has a particularly high affinity for the D4 dopamine receptor compared to most typical neuroleptics (Van Tol *et al.*, 1991). Doses of clozapine of 10mg/kg and above have been shown to bind to striatal D2 binding sites in rats (Saller *et al.*, 1987; Audinot *et al.*, 1993). It is therefore likely that the lower dose of 3mg/kg used here is preferentially blocking D4 sites in vivo rather than D2 sites. Therefore, it appears unlikely that the D4 receptor is involved in mediating the decrease in preprotachykinin mRNA in the rat striatum, following acute treatment with either haloperidol or fluphenazine.

Similarly, the antipsychotic methiothepin did not alter the levels of preprotachykinin mRNA in this study. As it is most likely that the typical neuroleptic agents studied caused a reduction in preprotachykinin gene expression through a mechanism involving blockade of dopamine receptors, a possible explanation for the lack of effect with methiothepin may be that the affinity of methiothepin for the dopamine D1 and D2 receptors is not sufficiently strong to decrease preprotachykinin gene expression in the striatum of the rat after acute administration. Another possible explanation could be that combined 5-HT antagonism together with dopamine receptor blockade ^{that} is preventing the reduction of preprotachykinin mRNA. Haloperidol and fluphenazine also possess 5-HT receptor antagonistic properties, but perhaps the blockade of dopamine neurotransmission by these typical neuroleptics overrides the ^{properties of} serotonin receptor blockade.

Short-term Regulation of Preprosomatostatin gene expression in vivo

3.14 The effect of Antipsychotic drug treatments on the level of Preprosomatostatin mRNA twenty-four hours after administration

In this study the typical neuroleptic drugs haloperidol and fluphenazine did not significantly alter the levels of preprosomatostatin mRNA in the caudal striatum or in the nucleus accumbens of the rat, twenty-four hours after administration. However, it has previously been reported that haloperidol can decrease somatostatin gene expression in the striatum, nucleus accumbens and in the frontal cortex of the rat after chronic treatment (Beal and Martin, 1984; Radke *et al.*, 1988; Salin *et al.*, 1989). It has also been suggested that dopamine may be involved in the regulation of somatostatin mRNA levels after chronic treatment (Salin *et al.*, 1989). The lack of effect with haloperidol or fluphenazine after twenty-four hours may reflect the time course necessary to affect the levels of this brain neuropeptide. It seems most likely that the decrease in somatostatin gene expression in the rat striatal and accumbens regions, observed after chronic treatment with the typical neuroleptic agent haloperidol arises after a prolonged time period, and thus may be associated with the appearance of extrapyramidal side effects, which also appear after prolonged drug use (Creese *et al.*, 1976; Seeman *et al.*, 1976; Baldessarini, 1985).

In contrast to this, a low dose of the atypical neuroleptic clozapine caused a significant increase in the levels of preprosomatostatin mRNA in both the caudal region of the striatum and in the nucleus accumbens of the rat after twenty-four hours. In chronic studies, it has previously been suggested that high doses of clozapine can cause an induction of preprosomatostatin mRNA in the rat nucleus accumbens (Salin *et al.*, 1989). These results provide the first evidence for a differential regulation of preprosomatostatin gene expression

after acute treatment with typical versus atypical antipsychotic drugs. It appears that the time course for the atypical agent, clozapine, to induce somatostatin gene expression is relatively more rapid than the time course for haloperidol, the typical neuroleptic, to cause a reduction in somatostatin mRNA levels. The underlying mechanisms involved in the clozapine-induced expression of somatostatin may be due to blockade of dopamine receptors. The low dose of clozapine used in this study may reflect the possibility that the D4 dopamine receptor is involved in this effect, as clozapine has a ten-fold higher affinity for the D4 receptor than for the D2 or D3 dopamine receptors, previously demonstrated in in vitro binding studies (Van Tol *et al.*, 1991). High doses of clozapine (greater than 10mg/kg) have been shown to bind to striatal D2 sites in rats (Saller *et al.*, 1987; Audinot *et al.*, 1993). The low dose of 3mg/kg used here, is preferentially blocking D4 sites in vivo rather than D2 sites. The atypical neuroleptic clozapine is also known to have relatively high affinity for some serotonin receptor subtypes, in particular the serotonin 5-HT_{2A} and 5-HT_{2C} receptors (Zifa and Fillion, 1992; Canton *et al.*, 1993). The distribution of the receptor mRNA's encoding both the 5-HT_{2A} and the 5-HT_{2C} serotonin receptors in the rat, has previously been demonstrated. Intermediate levels of both subtypes have been shown in the caudate-putamen, nucleus accumbens and substantia nigra pars compacta. However, in the caudate-putamen their distributions are quite different. The 5-HT_{2A} receptor is most abundant in the medial striatum, and the 5-HT_{2A} receptor mRNA has been shown to increase in the mediolateral direction. Expression of the 5-HT_{2C} receptor mRNA, present in the medial striatum has been shown to decrease in a mediolateral direction, and in addition, it was expressed in patches of higher density rather than an even distribution (Pompeiano *et al.*, 1993). Thus the possibility that clozapine is acting via a serotonergic mechanism to increase somatostatin gene expression in both the striatum and nucleus accumbens, after acute treatment, is not unlikely. However, both haloperidol and fluphenazine have been shown to have

some affinity for 5-HT_{2A/2C} receptors, but with much less potency than clozapine (Canton *et al.*, 1993). Also, it has recently been reported that intracerebroventricular injection of 5,7-dihydroxytryptamine, a selective serotonin neurotoxin, can increase preprosomatostatin mRNA levels in the rat striatum. From this, it has been suggested that serotonin has an inhibitory effect on the biosynthesis of somatostatin in the striatum (Bendotti *et al.*, 1993).

Antipsychotic drugs which have strong affinities for 5-HT₂ binding sites are known to be associated with a low incidence of extrapyramidal side effects (EPS) (Meltzer, 1988). The role of the 5-HT_{2A} and 5-HT_{2C} receptor sites are at present unclear. However, the antagonism of these sites could contribute to clozapine's low EPS profile. Clozapine has also been reported to antagonise the inhibitory effect of 5-HT₃ receptor agonists on the firing of cortical receptors in the rat (Ashby *et al.*, 1989). Blandina *et al.* (1988) have shown that 5-HT₃ receptor stimulation may facilitate striatal dopamine release. However, the extent to which the clozapine-induced release of striatal dopamine involves 5-HT₃ receptor mechanisms is at present uncertain.

A similar induction of preprosomatostatin mRNA was seen in both the rat striatum and nucleus accumbens regions after treatment with the non-selective 5-HT_{1/2} receptor antagonist, methiothepin. This result further substantiates the possibility of a role for a 5-HT receptor subtype in this response. Methiothepin has previously been demonstrated to have a very high affinity for the 5-HT_{2A/2C} receptors, as shown in binding studies (Leysen *et al.*, 1982; Conn and Sanders-Bush, 1986). Since the atypical antipsychotic drug clozapine also has a high affinity for both the 5-HT_{2A} and 5-HT_{2C}, it could be hypothesised that the blockade of either or both of these serotonin receptor subtypes is a contributing factor to the increase in preprosomatostatin mRNA observed after acute administration of clozapine and methiothepin.

A short time course of the effect of low doses of clozapine on preprosomatostatin mRNA, in the rat striatum was then performed in order to

further address this drug-specific acute effect. However, we observed no significant changes in the levels of preprosomatostatin mRNA up to three hours after the drug was administered.

3.15 Effect of Chronic Antipsychotic treatment on Preprosomatostatin mRNA levels

In this study we have also demonstrated that low doses of clozapine can induce preprosomatostatin mRNA in the rostral and caudal regions of the rat striatum and in the olfactory tubercles, after chronic administration. Previous evidence demonstrating an induction of somatostatin gene expression in the accumbens, involved the administration of much higher doses of clozapine than the dose used here. The low dose of 3mg/kg clozapine may be reflecting a preferential D4 dopamine receptor blockade rather than D2 receptor blockade in order to increase the levels of preprosomatostatin mRNA. However, in previous studies the typical antipsychotic drug haloperidol has been shown to have an opposite effect on somatostatin mRNA levels. Chronic treatment with haloperidol was shown to decrease somatostatin mRNA levels in the striatum, nucleus accumbens and frontal cortex in the rat (Salin *et al.*, 1989). This effect is thought to be the result of blockade of a stimulatory action of dopamine on somatostatinergic neurons mediated through dopamine D2-like receptors (Saller and Salama, 1986). Therefore, it seems unlikely that the effect of low doses of clozapine to increase somatostatin mRNA is mediated by blockade of dopaminergic transmission at D2 or D3 receptors. There is also evidence for an opposing action of typical versus atypical neuroleptics on muscarinic receptors in the striatum. The typical neuroleptic fluphenazine, has been shown to decrease muscarinic receptor density in the striatum, whereas in contrast, the atypical neuroleptic clozapine was shown to elevate muscarinic binding

densities in the striatum, cortex and in the hippocampus in the rat (Friedman *et al.*, 1983). The effect of clozapine to induce somatostatin gene expression is likely to involve blockade of the D4 dopamine receptor or blockade of another neurotransmitter system in the brain. A possible candidate would be the serotonergic system. As previously mentioned, clozapine is a very potent antagonist at certain serotonin receptor subtypes, in particular the 5-HT_{2A/2C} subtypes. The effect of clozapine to induce somatostatin gene expression in the olfactory tubercles may also be mediated by a serotonin receptor blockade. It has been shown that the olfactory tubercles have intermediate levels of mRNA for both the 5-HT_{2A} and 5-HT_{2C} receptor subtypes in the rat (Pompeiano *et al.*, 1993). Recently clozapine has also been demonstrated to have a relatively high affinity for the 5-HT₆ and 5-HT₇ serotonin receptor subtypes and it has been shown that the 5-HT₆ receptor is expressed at high levels in the rat striatum (Monsma *et al.*, 1993). Since a number of atypical neuroleptics have been shown to possess a high affinity for the 5-HT₆ receptor, it has been suggested that this receptor subtype may be involved in mediating the antipsychotic actions of certain neuroleptic drugs (Monsma *et al.*, 1993). In a recent report, fluphenazine has been demonstrated to have a high affinity for the cloned 5-HT₆ subtype, with a K_i value of less than 20nM, although haloperidol had essentially no affinity for the 5-HT₆ receptor (Roth *et al.*, 1993). In addition to its known affinity for certain 5-HT receptor subtypes, clozapine is well known to have an antagonising action on many different receptors, including α -adrenergic receptors and muscarinic receptors, which must still be considered in this effect of the drug.

It has been hypothesised that a disruption of somatostatin gene regulation may have a role in a number of identified psychoses in man, such as schizophrenia, parkinsonism, Huntington's disease and alcohol dementia. It has been demonstrated that individuals with Parkinson's disease have lower levels of somatostatin-like immunoreactivity in lumbar spinal fluid than their control

counterparts (Cramer *et al.*, 1989). Similarly a recent study has shown the reduction of somatostatin peptide in the neocortex of Parkinson's disease sufferers (Leake and Ferrier, 1993). A functional interaction between central nervous system somatostatin and dopamine systems has been implicated, as fluphenazine treatment has been shown to reduce cerebrospinal fluid somatostatin levels in patients with schizophrenia (Doran *et al.*, 1989). Another study has demonstrated a decrease in somatostatin-like immunoreactivity in the cerebrospinal fluid of chronic schizophrenic patients with a cognitive impairment (Reinikainen *et al.*, 1990). The development of schizophrenia has previously been suggested to involve autoimmune processes, such that blood serum samples taken from schizophrenic patients manifest a higher level of immune responsiveness to somatostatin, as compared to the control group (Rogaeva *et al.*, 1990). A further level of complexity in the development of schizophrenia has been implicated in this study, since the increased immune responsiveness in blood serum to somatostatin was evident in patients suffering from malignant and paranoid schizophrenia, but was not evident in patients suffering from a slow-progressive form of schizophrenia (Rogaeva *et al.*, 1990). This would suggest that different forms of the disease can differentially affect somatostatin levels. From this, it would appear that a number of physiological changes are necessary for the manifestation of schizophrenia, although the 'chemical balance' may be affected differently in different forms of the disease. In Huntington's chorea, a marked down-regulation of somatostatin receptors in the human striatum has been shown (Palacios *et al.*, 1990). It has been postulated that the down-regulated somatostatin receptors may be localised on a population of neurons which are at risk in Huntington's chorea (Palacios *et al.*, 1990). A reduction in cerebrospinal fluid somatostatin levels has also been demonstrated in alcoholic amnesic disorder or Korsakoff's psychosis (Martin *et al.*, 1989) and in senile dementia of the Alzheimer's type (Leake and Ferrier, 1993). Therefore it would seem that a decrease in

somatostatin levels is one of the major physiological changes that takes place in a number of cognitive disorders.

Short-term Regulation of Neuropeptide Y gene expression in vivo

3.16 The Effect of Antipsychotic drug treatment on the levels of Proneuropeptide Y gene expression twenty-four hours after administration

Neuropeptide Y, a peptide which shows many of the properties characteristic of a neurotransmitter or neuromodulator is known to be highly abundant in mammalian brain. It has previously been detected in the same neurons as catecholamines and enkephalins (Hunt *et al.*, 1981; Hokfelt *et al.*, 1983; Everitt *et al.*, 1984), and many neurons in the forebrain have been shown to contain both neuropeptide Y and somatostatin (Vincent and Johansson, 1983; Chan-Palay, 1987). In the striatum a small population of neurons are known to express both neuropeptide Y and somatostatin (Graybiel, 1990; Vuillet *et al.*, 1990). Therefore we were interested in the effect of acute treatment with the atypical neuroleptic clozapine and the serotonin receptor antagonist, methiothepin, on the levels of proneuropeptide Y mRNA in the striatum, as these drugs have previously been demonstrated to increase somatostatin mRNA levels in the striatum.

However, the results obtained indicate a lack of effect of both clozapine and methiothepin on proneuropeptide Y mRNA levels in either the rostral or caudal regions in the rat striatum. Due to the small number of animals per treatment group, any small changes in the level of neuropeptide mRNA may remain undetected despite a possible drug effect.

In previous studies selective D1 dopamine receptor blockade has been demonstrated to increase cellular levels of neuropeptide Y mRNA in the rat striatum (Morris, 1992). This is in contrast to the decrease in somatostatin mRNA levels that have been reported in the same population of neurons following chronic D2 receptor blockade (Weiss and Cheeselet, 1989; Salin *et al.*, 1989). It has therefore been suggested that dopamine is tonically and simultaneously exerting an opposite control on neuropeptide Y and somatostatin gene expression in the striatum (Morris, 1992). In view of this, it may not be surprising that both clozapine and methiothepin had no effect on neuropeptide Y gene expression. If neuropeptide Y and somatostatin are differentially regulated within the same cell, it would perhaps have been more likely to see a decrease in proneuropeptide Y mRNA levels, although our results did not show this to occur. Similarly the neuropeptides dynorphin and preprotachykinin which are expressed in striatonigral neurons have previously been shown to be differentially regulated by dopamine in the same cell population (Gerfen *et al.*, 1991).

3.17 The effect of chronic neuroleptic drug treatment on dopamine D1 receptor expression

It has long been known that chronic treatment with antipsychotic drugs can up-regulate dopamine receptor subtypes. The class of drugs known as 'typical neuroleptics' such as haloperidol, have previously been demonstrated to increase the mRNA encoding the D1A, the D2 and the D3 dopamine receptor subtypes (Buckland *et al.*, 1992). The levels of mRNA encoding the D1B/D5 receptor in the rat have been reported to remain unchanged after chronic haloperidol treatment (Buckland *et al.*, 1992). The results obtained from this study demonstrate that the typical neuroleptic fluphenazine, when

given chronically, can significantly increase D1A receptor mRNA levels in the rostral region, in the rat striatum. This is in accordance with previous reports suggesting that blockade of dopamine D1 receptors can increase D1 receptor mRNA (Creese *et al.*, 1992). The effect of fluphenazine did not prove to be significant in the caudal region of the striatum, despite a tendency towards an increase in D1A mRNA expression.

With the 'atypical' neuroleptic drug clozapine an elevation of D1A receptor mRNA levels was also observed in the rostral striatum, although the increase was less than that seen with fluphenazine. The dose of clozapine used in this study would not be expected to be occupying more than 30-35% of D1 receptors *in vivo*, according to the demonstrated affinities of clozapine at D1 compared to D4 dopamine receptors (Van Tol *et al.*, 1991). This therefore makes the observed induction D1 receptor mRNA in the striatum after chronic clozapine treatment relatively difficult to interpret. However, it is not impossible that only a small percentage of D1 receptors need to be occupied in order to increase D1 mRNA. In previous reports chronic treatment with high doses of clozapine has been shown to both increase (O' Dell *et al.*, 1990) and have no effect on the density of dopamine D1A receptors (Seeger *et al.*, 1982; Severson *et al.*, 1984; Lee and Tang, 1984; Rupniak *et al.*, 1984; Cohen and Lipinski, 1986; Buckland *et al.*, 1993). Other workers have studied the effect of clozapine on D2 receptor mRNA and receptor numbers. Wilmot and Szczepanik, (1989) found no increase in D2 receptor numbers in any rat brain regions. Similarly Van Tol *et al.* (1991) found no increase in D2 receptor mRNA levels in rat striatum following chronic treatment with clozapine. Although, chronic clozapine treatment has recently been associated with an increase in D3 receptor mRNA levels. Buckland *et al.*, (1993) have demonstrated that high doses of 30mg/kg clozapine can increase D3 receptor mRNA levels by five-fold after four days of treatment. In addition, they also found that the levels of D3 receptor mRNA dropped back to basal levels after

thirty-two days of treatment. In the previous studies that suggest clozapine has no effect on the levels of D1A receptor mRNA, it is likely that monitoring of mRNA levels occurred after twenty days or more. It is possible that the increase in D1A receptor mRNA levels that we observed after five days of treatment has declined back to basal levels after three weeks of treatment, similar to that observed for the D3 receptor. Therefore it would be necessary to monitor the level of D1A receptor mRNA more closely, over a prolonged time period, in order to clarify this result.

3.18 The effect of chronic neuroleptic drug treatment on the expression of the 5-HT6 receptor mRNA

Clozapine has previously been reported to have a high affinity for the serotonergic 5-HT6 receptor in binding studies (Monsma *et al.*, 1993). Therefore we were interested to see the effects of clozapine and the typical neuroleptic fluphenazine on the levels of 5-HT6 receptor mRNA in the rat striatum, following chronic drug treatment.

However, in this study both clozapine and fluphenazine had little effect on the levels of 5-HT6 mRNA in vivo after five days of chronic treatment. This would suggest that the 5-HT6 receptor is not up-regulated by chronic blockade with both typical and atypical neuroleptic drugs. Also it may indicate that the serotonergic control of this receptor subtype is low, in that there may be a low tonic activation of 5-HT6 receptors in the striatum.

Chapter 4

Regulation of Immediate Early Genes in vivo

INTRODUCTION

4.0 Neuroleptic Modulation of Immediate Early Genes In Vivo

At present, immediate early genes are thought to function by altering the rate of transcription of a target gene, by binding to a specific site on the late response gene. There has been a large amount of evidence gained to suggest that modulation of the level of dopaminergic activity can induce the immediate early genes *c-fos* and *zif/268*. Systemic treatment with dopaminergic agonists, in particular SKF38393, increases the levels of *c-fos* and *zif/268* after a 6-hydroxydopamine-induced lesion (Robertson *et al.*, 1990; Cole *et al.*, 1992), whereas evidence also exists showing that they are induced in the rat striatum after administration of haloperidol, a dopaminergic D2 receptor antagonist (Dragunow *et al.*, 1990; Nguyen *et al.*, 1992). This induction of both *c-fos* and *zif/268* by typical neuroleptics is likely to be mediated by the blockade of either D2 or D4 dopamine receptors, because drugs active on these receptors share the ability to increase the expression of these immediate early genes (Dragunow *et al.*, 1990; Cole *et al.*, 1992; Nguyen *et al.*, 1992; Robertson and Fibiger, 1992). By using the neuroleptic drug clozapine, which has a ten fold higher affinity for the dopamine D4 receptor than for the D2 receptor, given at low doses, we have attempted to investigate further the nature of the receptor involved in these responses.

Currently there is very little evidence to demonstrate the hypothesis that immediate early genes may provide the link to changes in late response genes. However, it has been suggested that there is a link between the haloperidol induction of *c-fos* and the subsequent increase in the levels of preproenkephalin mRNA (Sonnenberg *et al.*, 1989; Donovan *et al.*, 1992). More recently this circumstantial evidence has been questioned by results obtained by Konradi and

colleagues (1993). At present there is no indication as to the mechanisms leading to a decrease in preprotachykinin gene expression.

It is known that c-fos forms dimers with members of the jun family of immediate early genes to affect target gene transcription. The mRNA encoding two members of this family, c-jun and jun-D, is present constitutively in striatal neurons, and the corresponding proteins are likely to be available to form complexes with c-fos once it is synthesised. The mRNA encoding another immediate early gene belonging to this family, jun-B, is not normally detectable in striatal neurons. However, jun-B is potentially of interest, because in primary rat embryo cells it has been linked to the suppression, rather than the enhancement of downstream gene transcription (Chiu *et al.*, 1989; Schutte *et al.*, 1989). We were therefore interested to determine whether jun-B and jun-D is induced in the striatum and nucleus accumbens, by neuroleptic administration.

The ^{proteins encoded by the} newly cloned POU-domain -containing gene, RHS2 and the HNF/3 fork head gene BF-1, have recently been shown to possess a transcription regulatory region. However, as yet there has been very little research done on these genes. As both gene ^{products} have been shown to be present in the region of the striatum (Le Moine and Scott-Young, 1992; Tao and Lai, 1992), we have been interested in the effects of typical and atypical neuroleptic drug administration on the expression of these two genes in the rat striatum.

RESULTS

4.1 Time course of Fluphenazine effect on c-fos mRNA levels in the region of rostral and caudal striatum

It is known that the immediate early gene c-fos can be induced by administration of the relatively non-selective, typical neuroleptic drug haloperidol (Nguyen *et al* ,1992). Therefore in this study we have investigated the effects over time of the typical neuroleptic drug fluphenazine (6mg/kg), having high affinity for, and being a potent antagonist of the D2 dopamine receptor.

Following intraperitoneal administration of fluphenazine (6mg/kg) to male wistar rats (n= 3), it was observed that the levels of c-fos mRNA were increased significantly in the rostral striatum 15 minutes [F (4,20) =7.60, p= 0.008, p< 0.01] and 45 minutes, p< 0.01, after drug treatment compared to the animals treated at corresponding time points with physiological saline (n=2) [Figure 4.1.1]. After 45 minutes the levels of c-fos mRNA declined back to basal levels.

In the caudal striatum , a maximal and significant induction of c-fos mRNA levels after fluphenazine treatment occurred 45 minutes after drug administration. [Figure 4.1.2] The levels of c-fos mRNA were significantly different such that F (4,20) =8.67, p= 0.0067, p< 0.01, from saline treated animals. A similar decline back to basal levels was observed after 45 minutes. The pattern of induction of c-fos mRNA was primarily observed in the dorsolateral striatum.

Figure 4.1.1 : Effect of fluphenazine (6mg/kg) on c-fos mRNA levels in the rat rostral striatum. Fluphenazine (6mg/kg; n= 3) or vehicle (0.9% saline; n= 2) was injected i.p. and the animals killed after the specified time intervals. Results, obtained from film autoradiographs are expressed as mean relative optical density \pm s.e.m. The open triangle = saline treated groups, closed triangle = fluphenazine (6mg/kg) treated groups for the respective time periods. Significance was determined by one-way ANOVA, followed by the student's t-test. *p< 0.05.

Figure 4.1.2 : Effect of fluphenazine (6mg/kg) on c-fos mRNA levels in the rat caudal striatum. Fluphenazine (6mg/kg; n= 3) or vehicle (0.9% saline; n= 2) were injected i.p. and the animals killed after the specified time intervals. Results, obtained from film autoradiographs, are expressed as mean relative optical density \pm s.e.m. The open triangles = saline treated groups, closed triangles = fluphenazine (6mg/kg) treated groups for the respective time periods. Significance was determined by one-way ANOVA, followed by the student's t-test. *p< 0.05

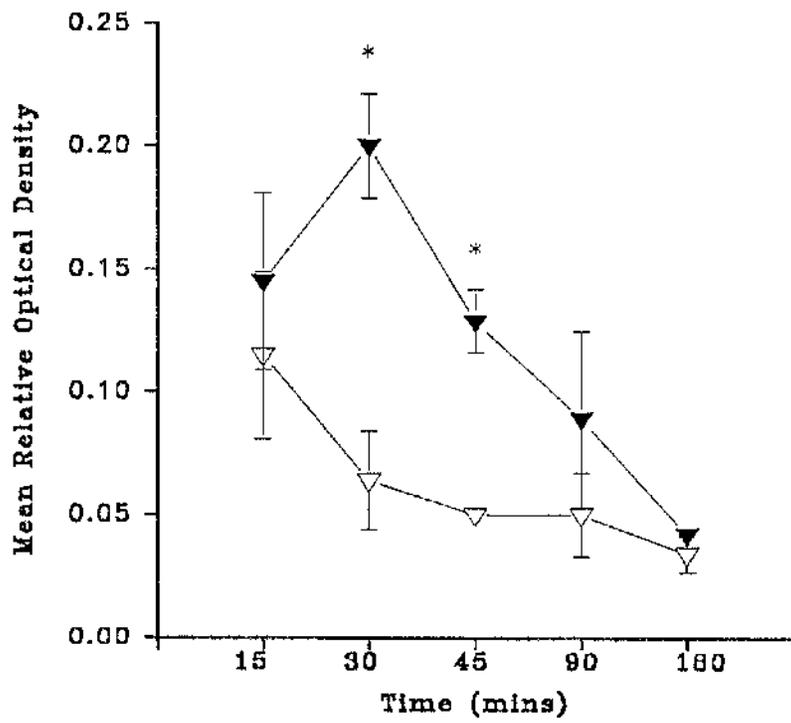
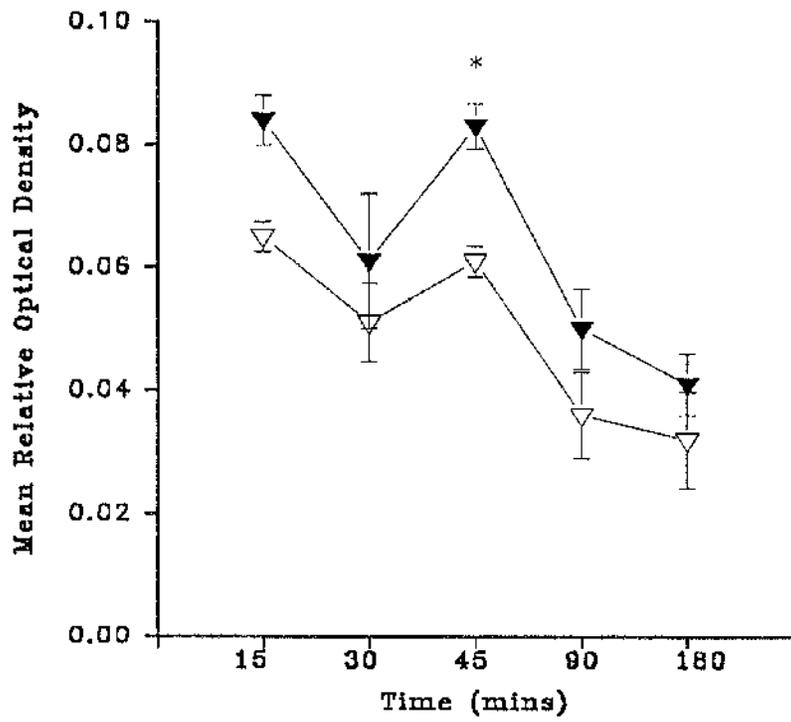
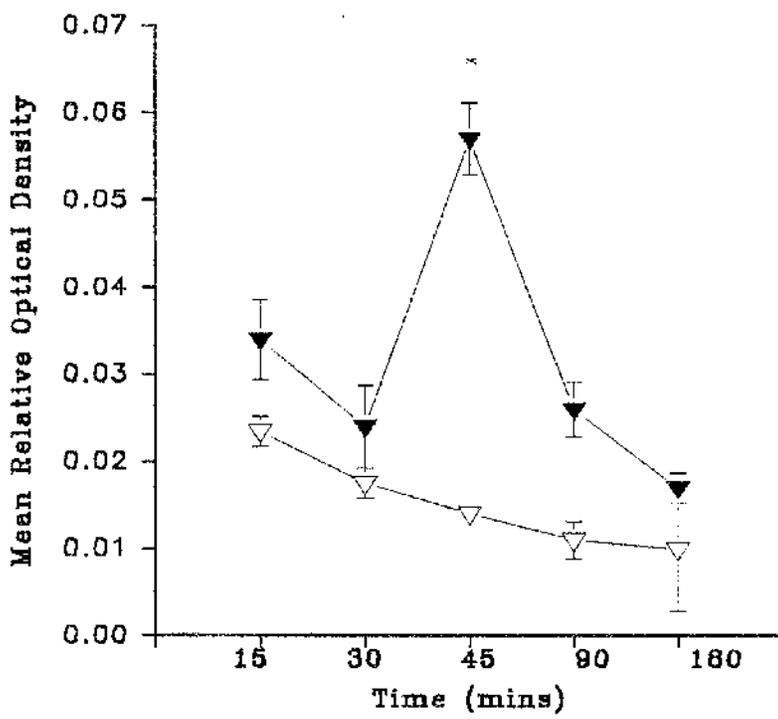
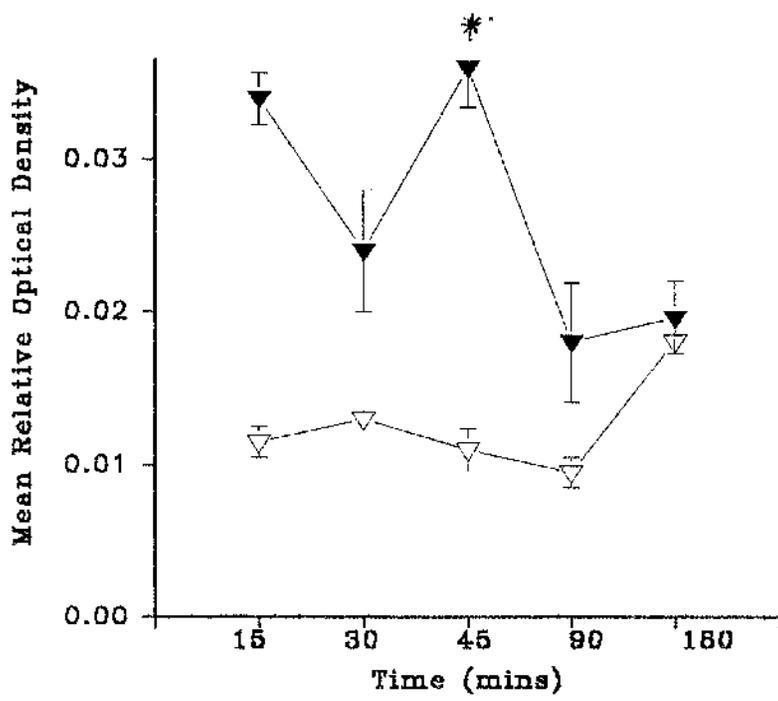


Figure 4.2.1 : Effect of fluphenazine (6mg/kg) on zif/268 mRNA in the rat rostral striatum. Fluphenazine (6mg/kg; n= 3) or vehicle (0.9% saline; n= 2) were injected i.p. and the animals killed after the specified time intervals. Results, obtained from film autoradiographs, are expressed as mean relative optical density \pm s.e.m. The open triangles = saline treated groups, closed triangles = fluphenazine (6mg/kg) treated groups for the respective time periods. Significance was determined by one-way ANOVA, followed by the student's t-test. * $p < 0.05$

Figure 4.2.2 : Effect of fluphenazine (6mg/kg) on zif/268 mRNA levels in the rat caudal striatum. Fluphenazine (6mg/kg; n= 3) or vehicle (0.9% saline; n= 2) were injected i.p. and the animals killed after the specified time intervals. Results, obtained from film autoradiographs, are expressed as mean relative optical density \pm s.e.m. The open triangles = saline treated groups, closed triangles = fluphenazine (6mg/kg) treated groups for the respective time periods. Significance was determined by one-way ANOVA, followed by the student's t-test. * $p < 0.05$.



4.2 Time course of Fluphenazine effect on zif/268 mRNA levels in the regions of the rostral and caudal striatum

Zif/268, a zinc-finger containing protein, is also known to be induced by treatment with the typical neuroleptic haloperidol in the striatum (Nguyen *et al.*, 1992). Therefore the time course effect on zif/268 mRNA induction by fluphenazine (6mg/kg, n= 3) was investigated.

The levels of zif/268 mRNA were significantly induced compared to saline treated animals, [F (4,20) =4.89, p= 0.04, p< 0.05] after 45 minutes in the rostral striatum. [Figure 4.2.1]. However, in the caudal striatum it was observed that zif/268 mRNA levels were induced 30 minutes after fluphenazine (6mg/kg, n= 3) administration. This increase was significantly different from saline treated animals (n= 2) at the corresponding time point, such that F (4,20) =3.50, p= 0.047, p< 0.05, DF= 1 [Figure 4.2.2]. Induction of zif/268 was significantly different from saline treated animals 45 minutes after drug administration, p< 0.05, following which, a gradual decline back to basal levels was observed. [Figure 4.2.2]. Again, the pattern of induction was primarily observed in the dorsolateral regions of the rat striatum.

4.3 Effect of acute neuroleptic drug administration on c-fos mRNA and zif/268 mRNA expression in the rat striatum and nucleus accumbens

<u>Drug Treatment</u>	<u>Striatum</u>	<u>Nucleus accumbens</u>
Saline	26.80±5.70	25.50±5.50
Haloperidol	144.00±19.00 *	101.70±24.50 *
Fluphenazine	60.60±21.10 *	18.50±2.10
Clozapine	23.70±6.30	20.70±3.70

Table 1: The expression of c-fos mRNA levels in the rat striatum and nucleus accumbens following treatment with vehicle (physiological saline), or the typical neuroleptics haloperidol (1mg/kg), and fluphenazine (3mg/kg) or the atypical neuroleptic clozapine (3mg/kg). Results, obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^3 \pm$ s.e.m., n=3-5. Significance was determined by one-way ANOVA, followed by the *post-hoc* Mann-Whitney U test. $p < 0.05^*$

<u>Drug treatment</u>	<u>Striatum</u>	<u>Nucleus accumbens</u>
Saline	46.75 \pm 5.76	41.70 \pm 5.42
Haloperidol	152.30 \pm 13.30 *	95.30 \pm 14.90
Fluphenazine	203.70 \pm 42.10 *	152.00 \pm 36.30 *
Clozapine	64.30 \pm 15.70	43.30 \pm 17.80

Table 2: The expression of zif/268 mRNA levels in the rat striatum and nucleus accumbens following treatment with vehicle (physiological saline), or the typical neuroleptic drugs haloperidol (1mg/kg), and fluphenazine (3mg/kg), or the atypical neuroleptic clozapine (3mg/kg). Results, obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^3 \pm$ s.e.m., n=3-5. Significance was determined by one-way ANOVA, followed by the *post-hoc* Mann-Whitney U test. $p < 0.05^*$

Animals were intraperitoneally injected with saline (n= 3), the typical neuroleptic drugs haloperidol (1mg/kg, n= 3) and fluphenazine (6mg/kg, n= 5)

and the atypical neuroleptic drug clozapine (3mg/kg; n= 3). After 30 minutes the levels of mRNA were measured using in situ hybridisation histochemistry. Prior to the use of the above drug concentrations, a lower dose of 1mg/kg was used for both fluphenazine and clozapine, however no induction of either c-fos or zif/268 mRNA was observed. The effects of a higher dose of 6mg/kg fluphenazine were also examined in rats, in the event that a higher dose may cause much more potent inductive effects. However, it was demonstrated that 6mg/kg fluphenazine caused a comparable increase in c-fos and zif/268 mRNA levels to the induction observed with the lower dose of 3mg/kg fluphenazine (data not shown).

The above tables demonstrate that low levels of both c-fos mRNA [Table 2] and zif/268 mRNA [Table 3] are observed following treatment with physiological saline, the basal zif/268 hybridisation signal being approximately double that for c-fos mRNA, in both the rat striatum and nucleus accumbens.

The levels of c-fos mRNA in the striatum 30 minutes after treatment with the typical neuroleptic haloperidol (1mg/kg) were significantly increased compared to animals treated with saline, such that $F = 50.56$, $p = 0.011$, $p < 0.05$ [Table 2]. Haloperidol had a tendency to increase c-fos mRNA expression in the nucleus accumbens, and the increase was significantly different from saline treated animals ($F = 15.58$, $p < 0.05$). Similarly fluphenazine (3mg/kg) significantly induced c-fos mRNA levels in the striatum ($F = 10.16$, $p < 0.05$), but not in the nucleus accumbens. Clozapine (3mg/kg) had no effect on the expression of c-fos in both the striatum and nucleus accumbens. [Table 2].

A similar pattern of induction was observed for zif/268 mRNA in the striatum. The increase in zif/268 mRNA levels after administration of haloperidol (1mg/kg) for 30 minutes was significantly different from control rats ($F = 42.63$, $p < 0.01$) [Table 3]. A tendency towards induction of zif/268 was observed in the nucleus accumbens following haloperidol (1mg/kg), although this increase failed to reach significance. Fluphenazine (3mg/kg) caused a

significant induction of *zif/268* mRNA in both the striatum ($F=8.85$, $p < 0.05$) and nucleus accumbens ($F=7.94$, $p < 0.05$). Clozapine (3mg/kg) had no effect on the levels of *zif/268* mRNA in both the striatum and the nucleus accumbens.[Table 3].

The expression of *c-fos* and *zif/268* mRNA had a tendency towards slightly higher levels in the dorsolateral region of the striatum, although a general induction was observed throughout the striatum. The levels expressed in the nucleus accumbens were distributed homogeneously over core and shell regions.

4.4 The effect of acute treatment with neuroleptics on the expression of *jun-B* mRNA and *jun-D* mRNA in the rat striatum and nucleus accumbens

In saline treated animals a very low hybridisation signal for *jun-B* mRNA was observed in all brain regions, similar to the low levels of *c-fos* mRNA following saline treatment. Following administration of the typical neuroleptic haloperidol (1mg/kg; $n=3$), a dramatic increase in the levels of *jun-B* mRNA was observed 30 minutes after drug treatment, in the rat striatum ($F=60.41$, $p < 0.05$).[Figure 4.4.1]. A clear induction of *jun-B* mRNA was also observed in the nucleus accumbens ($F=15.80$, $p < 0.05$), following haloperidol (1mg/kg). [Figure 4.4.2]. Similarly fluphenazine (3mg/kg, $n=8$), a typical neuroleptic having a high affinity for the dopamine D2 receptor, was shown to cause a significant induction in the expression of *jun-B* mRNA levels in the rat striatum ($F=4.22$, $p < 0.05$) [Figure 4.4.1, 4.4.5], and in the nucleus accumbens ($F=35.15$, $p < 0.05$) [Figure 4.4.2]. The pattern of *jun-B* expression in the striatum was primarily in the dorsolateral regions, whilst in the nucleus accumbens the tendency was towards a homogeneous distribution of *jun-B* mRNA. The atypical neuroleptic clozapine (3mg/kg, $n=6$) had no effect on the

Figure 4.4.1: Effect of acute administration of neuroleptic drugs on jun-B mRNA levels in the rat striatum. Haloperidol (1mg/kg), or fluphenazine (3mg/kg), or clozapine (3mg/kg), or vehicle (0.9% saline) were injected i.p. 30 minutes before sacrifice. Results, obtained from film autoradiographs, are expressed as mean relative optical density + s.e.m.; n=3-8. Significance was determined by ANOVA followed by the post-hoc Mann-Whitney U test. * p< 0.05 relative to saline group.

Figure 4.4.2: Effect of acute administration of neuroleptic drugs on jun-B mRNA levels in the rat nucleus accumbens. Haloperidol (1mg/kg), or fluphenazine (3mg/kg), or clozapine (3mg/kg), or vehicle (0.9% saline) were injected i.p. 30 minutes before sacrifice. Results, obtained from film autoradiographs, are expressed as mean relative optical density + s.e.m.; n=3-8. Significance was determined by ANOVA, followed by the post-hoc Mann-Whitney U test. *p< 0.05 relative to saline group.

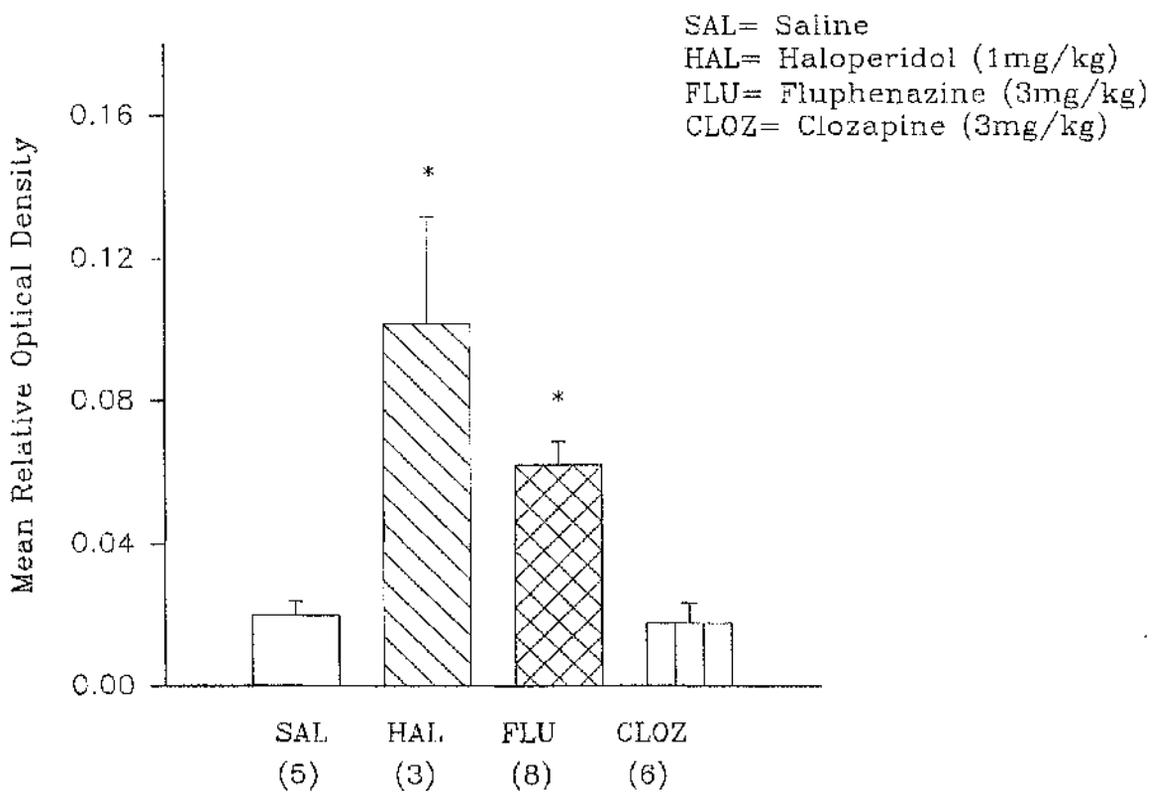
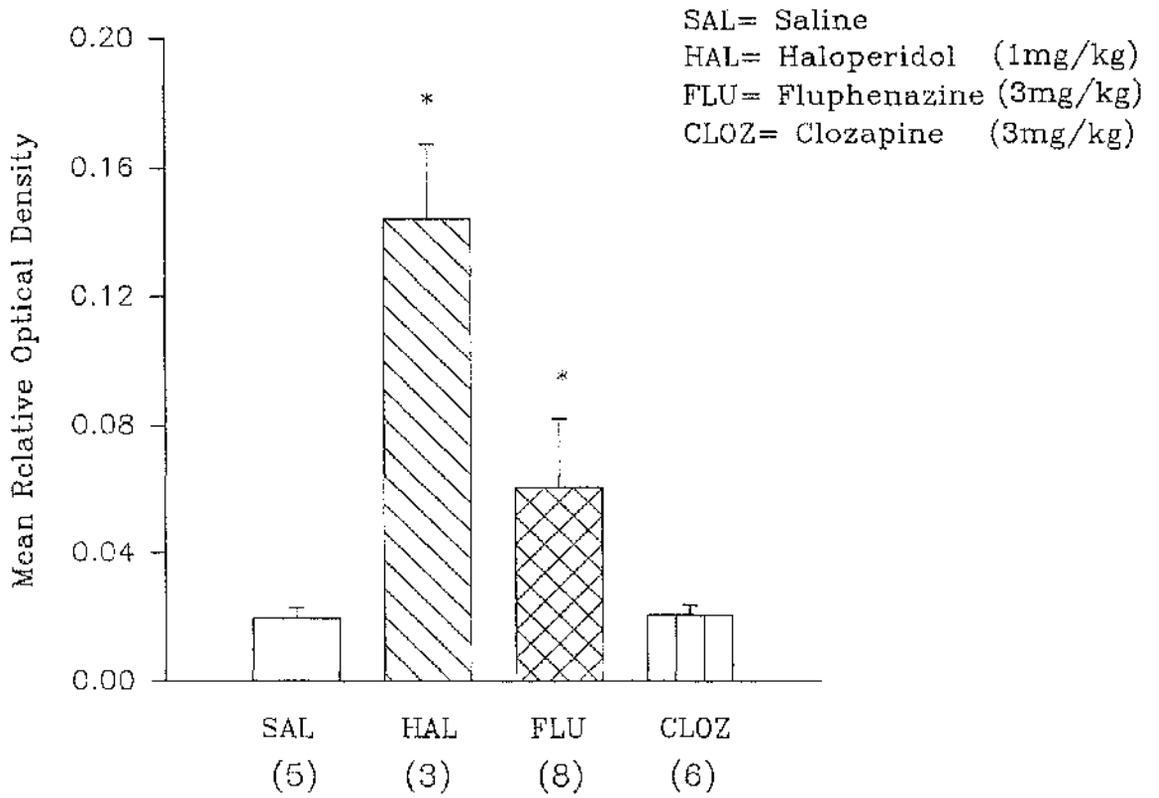


Figure 4.4.3: Effect of acute administration of neuroleptic drugs on jun-D mRNA levels in the rat striatum. Haloperidol (1mg/kg), or fluphenazine (3mg/kg), or clozapine (3mg/kg) or vehicle (0.9% saline) were injected i.p. 30 minutes before sacrifice. Results, obtained from film autoradiographs are expressed as mean relative optical density + s.e.m.; n=3-8. Significance was determined by ANOVA, followed by the post-hoc Mann-Whitney U test. * p< 0.05 relative to saline group.

Figure 4.4.4 : Effect of acute administration of neuroleptic drugs on jun-D mRNA levels in the rat nucleus accumbens. Haloperidol (1mg/kg), or fluphenazine (3mg/kg), or clozapine (3mg/kg), or vehicle (0.9% saline) were injected i.p. 30 minutes before sacrifice. Results, obtained from film autoradiographs, are expressed as mean relative optical density + s.e.m.; n=3-8. Significance was determined by ANOVA followed by the post-hoc Mann-Whitney U test. * p< 0.05 relative to saline group.

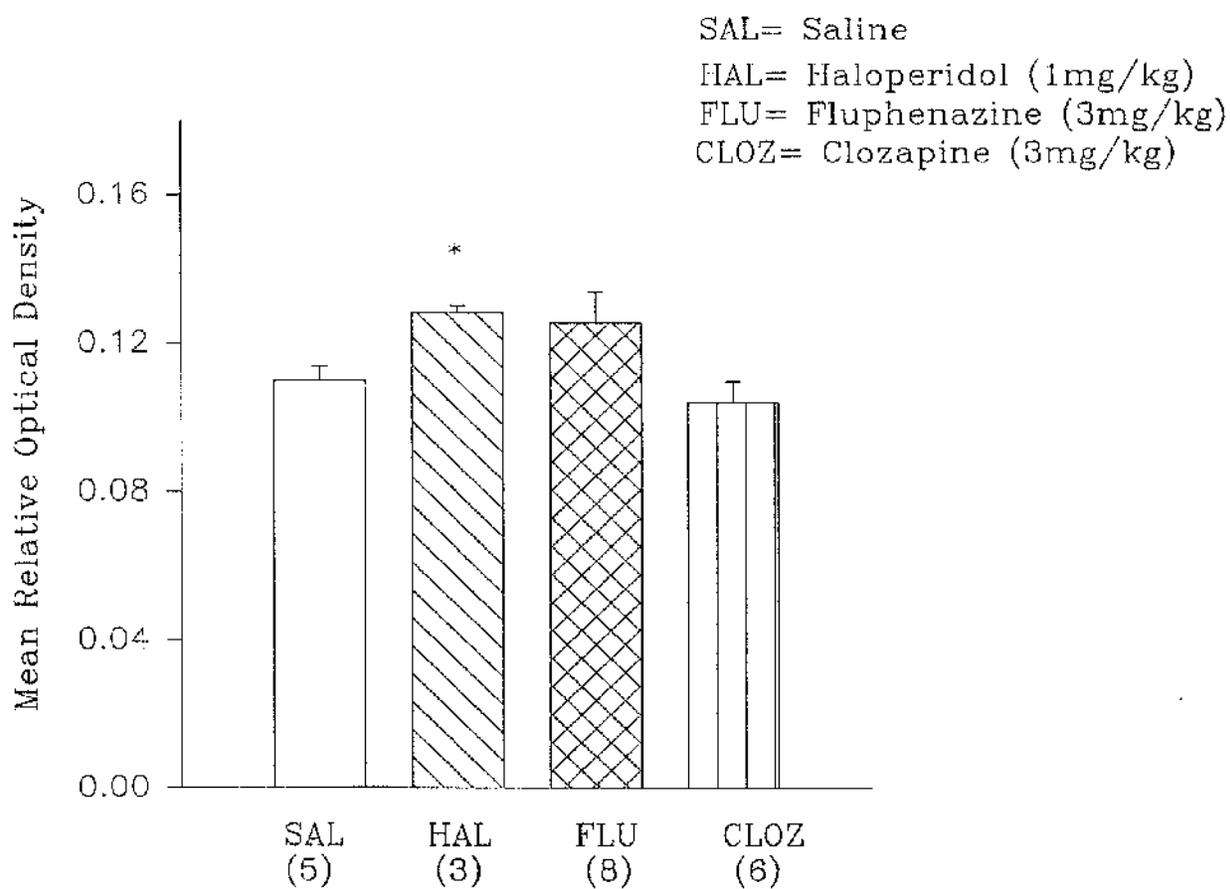
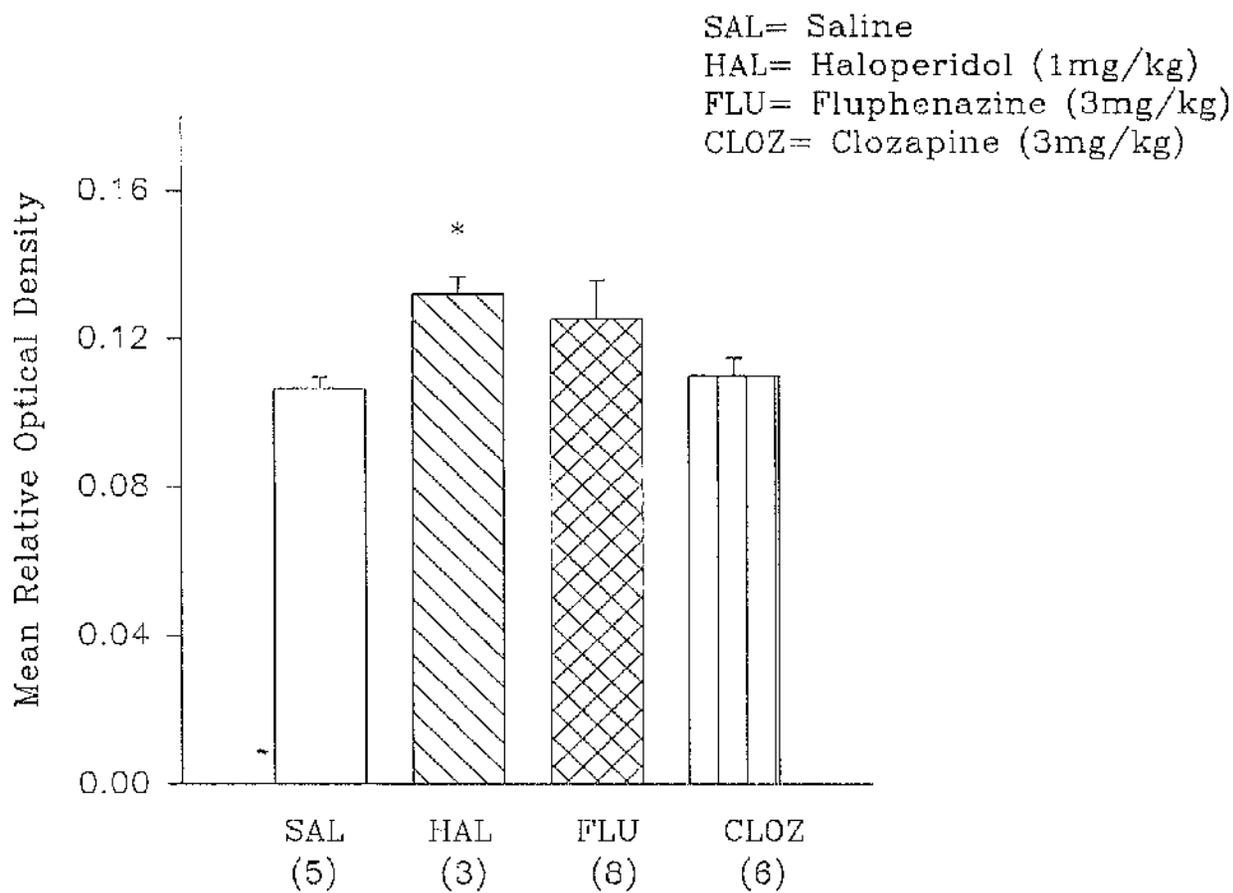
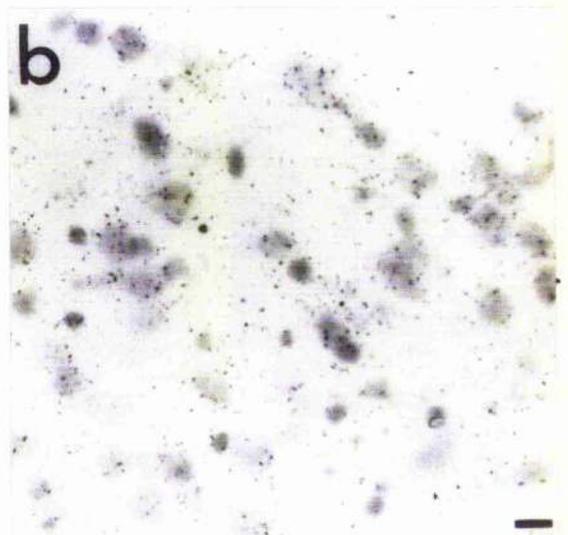
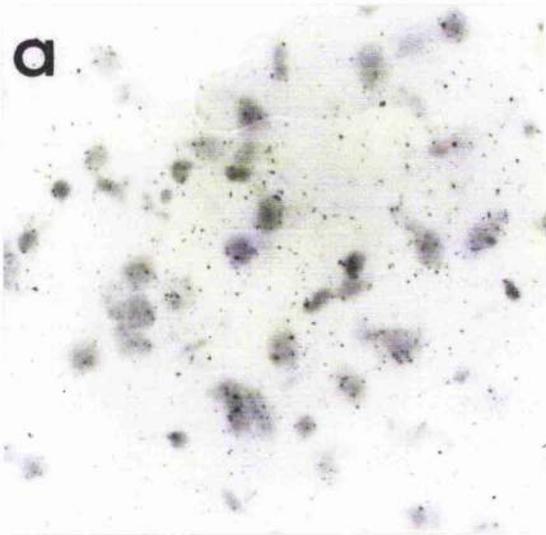
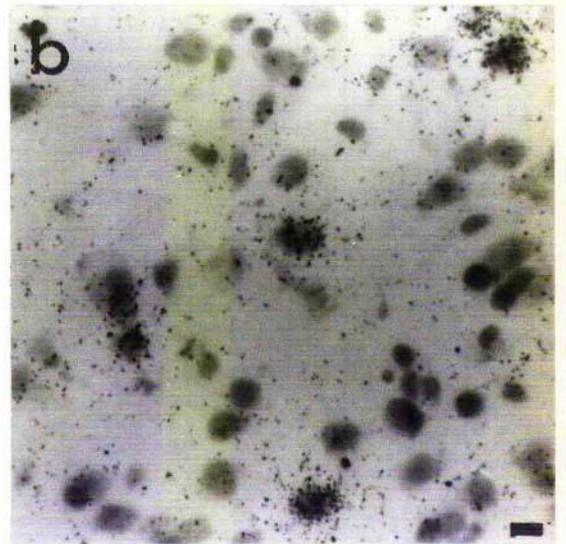
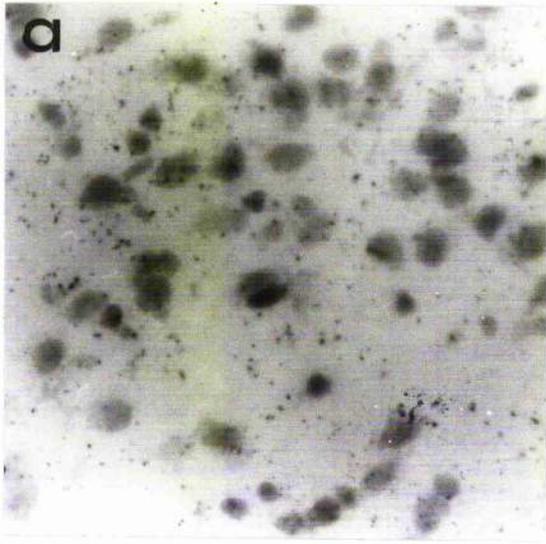


Figure 4.4.5 : Effect of fluphenazine on the expression of jun-B mRNA in the rat striatum. Photomicrographs show the distribution of silver grains over cell bodies in the caudal striatum. On the left, the animal was treated with vehicle (0.9% saline) (A), and on the right, with fluphenazine (3mg/kg), injected i.p. The animals were killed 30 minutes later. Scale bar represents 25 μ m.

Figure 4.4.6 : Effect of haloperidol on the expression of jun-D mRNA in the rat striatum. Photomicrographs show the distribution of silver grains over cell bodies in the caudal striatum. On the left, the animal was treated with vehicle (physiological saline) (A), and on the right , with haloperidol (1mg/kg) (B), administered i.p. The animals were killed 30 minutes later. Scale bar represents 25 μ m.



expression of jun-B mRNA levels in either the striatum or nucleus accumbens compared to saline treated control animals.

Another member of the jun family of immediate early genes, jun-D, was investigated. In comparison to the other IEGs under investigation, jun-D is anomolous in that it is basally expressed in much higher levels throughout the brain. The expression of jun-D mRNA was shown to be increased in the striatum 30 minutes after treatment with haloperidol (1mg/kg, n= 3) [Figure 4.4.3, 4.4.6], compared to saline treated control animals. This induction was shown to be statistically significant relative to control ($F=20.70$, $p < 0.05$). Likewise a similar increase in jun-D mRNA levels were seen in the nucleus accumbens after haloperidol (1mg/kg, n= 3) ($F=19.00$, $p < 0.05$) [Figure 4.4.4]. However fluphenazine (3mg/kg, n= 8) and clozapine (3mg/kg, n= 6) had no observed effect on the expression of jun-D mRNA in either the striatum or nucleus accumbens.

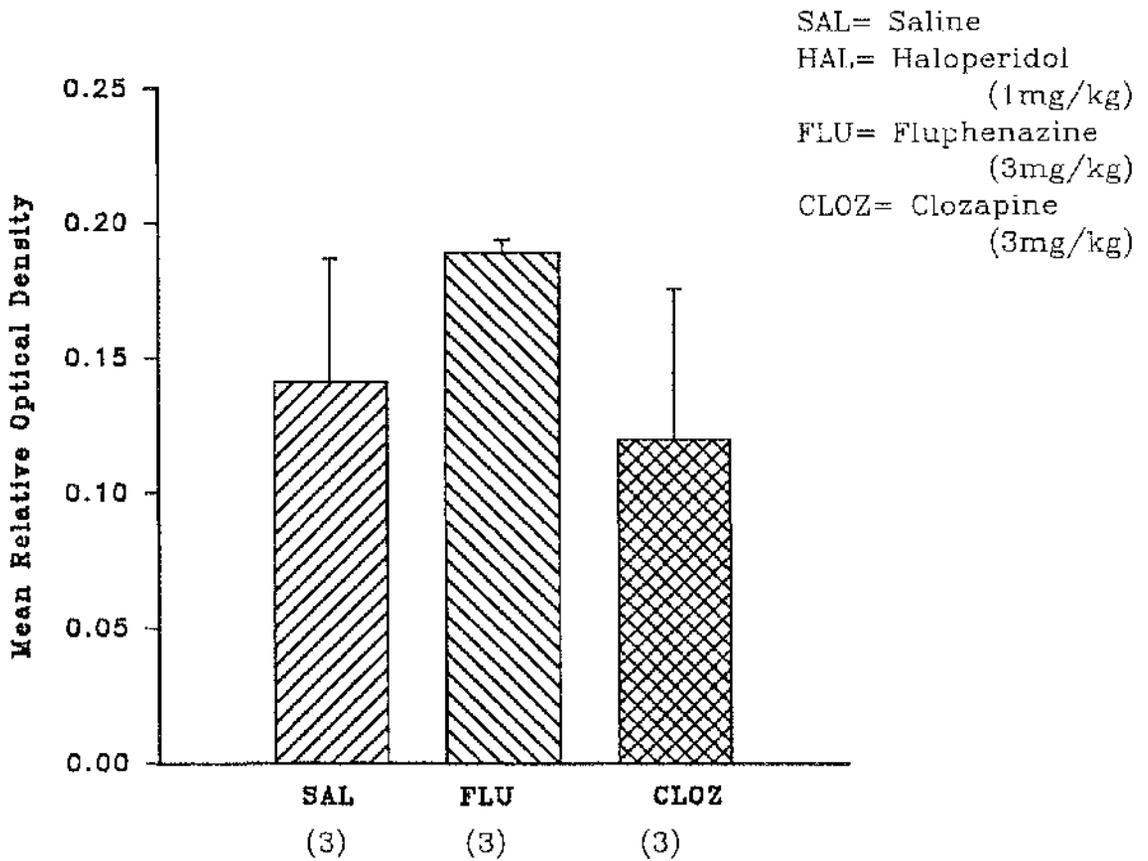
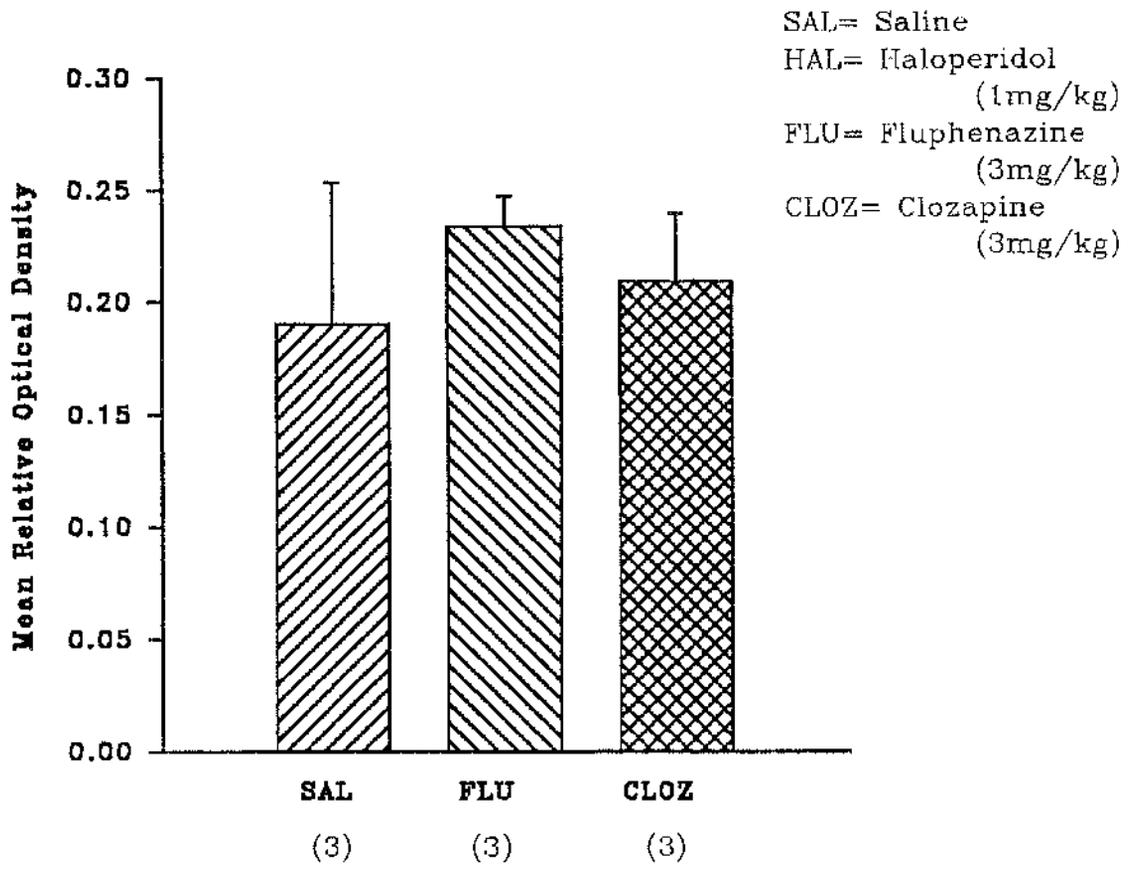
4.5 Effect of acute treatment with neuroleptics on the transcription factor genes BF-1 and RHS2.

The recently cloned homeobox genes BF-1 and RHS2 have been shown to be expressed at moderate levels in the striatum. (Tao and Lai, 1992; Le Moine and Young, 1992). Therefore, we have investigated the effect of neuroleptic drugs on the levels of expression of these genes in the rat striatum.

The hybridisation signal obtained for BF-1 and RHS2 gene expression was quite high in saline treated animals (n= 3). Thirty minutes after the administration of the typical neuroleptic drug fluphenazine (3mg/kg, n= 3) or the atypical neuroleptic clozapine (3mg/kg, n= 3), there was no significant

Figure 4.5.1 : Effect of neuroleptic drug treatment on the expression of BF-1 mRNA in the rat caudal striatum. Three animals per group were injected i.p. with vehicle (0.9% saline), fluphenazine (3mg/kg) or with clozapine (3mg/kg). After 30 minutes the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m. The number of animals per group is shown in parentheses.

Figure 4.5.2 : Effect of neuroleptic drug treatment on the expression of RHS2 mRNA in the rat caudal striatum. Three animals per group were injected i.p. with vehicle (0.9% saline), or fluphenazine (3mg/kg), or with clozapine (3mg/kg). After 30 minutes their brains were processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m. The number of animals per group is shown in parentheses.



change in the levels of expression of BF-1 mRNA [Figure 4.5.1] or RHS2 mRNA [Figure 4.5.2] compared to saline treated control animals.

4.6 Effect of acute neuroleptic drug treatment on the expression of BF-1 mRNA in the rat striatum 24 hours later.

The levels of BF-1 mRNA expressed in saline treated animals were generally distributed in relatively high levels throughout the brain, particularly in the forebrain regions. Following treatment with haloperidol (1mg/kg, n= 3), fluphenazine (3mg/kg, n= 3), or clozapine (3mg/kg, n= 3), the levels of BF-1 mRNA in the rostral striatum were unchanged compared to the levels expressed in saline treated control animals. [Figure 4.6.1]. However, in the rat caudal striatum, it was observed that intraperitoneal injection of clozapine (3mg/kg, n= 3) caused an increase in the levels of BF-1 mRNA after 24 hours. This induction by clozapine was shown to be significantly different from saline treated animals [$F(3,8) = 4.93, p = 0.038, p < 0.05$] [Figure 4.6.2]. Haloperidol (1mg/kg, n= 3) and fluphenazine (3mg/kg, n= 3) had no effect on BF-1 mRNA levels in the caudal striatum 24 hours after drug administration. [Figure 4.6.2].

4.7 Effect of acute neuroleptic drug treatment on the expression of the POU-domain containing gene RHS2 in the rat striatum

Low levels of RHS2 mRNA were expressed in both the rostral and caudal striatum, in saline treated rats. Treatment with haloperidol (1mg/kg, n= 3), fluphenazine (3mg/kg, n= 3) or clozapine (3mg/kg, n= 3) had no significant effect on the levels of RHS2 mRNA in the rostral striatum of the rat [Figure 4.7.1]. Similarly in the rat caudal striatum, treatment with haloperidol (1mg/kg, n= 3), fluphenazine (3mg/kg, n= 3) or with clozapine (3mg/kg, n= 3) caused no

Figure 4.6.1 : Effect of acute neuroleptic drug treatment on the expression of BF-1 mRNA in the rat rostral striatum. Animals were injected i.p. with vehicle (0.9% saline), or haloperidol (1mg/kg), or fluphenazine (3mg/kg), or with clozapine (3mg/kg). After 24 hours, the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m.; the number of animals per group is shown in parentheses.

Figure 4.6.2: Effect of acute neuroleptic drug treatment on the expression of BF-1 mRNA in the rat caudal striatum. Animals were injected i.p. with vehicle (0.9% saline), or haloperidol (1mg/kg), or fluphenazine (3mg/kg), or with clozapine (3mg/kg). After 24 hours, the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m.; the number of animals per group is shown in parentheses. Statistical significance was determined by one-way ANOVA, followed by post-hoc Dunnett's test. $p < 0.05^*$ relative to saline group.

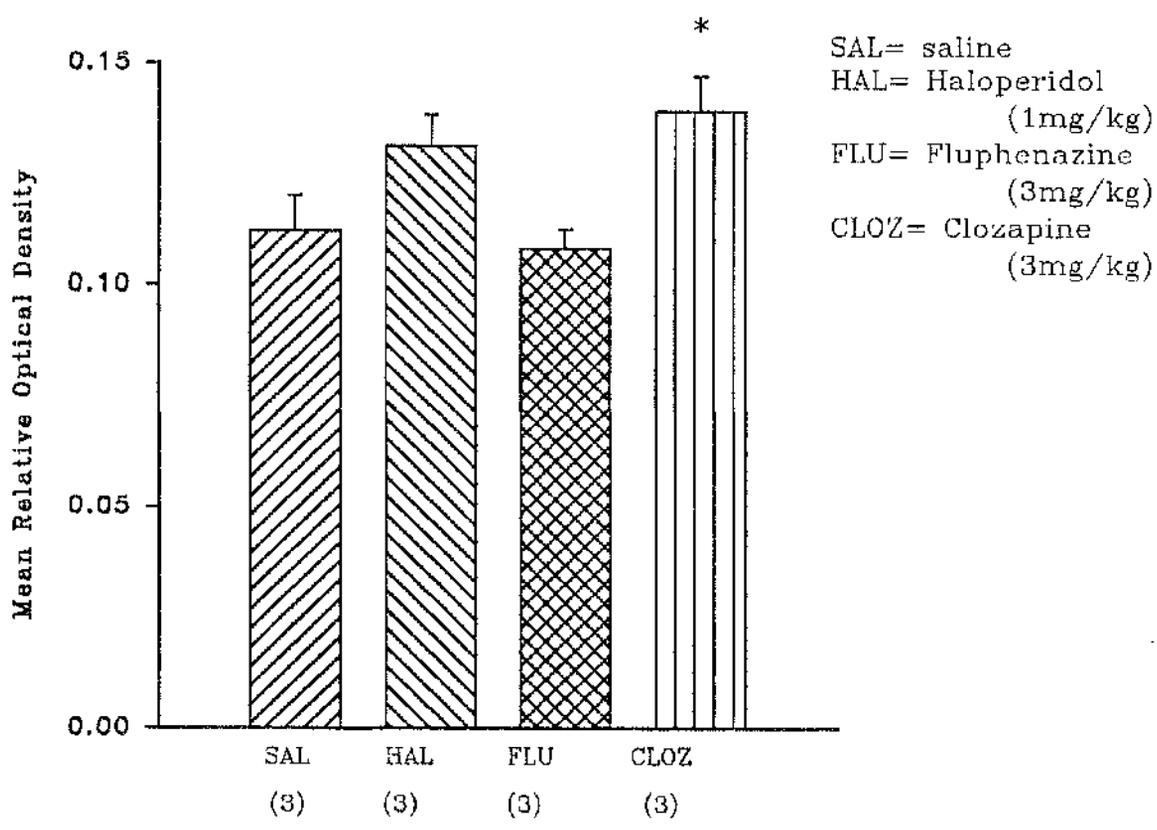
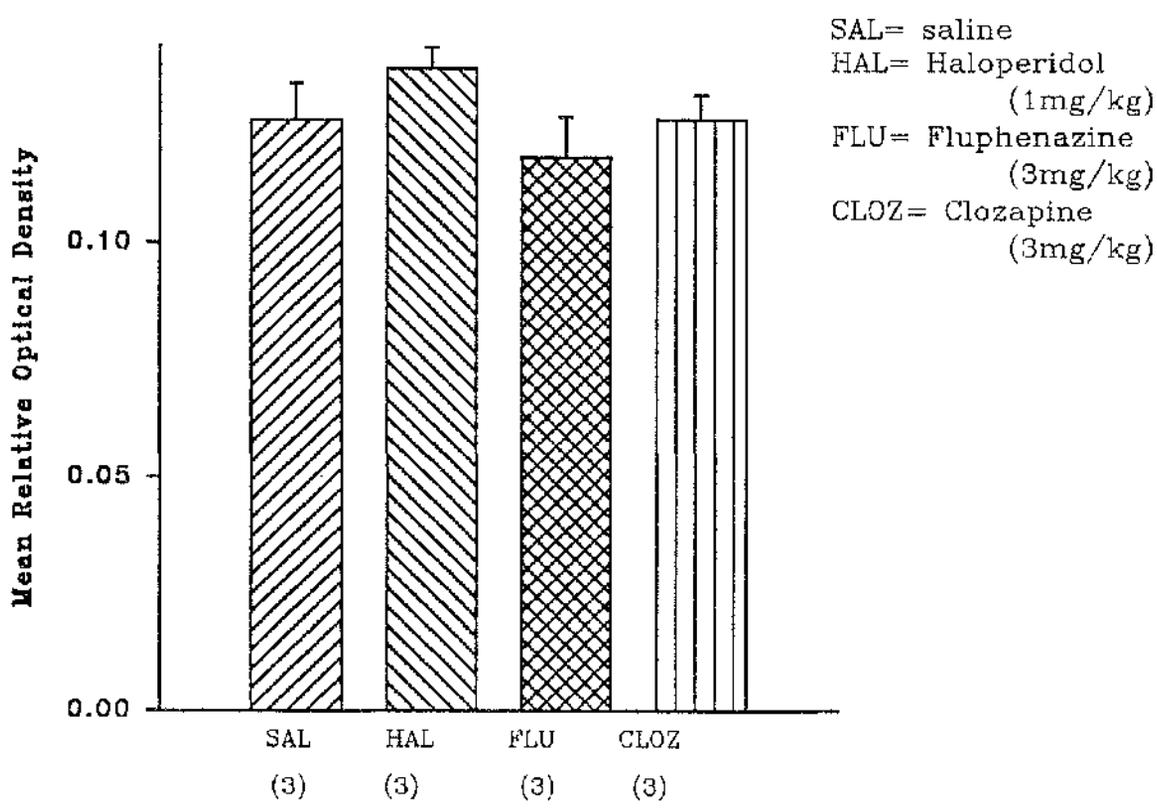
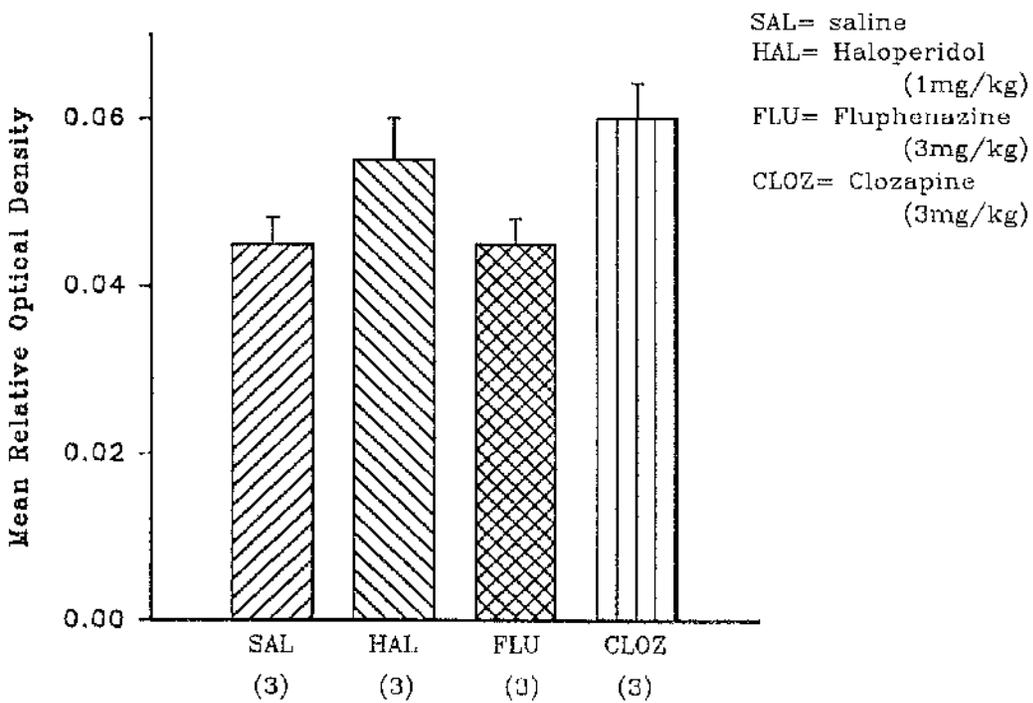
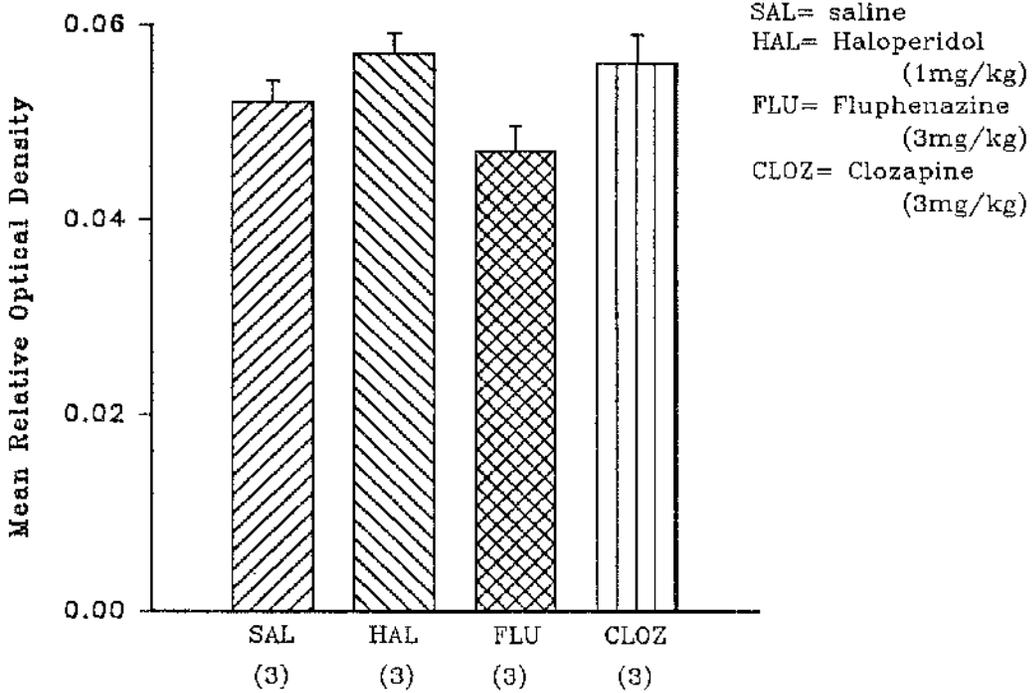


Figure 4.7.1 : Effect of acute neuroleptic drug treatment on the expression of RGS2 mRNA in the rat rostral striatum. Animals were injected i.p. with vehicle (0.9% saline), or haloperidol (1mg/kg), or fluphenazine (3mg/kg), or with clozapine (3mg/kg). After 24 hours, the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m.; the number of animals per group is shown in parentheses.

Figure 4.7.2 : Effect of acute neuroleptic drug treatment on the expression of RGS2 mRNA in the rat caudal striatum. Animals were injected i.p. with vehicle (0.9% saline), or haloperidol (1mg/kg), or fluphenazine (3mg/kg), or with clozapine (3mg/kg). After 24 hours, the animals were killed and their brains processed for in situ hybridisation. Results, obtained from autoradiographic film, are expressed as mean relative optical density + s.e.m.; the number of animals per group is shown in parentheses.



significant changes in the levels of RHS2 mRNA compared to saline treated animals in the rat [Figure 4.7.2].

DISCUSSION

Regulation of Immediate Early Genes in vivo

4.8 Time Course of fluphenazine effect on c-fos and zif/268 mRNA levels

The typical neuroleptic drug fluphenazine, having a high affinity for dopaminergic D2 receptors, was shown to maximally induce c-fos mRNA levels in the rat striatum forty-five minutes after the drug was given, and the levels were reduced back to basal levels after three hours. This finding is similar to previous reports that suggest a peak induction of c-fos can occur from thirty to sixty minutes after an acute treatment with psychostimulants or neuroleptic drugs (Robertson *et al.*, 1989; Cole *et al.*, 1992; Nguyen *et al.*, 1992; Bhat *et al.*, 1992). Our results also showed a significant increase in c-fos mRNA after fifteen minutes in the rostral region of the rat striatum, but not in the caudal regions. In a former study, zif/268 mRNA levels have been shown to be significantly increased after fifteen minutes (Cole *et al.*, 1992), and therefore the observed induction of c-fos mRNA fifteen minutes after fluphenazine treatment is not unlikely. It may be that the drug fluphenazine is absorbed more quickly, and thus an induction of c-fos gene expression is occurs more rapidly. Differences in the rate at which drugs are absorbed across the blood-brain barrier are likely to reflect small differences in peak expression time.

The immediate early gene zif/268 was also demonstrated to be induced maximally in the caudal region of the striatum after thirty minutes, and the levels declined back to basal levels after three hours. In the rostral regions of the rat striatum maximal zif/268 expression was observed after forty-five minutes which then decreased back to basal levels over a three hours time period. In previous reports it has been shown that zif/268 is induced maximally by thirty to forty-five minutes after neuroleptics (Cole *et al.*, 1992; Nguyen *et*

al., 1992), although an induction can be measured as early as fifteen minutes (Cole et al., 1992). Our results are therefore consistent with previous studies (Cole et al., 1992; Nguyen et al., 1992; Bhat et al., 1993).

These studies were undertaken in order to reproduce the current evidence suggesting that 'typical' neuroleptics can induce both c-fos and zif/268 gene expression in the rat striatum and accumbens regions. In addition we were interested in the effects of certain 'typical' and 'atypical' antipsychotic drugs on the expression of two members of the jun family of immediate early genes, namely jun-B and jun-D, and to elucidate the receptor mechanisms that precede the induction of these immediate early genes.

4.9 Effect of acute Typical Neuroleptic drug treatment on c-fos and zif/268 gene expression in the rat striatum and nucleus accumbens

The results of this study show that acute injections of the typical antipsychotic drugs haloperidol and fluphenazine can cause the induction of a number of immediate early genes including c-fos and zif/268 in the striatal region of the rat brain. A considerable induction of zif/268 mRNA, and to a lesser extent c-fos mRNA, was observed in the striatum. The induction of both c-fos and zif/268 mRNA by these typical antipsychotics followed a dorsolateral pattern. In addition, haloperidol was also shown to induce c-fos mRNA levels homogeneously in the nucleus accumbens. This is in agreement with previous reports suggesting that haloperidol can induce fos protein in both the core and shell regions, as well as in the striatum (Deutch et al., 1992; Robertson and Fibiger, 1992). However, the level of c-fos mRNA was considerably less in the accumbens region after fluphenazine, compared with the levels seen in the striatum after fluphenazine administration. Merchant and Dorsa (1993),

observed no induction of c-fos mRNA in the nucleus accumbens after administration of typical antipsychotic agents. Conversely, zif/268 mRNA was dramatically induced by fluphenazine in the nucleus accumbens, the effect being quantitatively greater than that of haloperidol. It is clear from this that the immediate early genes under investigation have different induction patterns after typical neuroleptic drug administration. This may reflect stimulation of different cell populations within the caudate-putamen. The inductive effect of haloperidol on both c-fos and zif/268 is consistent with previous reports demonstrating an increase in the expression of these immediate early genes by haloperidol in the caudate-putamen (Dragunow et al., 1990; Miller, 1990; Nguyen et al., 1992; Robertson et al., 1992).

As many antipsychotic drugs are known to be potent dopamine receptor antagonists, it has been suggested that the induction of certain immediate early genes such as c-fos and zif/268 may be mediated by blockade of dopamine neurotransmission.

More recent evidence also indicates that the ^ulate response gene[»] dynorphin acting through kappa opioid receptors can inhibit the cocaine-induced expression of c-fos, mediated by stimulation of D1 receptors, in striatonigral neurons (Steiner and Gerfen, 1995). This suggests that the striatal dynorphin system acts to inhibit the dopamine input to striatonigral neurons, implying that endogenous feedback mechanisms can control the level of the immediate early gene, c-fos, that is expressed. This reveals a further complexity to cellular events, and in the future may provide an insight into the manifestation of certain disease states.

4.10 Effect of acute Typical Neuroleptic drug treatments on jun-B and jun-D gene expression in the rat striatum and nucleus accumbens

We report here for the first time that haloperidol can dramatically induce the expression of jun-B mRNA in the rat striatum and nucleus accumbens. The level of expression of jun-B mRNA in the striatum was similar to the levels expressed for c-fos mRNA and zif/268 mRNA after administration of haloperidol, compared with the very low levels expressed in saline-treated control animals. The large proportion of cells positively labelled for jun-B in the striatum suggest that jun-B is induced primarily in the medium-sized spiny neurons, because this cell type is known to represent 95% of striatal neurons (Graybiel *et al.*, 1986). The induction due to haloperidol followed a rostral-caudal decline that matches the proposed rostral-caudal decline in the density of D2 dopamine receptors in the striatum (Boysen *et al.*, 1986). Therefore, this evidence suggests that the induction of jun-B is predominantly modulated by D2 dopamine receptors.

Whereas this is the first demonstration that jun-B is induced by neuroleptic treatment, there is some evidence for the induction of jun-B after acute administration of cocaine (Hope *et al.*, 1992), amphetamine, and apomorphine (Cole *et al.*, 1992). It is well known that these drugs have agonist properties at a number of receptors, including dopamine receptors. Therefore our evidence suggests that the expression of jun-B is regulated according to the level of dopamine activity.

This study has also revealed another member of the jun family of immediate early genes to be induced by typical neuroleptic drug administration. After haloperidol, a small but significant induction of jun-D mRNA was observed in both the striatum and nucleus accumbens of the rat. High basal

levels of jun-D mRNA exist throughout the brain, which makes this immediate early gene different from jun-B. Similar to jun-D, high levels of basal expression of c-jun are also present in certain brain areas. This is the first evidence that the level of expression of the jun-D gene can be modulated by neurotransmitter activity. The induction of jun-D after haloperidol administration suggests that jun-D, as well as jun-B, may be interacting with fos proteins to regulate the expression of downstream genes. In a manner similar to jun-B, expression of jun-D mRNA followed a dorsolateral pattern of distribution in the striatum and a homogeneous distribution in the nucleus accumbens after haloperidol administration.

The effect of fluphenazine, also having a typical antipsychotic drug profile, was clearly very similar to that of haloperidol. Fluphenazine, given at relatively high doses was shown to induce jun-B mRNA. A dramatic induction of jun-B expression was observed in the striatum and a similar level of induction was observed for jun-B mRNA in the nucleus accumbens after fluphenazine. As typical neuroleptics are potent dopamine receptor antagonists in vivo, this further suggests that blockade of dopamine transmission may be involved in mediating this effect.

4.11 Effect of the Atypical neuroleptic, clozapine on the expression on the immediate early genes c-fos, zif/268, jun-B and jun-D and the possible receptor mechanism(s) mediating immediate early gene induction by antipsychotic drugs

The results obtained demonstrate that low doses of clozapine have no effect on the expression of c-fos, zif/268, jun-B, or jun-D mRNA levels in either the rat striatum or nucleus accumbens after acute administration.

Previous reports have demonstrated that higher doses of the atypical antipsychotic clozapine can induce fos-immunoreactivity in the shell region of the nucleus accumbens, but not in the striatum (Deutch *et al.*, 1992; Robertson

and Fibiger, 1992). It is currently thought that the accumbens is associated with the mesolimbic system and is one of the prime targets for neuroleptic drugs. Behavioural studies have shown that chronic clozapine treatment selectively enhances apomorphine-induced locomotor activity, behaviour associated with dopamine mesolimbic activation. Whereas, treatment with haloperidol significantly enhanced apomorphine-induced chewing and sniffing stereotypies, thought to be associated with dopamine nigrostriatal activation (Seeger *et al.*, 1982). The extrapyramidal side effects associated with many typical neuroleptic drug treatments may be attributed to their activity on motor systems within the caudate-putamen. Therefore, the pattern of expression of c-fos and other immediate early genes may provide an important insight into the regional mechanisms of atypical antipsychotics. In this study low doses of the atypical antipsychotic drug clozapine had no effect on the levels of expression of c-fos, zif/268, jun-B and jun-D mRNA in either the striatum or nucleus accumbens of the rat brain. It has previously been shown through in vitro binding studies that fluphenazine has a greater affinity for the D2 dopamine receptor than the D4 dopamine receptor (Van Tol *et al.*, 1991). Clozapine, however, has a ten-fold greater affinity for the D4 site compared with the D2 site (Van Tol *et al.*, 1991). Clozapine has been shown to bind appreciably to striatal D2 sites in rats only at doses of 10mg/kg and above (Saller *et al.*, 1987; Audinot *et al.*, 1993). It is therefore likely that the lower dose of 3mg/kg used in these acute experiments is preferentially blocking D4 sites in vivo rather than D2 sites. This is supported by the evidence that a dose of 3mg/kg clozapine blocks a substantial proportion of the serotonin 5-HT₂ receptors in vivo (Canton *et al.*, 1993). Indeed clozapine has a similar affinity for D4 and 5-HT₂ sites (Van Tol *et al.*, 1991; Zifa and Fillion, 1992), implying that this dose will produce D4 antagonism in vivo. At higher doses of clozapine (10-20mg/kg), an induction of zif/268 has been reported in the caudate-putamen (Nguyen *et al.*, 1992). This therefore suggests that at higher doses, clozapine is acting at D2 receptors to

produce this effect. As clozapine had no effect on the immediate early genes under investigation, this would imply that the increased expression observed after typical neuroleptic drug administration is most likely due to a D2 dopamine receptor blockade rather than a D4 effect.

D1 dopamine receptor blockade may be considered to have negligible contribution to the effect on immediate early gene expression by any of the neuroleptics used in these studies. It has been shown that low doses of fluphenazine occupy primarily D2 dopamine receptors *in vivo* (Saller *et al.*, 1987). Dragunow *et al.* (1990), have demonstrated previously that the specific D1 dopamine receptor antagonist SCH23390 did not induce *c-fos* protein in striatal neurons. Therefore, it is likely that the effect of haloperidol, fluphenazine or clozapine on D1 or D5 dopamine receptors does not contribute to the clear induction of the immediate early genes *c-fos*, *zif/268*, *jun-B*, and *jun-D* seen after drug administration. However, the affinity of fluphenazine for D1 receptors is relatively high; therefore, it may be considered possible that the induction of *jun-B* mRNA after fluphenazine treatment is reduced in comparison with the induction observed after haloperidol, due to the ability of fluphenazine to antagonize D1 receptors.

Despite their high affinity for dopamine receptors, neuroleptics also have considerable affinity for many serotonin receptors. From both *in vivo* and *in vitro* binding studies, clozapine is known to be a potent antagonist at serotonin 5-HT₂ and 5-HT_{1C} receptors (Zifa and Fillion, 1992; Canton *et al.*, 1993). It is likely that clozapine administered at a dose of 3mg/kg would be blocking approximately 50% of 5-HT_{2/1C} receptors (Canton *et al.*, 1993). Likewise the typical neuroleptics haloperidol and fluphenazine were shown to have some antagonist binding properties at 5-HT₂ receptors, but with much lower affinity than clozapine. The relative potencies of these drugs, together with a lack of effect of clozapine at a dose likely to be blocking 5-HT_{2/1C} receptors *in vivo*, would therefore suggest that 5-HT receptors are not

involved in the effects reported here. Nevertheless, the consequences of neuroleptics having simultaneous effects on multiple transmitter systems must be taken into consideration.

A number of mechanisms have been proposed that could explain the increased expression of many immediate early genes by typical neuroleptic drug administration. One possible interpretation of these results is that haloperidol and fluphenazine produce their effects by increasing the release of dopamine from nigral-striatal terminals, which then acts on D1 dopamine receptors to induce immediate early genes. Related to this idea is the more likely possibility that D2 dopamine antagonists induce immediate early gene mRNA in striatal neurons by allowing unopposed D1 receptor stimulation. After a 6-hydroxydopamine lesion, the specific D1 dopamine receptor agonist SKF 38393 has been shown to increase c-fos in striatal neurons of the rat brain (Robertson *et al.*, 1990). The link between D1 receptor activation and the induction of immediate early genes, has been further supported by Cole *et al.* (1992), who have demonstrated that administration of SKF 38393, but not the specific D2 receptor agonist LY171,555, causes an induction of c-fos, zif/268 and jun-B in the caudate-putamen after a 6-hydroxydopamine lesion, and this can be abolished by pretreatment with SCH 23390. Recent evidence also suggests that SCH 23390 decreases the mRNA levels of zif/268 in the adult intact striatum of the mouse (Mailleux *et al.*, 1992). Therefore the current evidence implies strongly that the induction of a number of immediate early genes including c-fos, zif/268 and jun-B may be the result of a stimulation of the D1 dopamine receptor. Previous reports have also demonstrated that destruction of mesotelencephalic dopaminergic neurons with 6-hydroxydopamine abolishes the increase in fos expression in the striatum and nucleus accumbens produced by haloperidol (Robertson and Fibiger, 1992). This provides further support for the induction of c-fos after D1 dopamine receptor activation. However, it should be noted that SCH23390 does not

block the induction of c-fos by D2 receptor antagonism (Robertson *et al.*, 1992; Merchant and Dorsa, 1993). Therefore the precise mechanisms operating remain unclear.

Our results suggest that neuroleptic drug administration induce a pattern of immediate early gene induction more complex than has been appreciated until now. This includes the induction, at the mRNA level, of two members of the jun family of leucine zipper proteins, jun-B and jun-D. The results also suggest that these effects occur in response to blockade of D2 rather than D4 receptors. It would appear that a fine balance exists in vivo between D1 and D2 receptor stimulation. When the balance is disturbed, as in psychotic disorders, the expression of a number of immediate early genes may be affected. As yet, the consequences of these transient changes on the expression of downstream genes is not fully understood. However, a link has been suggested between c-fos and proenkephalin gene regulation (Sonnenberg *et al.*, 1989). Although, more recently, this idea has been argued against, as it has been implied that it is not the fos protein that interacts with the enkephalin enhancer (Konradi *et al.*, 1993). As the protein CREB has also been shown to act as a transcription factor, this may be a more likely candidate.

4.12 Effect of acute neuroleptic drug treatment on the expression of BF-1 and RHS2 transcription factor genes

Our results show that the expression of the recently cloned transcription factor genes RHS2 and brain factor-1 (BF-1) were not affected thirty minutes after treatment with either the typical neuroleptic drug fluphenazine, or the atypical neuroleptic drug clozapine, compared to saline treated controls. However, the expression of BF-1 mRNA after twenty-four hours was significantly induced by low doses of the atypical neuroleptic clozapine in the

medial striatum, but not by fluphenazine. There was a tendency towards an induction of RHS2 mRNA in the striatum after clozapine treatment, although the levels failed to reach significance. These results provide the first evidence that neuroleptics can induce BF-1 gene expression, at the mRNA level, in the adult rat brain.

In the developing mammalian brain, three separate regions known as the forebrain, midbrain and hindbrain are initially formed from the rostral part of the neural tube. The most rostral forebrain region, the telencephalon, gives rise to the structures of the cerebral hemispheres, which include the cerebral cortex and basal ganglia. Previous studies have demonstrated that the expression of BF-1, a member of the HNF-3/fork head gene family, is restricted to the telencephalon region in the developing rat brain (Tao and Lai, 1992). Therefore it has been suggested that BF-1 may have a role as a transcriptional regulator in the complex signalling system that coordinates the precise temporal and spatial patterns of differential gene expression in multicellular organisms. Northern blotting analysis has shown that the BF-1 gene is expressed in the cortex, olfactory bulb, hippocampus and caudate-putamen in the adult rat brain, the levels of expression being four-fold lower than in the fetal rat brain (Tao and Lai, 1992). More recently, the identification of a gene in human fetal tissue having high homology at the nucleotide level to brain factor-1 in the rat, has been shown to be expressed in the telencephalon, with strong expression being observed in the dentate gyrus and hippocampus. The expression of human BF-1, also referred to as HFK1, was also found to be restricted to neuronal cells in the telencephalon (Murphy *et al.*, 1994). The basal expression of BF-1 mRNA in the adult rat striatum showed that a large proportion of cells were positively labelled for BF-1, which suggests that BF-1 is expressed primarily in the medium-spiny neurons, because this cell type has previously been demonstrated to represent 95% of striatal neurons (Graybiel *et al.*, 1986). Therefore, it may

be suggested that the induction of BF-1 mRNA seen after clozapine treatment, is primarily in the medium-spiny neuronal cell population within the striatum.

The low dose of clozapine used in this study may reflect a mechanism involving either a dopamine D4 receptor stimulation or a 5-HT receptor subtype stimulation, which has previously been discussed in earlier studies using the same dosage of clozapine. Although clozapine is known to have a high affinity for many receptor types, including adrenergic, histaminergic and muscarinic receptors and therefore a variety of different receptors may be contributing to the observed induction of BF-1 after low doses of clozapine.

This is the first evidence that the a member of the HNF-3/ fork head gene family, brain factor-1, can be modulated by neuroleptic drug administration. Since the atypical neuroleptic, clozapine, but not the typical neuroleptic fluphenazine was able to induce BF-1 mRNA in the striatum, this may reflect a previously undiscovered role for BF-1 in the control of antipsychotic disorders, and a possible contributory factor in the lack of extrapyramidal side effects observed clinically after clozapine treatment. However, the potential target genes, whose level of expression is dependent on BF-1 induction, remains to be identified.

Chapter 5

Striatal Neuronal Cultures

INTRODUCTION

5.0 Striatal Neurone Morphology

As our understanding of the molecular mechanisms regulating cellular activity has grown, the information that can be obtained from *in vivo* experiments has become more difficult to interpret, and the importance of *in vitro* systems has consequently increased. The intercellular and subcellular processes which coordinate and control the activity of neurones in particular brain regions such as the striatum can frequently only be studied in tissue maintained *in vitro*, where reagents can be applied in known concentrations, directly onto the cells. This has been emphasised by the recent identification of novel dopamine and serotonin receptors which are expressed at relatively high levels in striatal tissue (Sunahara *et al.*, 1991; Van Tol *et al.*, 1991; Monsma *et al.*, 1993). The absence of selective ligands for these receptors means that their role in regulating the activity of striatal neurons can essentially only be studied *in vitro*, where the drugs can be applied in precise concentrations, and hence relative selectivity can be calculated exactly.

It has been clear for some years that striatal neurons, dissociated from embryonic rat brain and maintained under cell culture conditions, can survive well *in vitro* for a number of weeks (Messer, 1981; Kessler, 1986; Weiss *et al.*, 1986; Surmeier *et al.*, 1988). These primary striatal neuron cultures have proved to be a useful system for electrophysiological and second messenger studies, looking at the effects of stimulation of receptors for dopamine, GABA, glutamate and growth factors (Murphy *et al.*, 1987; Dumuis *et al.*, 1988; Dubinsky, 1989; Misgeld and Dietzel, 1989; Sproson and Woodruff, 1990; Maria *et al.*, 1992; Simpson and Morris, 1994). However, there is evidence that dopamine is likely to have different effects on the different classes of neuron in the striatum *in vivo* (Morris *et al.*, 1988; Le Moine *et al.*, 1991; Pollack and

Wooten, 1992). It is clearly of considerable importance to know which of the neuronal types normally present in the striatum are also present in the primary cultures. It may be that some classes of neuron do not survive at all, and that the intracellular effects that can be studied are those occurring in only one or two different neuronal types. Conversely, all classes of neuronal types may be present *in vitro*, in which case these primary cultures may provide a suitable model for studying the regulation of activity of the whole range of striatal neurons.

Approximately 95% of striatal neurons *in vivo* are thought to be GABA-ergic projection neurons additionally containing either enkephalin peptides or dynorphin and tachykinin peptides (Bolam *et al.*, 1985; Graybiel, 1990; Besson *et al.*, 1990). The remainder comprise possibly three populations of interneurons releasing acetylcholine, or GABA and neuropeptide Y, or somatostatin, neuropeptide Y and nitric oxide (Bolam *et al.*, 1984; Phelps *et al.*, 1985; Graybiel, 1990; Vuillet *et al.*, 1990). Despite the importance of a comprehensive phenotypical characterisation of striatal neurons in culture, there is to date only limited evidence as to what classes of neurons are present *in vitro*. Extracts of primary striatal cultures have previously been shown to contain immunoreactive somatostatin and substance P (Kessler, 1986), and proenkephalin mRNA (Schwartz and Simantov, 1988; Vilijn *et al.*, 1988), while analysis at the cellular level has shown the presence of GABA (Messer, 1981; Weiss *et al.*, 1986; Surmeier *et al.*, 1988), along with immunoreactive leu-enkephalin (Surmeier *et al.*, 1988), which could arise from either the expression of the prodynorphin gene, or the proenkephalin gene, and immunoreactive somatostatin (Williams *et al.*, 1991). Of these, only GABA, immunoreactive leu-enkephalin and immunoreactive somatostatin have been localised to neurons rather than glia (Surmeier *et al.*, 1988; Williams *et al.*, 1991). Striatal glia in culture are known to express the enkephalin gene at high levels (Vilijn *et*

al., 1988), so it remains an open question to what extent the other neuroactive substances detected in culture extracts are derived from non-neuronal cells.

Therefore, before striatal neurons in culture can be used as a model for the in vivo situation, it is important to characterise the cell types that are present in vitro.

RESULTS

5.1 Characterisation of rat striatal neuronal cultures

In striatal neurone cultures grown in DMEM, a high proportion of cells were identified as astrocytes on the basis of their large, flat morphology and their characteristic immunoreactivity for glial fibrillary acidic protein (GFAP), revealing large unstained nuclei, as shown in figure 5.1.1. To reduce astrocytic contamination of the cultures, a range of different concentrations of cytosine arabinoside was added to the culture medium. It was found that cytosine arabinoside ($20\mu\text{M}$), optimally reduced astrocytic contamination of the cultures to less than 5% of the total cell population without dramatically affecting the numbers of neurones. Cytosine arabinoside treatment was therefore included in all subsequent experiments. Neuronal cells were clearly distinguishable from astrocytes on the basis of their morphology. However, their identity was confirmed by using neurone-specific enolase (NSE)- immunostaining, together with in situ hybridisation with an oligonucleotide probe specific for microtubule-associated protein 2 (MAP 2) mRNA, found only in neuronal cell types. A variety of neuronal morphologies could be identified in neurones stained with the anti-NSE antiserum, including bipolar cells with oval somata, multipolar cells with oval somata, and cells with triangular somata giving rise to large numbers of dendritic processes. [Figure 5.1.2] revealing the diversity of cell types within the striatum. Following in situ hybridisation with an oligonucleotide probe for MAP 2 mRNA, neuronal cells counterstained with cresyl violet, can be seen to express MAP 2 message, compared to the complete absence of MAP 2 mRNA following hybridisation of cultures with a MAP 2 sense oligonucleotide probe (not shown).

A large number of cells in culture, approximately 30-40%, were stained using the met-enkephalin antiserum [Figure 5.1.3], characteristically staining the cell soma. Cells from each of the morphological groups were identified as met-enkephalin immunoreactive neurons. In accordance with this result, a relatively high proportion of neurones in culture (30-40%), were found to contain proenkephalin (Penk) mRNA, when assayed by in situ hybridisation histochemistry [Figure 5.1.4]. After hybridisation with a 25-fold excess of unlabelled Penk oligonucleotide included in the hybridisation buffer, no labelling was observed, confirming the specificity of the hybridisation reaction. The cells containing immunoreactive-met-enkephalin or those expressing Penk mRNA were generally of medium size (12-20 μ m diameter). In comparison, cortical cultures grown under the same conditions contained no met-enkephalin immunoreactive neurons (not shown).

A smaller number of neurones were also found to express preprotachykinin (PPT) mRNA [Figure 5.1.5], although the hybridisation signal was much weaker than with the Penk probe. No specific signal could be detected with an oligonucleotide probe specific for prodynorphin mRNA at any time point from 7 DIV to 21 DIV.

Following hybridisation of striatal cultures with the somatostatin antisense probe, only a small proportion of neurones were labelled, but the intensity of the hybridisation signal was very strong. [Figure 5.1.6]. Cells labelled with the somatostatin probe were generally larger (15-25 μ m), than those labelled for Penk or PPT. Again, the hybridisation signal was completely removed by the inclusion of a 25-fold excess of unlabelled somatostatin oligonucleotide in the hybridisation reaction. No specific signal was seen when sense probes were used under identical conditions [Figure 5.1.7]. The NPY probe was expressed in the same number of cells as the somatostatin probe, although the hybridisation signal was considerably weaker than that observed for the somatostatin probe [Figure 5.1.8], and similarly a 25-fold excess of

unlabelled probe when added to the hybridisation reaction, abolished the hybridisation signal.

A very small number of striatal neurons in culture also stained positively for acetylcholinesterase [Figure 5.1.9]. These cells tended to be larger than average (20-25 μ m diameter) and stained very darkly for acetylcholinesterase. Characteristically, these cells gave rise to a number of heavily-stained processes that appeared to make contact with many neighbouring cells.

A relatively small percentage (< 5%) of cultured striatal neurons were demonstrated to contain NADPH-diaphorase activity [Figure 5.1.10]. These cell types generally had large triangular or oval somata (20-25 μ m diameter), and appeared similar in terms of their gross morphology to the neurones exhibiting NADPH-diaphorase activity in sections of adult rat striatum [Figure 5.1.11]. The cultured NADPH-diaphorase-positive neurones gave rise to long processes with axon-like morphology, with an extensive plexus of connections with neighbouring cells. These "axons" were characterised by the presence of pronounced varicosity-like swellings along their length, which loosely resembles the appearance of the diaphorase-positive fibres in adult rat striatal tissue.

Figure 5.1.1 : Primary striatal cultures ,in the absence of cytosine arabinoside, processed for immunocytochemistry using antisera against glial fibrillary acidic protein (GFAP). The stained cells are large and flat with unstained nuclei. Scale bar represents 25 μ m.



Figure 5.1.2: Primary striatal cultures processed for immunocytochemistry using antisera against neurone specific enolase (NSE). Groups of neurones show strong intensity staining for NSE, including a variety of different morphological types (bipolar, multipolar, triangular). Scale bar represents 25 μ m.



Figure 5.1.3 : Primary rat striatal cultures processed for immunocytochemistry using antisera against met-enkephalin. The cell bodies of neurones in culture stained positively for met-enkephalin. The characteristics of enkephalin positive neurones tended to be those with oval somata and a bipolar morphology. Scale bar represents 25 μ m.

Figure 5.1.4 : Cultured striatal neurones following hybridisation with a probe specific for proenkephalin (Penk) mRNA. Exposure time 2 weeks. Photomicrographs show cultured striatal neurones expressing the Penk gene, as seen by the distribution of silver grains over certain cell bodies. Approximately 40-50% of cells were labelled positively with the Penk probe. Scale bar represents 25 μ m.



Figure 5.1.5 : Cultured rat striatal neurones following hybridisation with a radiolabelled probe specific for preprotachykinin (PPT) mRNA. Exposure time approximately 6 days. Photomicrographs show striatal neurones counterstained in cresyl violet. Approximately 25-35% of cells were labelled positively for PPT mRNA. Scale bar represents 25 μ m.

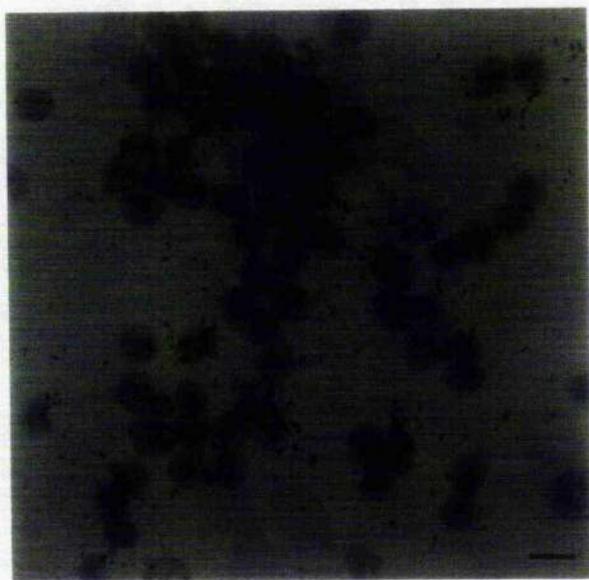


Figure 5.1.6 : Cultured rat striatal neurones following hybridisation with a radiolabelled probe specific for preprosomatostatin mRNA. Exposure time 4 days. Photomicrographs show striatal neurones counterstained in cresyl violet. A relatively small percentage of cells in culture were labelled positively for preprosomatostatin mRNA, as shown by an increase in the density of silver grains over certain cell bodies. Characteristically, the positively labelled cells were of larger size. Scale bar represents 25 μ m.

Figure 5.1.7 : Cultured rat striatal neurones following hybridisation with a radiolabelled somatostatin 'sense' probe. Exposure time 5 days. Photomicrographs show striatal neurones counterstained in cresyl violet. Scale bar represents 25 μ m.

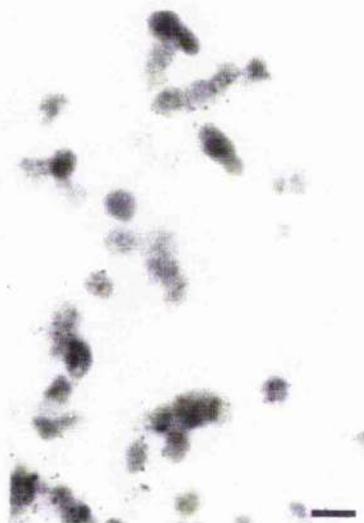
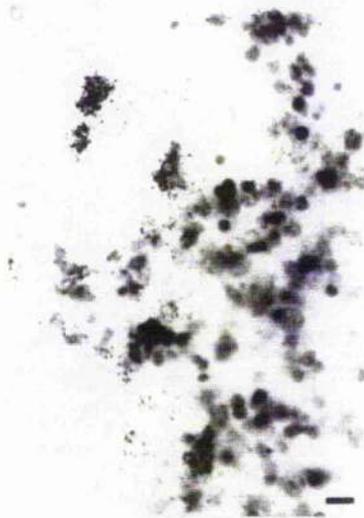


Figure 5.1.8 : Cultured rat striatal neurones following hybridisation with a radiolabelled probe specific for proneuropeptide Y mRNA. Exposure time 5 days. Photomicrographs show striatal neurones counterstained in cresyl violet. A small percentage of cells were labelled positively for proneuropeptide Y mRNA, as shown by an increase in the number of silver grains over certain cell bodies. However the intensity of labelling was generally much lower than for preprosomatostatin mRNA. Scale bar represents 25 μ m.

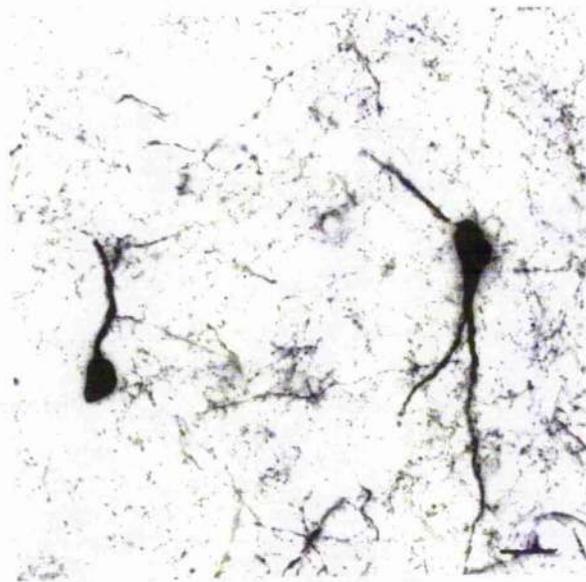
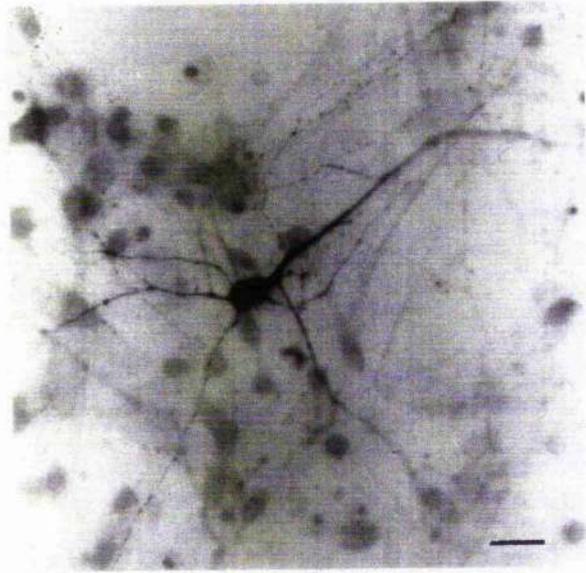


Figure 5.1.9 : Primary rat striatal neurone cultures processed for acetylcholinesterase staining. A small percentage of cells stained positively for acetylcholinesterase in culture, observed as a brown stain. Again both cell soma and dendrites were stained intensely and characteristically the labelled cells were much larger with many dendritic processes. Scale bar represents 25 μ m.



Figure 5.1.10 : Primary rat striatal neurone cultures processed for NADPH-diaphorase staining. Cultures were incubated with phosphate buffer containing NADPH and nitro blue tetrazolium for 1 hour at 37°C. A small percentage of cells in culture stained positively for NADPH-diaphorase, which can be seen here as a blue stain. Both the cell soma and dendrites stained positively. The population of cells that were NADPH-diaphorase positive tended to be large with numerous dendritic processes. Scale bar represents 25µm.

Figure 5.1.11 : Adult rat cryostat sections of caudal striatum processed for NADPH-diaphorase staining. Sections were incubated with phosphate buffer containing NADPH and nitro blue tetrazolium for 1 hour at 37°C. A small percentage of cells in adult rat sections stained positively for NADPH-diaphorase, seen here as a blue stain. Both cell soma and dendrites stained positively. Scale bar represents 25µm.



DISCUSSION

5.2 Phenotypic characterisation of rat striatal neurones in primary culture

Striatal neurones in primary culture have already proven to be a useful model system for studying the molecular events associated with striatal function. The use of primary cultures may be of particular importance with the rapid discovery of newly cloned receptors, many of which are known to be expressed in striatal tissue, such as the newly cloned dopaminergic D4 receptor (Van Tol *et al.*, 1991) and serotonergic 5-HT₆ receptor (Monsma *et al.*, 1993). In this study we have therefore examined primary striatal cultures for the presence of the various neurochemical phenotypes which are found in the rat striatum *in vivo*. The use of *in situ* hybridisation, rather than immunohistochemistry, allows the unequivocal identification of the pattern of gene expression in these cultured neurons.

The specificity of the hybridisation protocol is suggested in various pieces of evidence. We have shown repeatedly that in normal tissue, our hybridisation sequences allow the specific localization of the mRNA sequences to which the probes are complementary (Morris, 1989; Morris *et al.*, 1990; Sirinathsinghji *et al.*, 1990; Morris and Hunt, 1991; Morris, 1992). The high stringency conditions we employ prevent the hybridisation of the probes to any related sequences in the tissue. The lack of any signal in striatal cultures when sense probes are used, and the observation that the hybridisation signal with the antisense probes can be inhibited competitively by a 25-fold excess of the unlabelled oligonucleotide, provide further evidence that the hybridisation signals that we detect represent the authentic complementary mRNA.

Previous work has established that a high proportion of cultured striatal neurons are GABA-ergic (Messer, 1981; Weiss *et al.*, 1986; Surmeier *et al.*,

1988). This is consistent with the *in vivo* situation. The majority of these GABA-ergic neurons *in vivo* are thought to be projection neurons, where GABA coexists with either dynorphins and tachykinins, or enkephalins (Besson *et al.*, 1990). It could therefore be predicted that the cultures should contain a high percentage of neurons expressing the proenkephalin, preprotachykinin and prodynorphin genes.

In primary striatal neuron cultures, it was observed that many neurons stained positively with an antiserum raised against met-enkephalin and there was negligible cross-reactivity to any of the peptide products of the prodynorphin (Proenkephalin B) gene. Since these cells closely resemble those demonstrated by Surmeier *et al.*, (1988), to contain immunoreactive-leu-enkephalin, this suggests that these neurons express the proenkephalin rather than the prodynorphin gene. This was confirmed by *in situ* hybridisation analysis using a proenkephalin oligonucleotide. These results indicate that a high proportion of cells in these cultures express the proenkephalin phenotype.

An exact determination of the percentage of neurons expressing a particular phenotype *in vitro* is impossible. Although many neurons remain spread evenly over the substrate after seeding, other neurons migrate at an early stage to form regions where the neurons are clumped together, (Schwartz and Simantov, 1988; Surmeier *et al.*, 1988), and accurate counting cannot be performed. However, a rough estimate of the proportion of neurons containing either proenkephalin mRNA or immunoreactive-met-enkephalin in culture suggests that enkephalinergic cells are the predominant cell type *in vitro*, as *in vivo*, although the proportion is rather lower than the 60% of striatal neurons that are known to be enkephalinergic *in vivo*. On the basis of the number of cells involved, it is likely that these cultured neurons expressing the proenkephalin phenotype are identical to the GABA-ergic cells already identified in primary culture (Messer, 1981; Weiss *et al.*, 1986; Surmeier *et al.*, 1988).

A considerable number of neurons in culture were also found to express preprotachykinin mRNA, although the proportion appears to be slightly lower than the proportion which are enkephalinergic. Considering the impossibility of making exact determinations of the percentage of labelled neurons in primary striatal culture, this would be approximately consistent with the reports that roughly 40-50% of striatal neurons *in vivo* contain immunoreactive substance P (Besson *et al.*, 1990). It therefore seems clear that neuronal rather than non-neuronal cells are the source of immunoreactive-substance P detected in extracts of primary striatal cultures (Kessler, 1986).

In situ hybridisation is not a truly quantitative technique, in the sense that the kinetics and equilibrium position of the hybridisation reaction are unknown for each oligonucleotide. The absolute levels of a particular mRNA species cannot be determined, and so it is not possible to compare the abundance of different mRNA species, unless it is assumed that the hybridisation reaction has an identical efficiency for each probe/mRNA combination. This assumption is likely to be valid, but a formal proof would be essentially impossible. If this assumption is made, then the results suggest that the neuronal levels of preprotachykinin in cultured striatal neurons are markedly lower than the levels of preproenkephalin mRNA. This appears to be the situation *in vivo* (Young *et al.*, 1986; Sirinathsinghji *et al.*, 1990), although the relative difference between the hybridisation signals with the two probes is probably greater in the cultures. Removal of the dopaminergic innervation of the striatum *in vivo* increases the levels of proenkephalin mRNA and decreases the levels of protachykinin mRNA (Young *et al.*, 1986; Morris *et al.*, 1989). This potentially exaggerated difference (based on the above assumption) between the apparent levels of proenkephalin mRNA and preprotachykinin mRNA in culture may, if correct, be due to the absence of dopamine in the environment of the cultured neurons.

We found that no striatal neurons in culture contained detectable levels of prodynorphin mRNA. This is in agreement with an earlier study analysing extracts of primary cultures (Vilijn *et al.*, 1988). This is interesting, when it is considered that striatal tachykinins and dynorphins are thought to be located predominantly in the same population of neurons (Besson *et al.*, 1990). The presence of neurons expressing the preprotachykinin phenotype *in vitro* suggests that this class of neurons is able to survive under these conditions. It may be that some developmental signal, for example the secretion of a particular growth factor, is required for the appearance of the prodynorphin phenotype in the cells, and that this signal is not present in our culture medium. Alternatively, prodynorphin mRNA may be expressed, but at levels below the limit of sensitivity of the hybridisation reaction. However, this is unlikely, given that the protocol is highly sensitive (Morris, 1992).

The medium aspiny neurons which represent only a few percent of the total number of neurons in the normal striatum, are thought to contain somatostatin, neuropeptide Y and NADPH-diaphorase (Vincent and Johansson, 1983; Smith and Parent, 1986; Graybiel, 1990). We report here that neurons expressing the somatostatin gene are present in primary striatal cultures. We also observe many neurons that express the neuropeptide Y gene, and in so far as comparisons of the amounts of different mRNA species are possible, the neuropeptide Y gene appears to be expressed, at the mRNA level, at a lower level than the somatostatin gene. This parallels the situation in the normal striatum (Morris, 1989; 1992). Equally, within the limitations outlined above, it appears that considerably fewer neurons in culture contain somatostatin mRNA or neuropeptide Y mRNA, as compared to proenkephalin mRNA.

These results provide the first evidence that neurons in striatal culture also contain NADPH-diaphorase. While it has yet to be shown that somatostatin mRNA, neuropeptide Y mRNA and NADPH-diaphorase are all

contained in the same cell *in vitro*, our data demonstrate that the cultures contain neurons expressing each of these phenotypes. It now seems clear that the NADPH-diaphorase reaction detects the enzyme responsible for nitric oxide synthase (Hope *et al.*, 1991). Release of nitric oxide from primary striatal cultures following glutamate receptor stimulation has recently been described, (Marin *et al.*, 1992). Our results confirm that the origin of this release is likely to be neuronal. Nitric oxide is synthesised in the cytoplasm by the nitric oxide synthase/NADPH-diaphorase enzyme, and is then thought to diffuse randomly across membranes to effect any cells in the surrounding area. The evidence that nitric oxide is involved in a number of biochemical and physiological processes in the central nervous system, is increasing rapidly (Garthwaite, 1991).

The morphology of the NADPH-diaphorase-positive neurons appears to correspond surprisingly closely to the morphologies identifiable in intact tissue. The intense Golgi-like staining of the processes of the neurons containing NADPH-diaphorase enabled us to ascertain the extent of their connections with other cells in the culture wells. In every case, the diaphorase-positive neurons gave rise to a complex branching network of fibres covered in varicosity-like swellings. This reproduces closely the appearance of intact striatal tissue, which is a dense interwoven mesh of diaphorase-positive fibres with varicosities. We were able to observe these varicosities in contact with unstained neurons. For traditional neurotransmitters, evidence of synaptic specialisations, both pre- and post-synaptic, in electron micrographs, would be required for evidence of cell-cell communication in culture. However, the non-synaptic release of nitric oxide, and its ability to cross cellular membranes, means that a rough proximity of NADPH-diaphorase-containing fibres with another cell would be sufficient to allow intercellular communication of signals. These results therefore provide strong evidence that nitric oxide release from these cells may affect other neurons in culture.

The adult intact striatum also contains relatively rare cholinergic cells, larger than the other striatal neurons, and thought to be interneurons (Bolam *et al.*, 1984; Phelps *et al.*, 1985; Graybiel, 1990). In primary striatal culture we found that a very small percentage of striatal neurons stained heavily for acetylcholinesterase, which is an indication, although not an absolute proof that the cells are cholinergic. In other brain regions there are neurons which contain acetylcholinesterase, but no choline acetyl-transferase, and therefore do not synthesise the transmitter acetylcholine. However, in the striatum *in vivo*, it appears that acetylcholinesterase is expressed only in cholinergic neurons (Walker *et al.*, 1987). The use of acetylcholinesterase as a marker for cholinergic neurons has the advantage that, in contrast to choline acetyl-transferase-immunostaining, the entire neuritic structure of the neurons is revealed.

The acetylcholinesterase-positive neurons that we observed in culture were larger than most of the other neurons, consistent with the situation in the intact striatum. The entire neuropil in normal striatal tissue stains very heavily for acetylcholinesterase, suggesting that neurites containing acetylcholinesterase *in vivo* are virtually ubiquitous. It was noticeable that a small number of acetylcholinesterase-positive cells in culture displayed an intricate and widespread array of neurites, extending potentially to affect large areas of the culture well. It is therefore possible that this feature of the acetylcholinesterase-positive neurons *in vitro* mirrors an aspect of the functional cytoarchitecture of similar neurons *in vivo*.

The results shown here suggest that both NADPH-diaphorase-positive and the acetylcholinesterase-positive neurons make extensive connections with other neurons in culture. There is electrophysiological (Misgeld and Dietzel, 1989; Dubinsky, 1989) and biochemical (Weiss *et al.*, 1986) evidence to support the presence of synaptic interactions between cultured striatal neurons. It should therefore be remembered that any neurotransmitter effects being

studied in such cultures may occur trans-synaptically, as well as directly on the cells under investigation.

Overall, these results suggest that primary cultures of striatal neurons may provide a surprisingly good model of the intact striatum. Neurons expressing proenkephalin mRNA, protachykinin mRNA, somatostatin mRNA and neuropeptide Y mRNA are all present, along with neurons staining for NADPH-diaphorase and acetylcholinesterase. The only neurotransmitter gene that apparently fails to be expressed from the striatal repertoire is the prodynorphin gene. Therefore primary cultures are a relatively good model of the intact striatum and will provide a suitable model for the identification of factors regulating the expression of these genes in culture.

Chapter 6

Dopaminergic Regulation of Immediate Early Genes in vitro

INTRODUCTION

6.0 Dopaminergic Regulation of Immediate Early Genes In Vitro

Many cell types in the mature central nervous system respond to changes in their local neurochemical environment by altering their pattern of gene expression (Goodman, 1990; Morris, 1993). These alterations in gene expression then persist for some time, allowing the cells to sustain a long-term response to the original stimulus. Perturbation of the normal level of dopaminergic transmission in the rat striatum, either by the administration of neuroleptics, which antagonise the actions of dopamine, or by lesioning the nigrostriatal pathway, results in sustained changes in the levels of peptide neurotransmitter mRNAs (Bannon *et al.*, 1986; Merchant and Dorsa, 1993; Morris *et al.*, 1988; Morris and Hunt, 1991; Young *et al.*, 1986), associated with parallel changes in the rate of neuropeptide release (Lindfors *et al.*, 1989).

Immediate early genes encode proteins whose structures fall into a number of distinct classes. They are rapidly induced in response to a variety of different stimuli such as growth hormones, increased synaptic activity and drug administration, and their proteins are thought to bind to specific sites in the genome to modulate the rate of transcription of downstream genes (Morgan and Curran, 1987; Sheng and Greenberg, 1990). Induction of immediate early genes is likely to provide a link between events at the cell membrane and long term alterations in neuronal gene expression. The immediate early genes *c-fos* (Morgan and Curran, 1987) and *zif/268* (Millbrandt, 1987), also known as *erg1*, *Krox 24*, and *NGFIA*, are thought to be involved in functional plasticity in rat striatal neurons, in that they are induced *in vivo* by treatment with neuroleptic drugs (Dragunow *et al.*, 1990; Nguyen *et al.*, 1992; Robertson and

Fibiger, 1992; see also chapter 3). It is therefore of considerable interest to determine the mechanisms through which dopamine receptors are linked to the regulation of gene expression. A number of different dopamine receptors are known to be present in the striatum. The D1 and D5 (D1b) dopamine receptors correspond to the traditional D1-like receptors (Monsma *et al.*, 1990; Sunahara *et al.*, 1991; Tiberi *et al.*, 1991) in that they show high affinity for the agonist SKF38393 and the antagonist SCH23390, and low affinity for D2 "selective" drugs such as quinpirole and eticlopride. Both D1 and D5 receptors are positively coupled to adenylate cyclase. Therefore the stimulation of D1-like receptors results in an increase in intracellular cAMP levels, with the consequent activation of cAMP-dependent protein kinase (PKA). However, there is also evidence that D1-like receptors can be linked to a stimulation of phospholipase C and consequently protein kinase C (PKC) activation (Felder *et al.*, 1989; Mahan *et al.*, 1990; Frail *et al.*, 1993). The D2, D3, and D4 dopamine receptors correspond to the traditional D2-like receptor, in that they show relatively high affinity for the agonist quinpirole and the antagonist eticlopride, along with low affinity for SKF38393 and SCH23390. The D2 and D4 receptors are coupled to an inhibition of adenylate cyclase activity (Seeman and Van Tol, 1993).

At present there is very little evidence on the dopaminergic regulation of gene expression in striatal neurons maintained in cell culture, although there is some evidence for neurotransmitter regulated induction of immediate early genes in cell culture conditions. Stimulation of the N-methyl-D-aspartate receptor in primary cultures of cerebellar neurons has been shown to trigger the induction of c-fos and other immediate early genes (Szekely *et al.*, 1987; Szekely *et al.*, 1989; Szekely *et al.*, 1990). More recently it has been shown that glutamate can increase c-fos, c-jun, jun-B, and zif/268 mRNAs by binding to selective glutamatergic receptor subtypes in cultured cortical and striatal neurons, and this induction is thought to be mediated by the enzyme PKC

(Vaccarino *et al.*, 1992). Several neurotransmitters activate phosphoinositide hydrolysis in central nervous system neurons as the first step of a transmembrane signalling cascade that may lead to neuronal circuit modulation (Berridge *et al.*, 1982; Gonzales and Crews, 1984; Batty *et al.*, 1985; Schoepp and Johnson, 1988). Ambrosini and Meldolesi (1989) have shown that quisqualate and muscarinic receptor-induced phosphoinositide hydrolysis in primary cultures of striatal and hippocampal neurons involves differential mechanisms of activation. Activators of the cAMP second messenger pathway in primary cultures of neurons from rat cerebral cortex have also been shown to induce c-fos and other immediate early gene transcription factors directly (Vaccarino *et al.*, 1993). Indeed, the D1-dopaminergic agonist, SKF38393, administered for short time periods, has been demonstrated to increase mRNA levels for the immediate early genes c-fos, jun-B, and zif/268 in cortical neurons (Vaccarino *et al.*, 1993). Stimulation of glutamatergic NMDA receptors and metabotropic quisqualate receptors in cultured striatal neurons has previously been shown to increase the release of arachidonic acid, and therefore activate the arachidonic acid cascade (Dumuis *et al.*, 1988; 1990).

There is little information on the regulation of late response genes in cell culture. However, Giraud and colleagues (1991) have demonstrated that preproenkephalin gene transcription can be activated by the cAMP-dependent protein kinase A and protein kinase C pathways in striatal cell cultures, with the use of selective drugs to target the respective enzymes. Similarly, treatment of striatal astrocytes with isoprenaline, or with forskolin or 8-bromo-cAMP, agents which directly increase intracellular cAMP, have been shown to elevate the level of preproenkephalin mRNA (Batter and Kessler, 1991).

Therefore in this study, we have attempted to reproduce the dopaminergic stimulation of c-fos and zif/268 gene expression in primary cultures of embryonic rat striatal neurons, in order to identify the intracellular pathway(s) and second messenger system(s) involved. As much controversy

surrounds the location of dopaminergic D1 and D2 receptors in the caudate-putamen. Two proposals are currently well supported. Firstly, that the two subtypes of dopamine receptor are located on separate populations of cells, on the basis that certain neuropeptides have been colocalised with a particular subtype but not the other and secondly that both D1 and D2 subtypes are expressed on the same cell. In the following studies we have addressed this question using the cultured striatal neuron model.

RESULTS

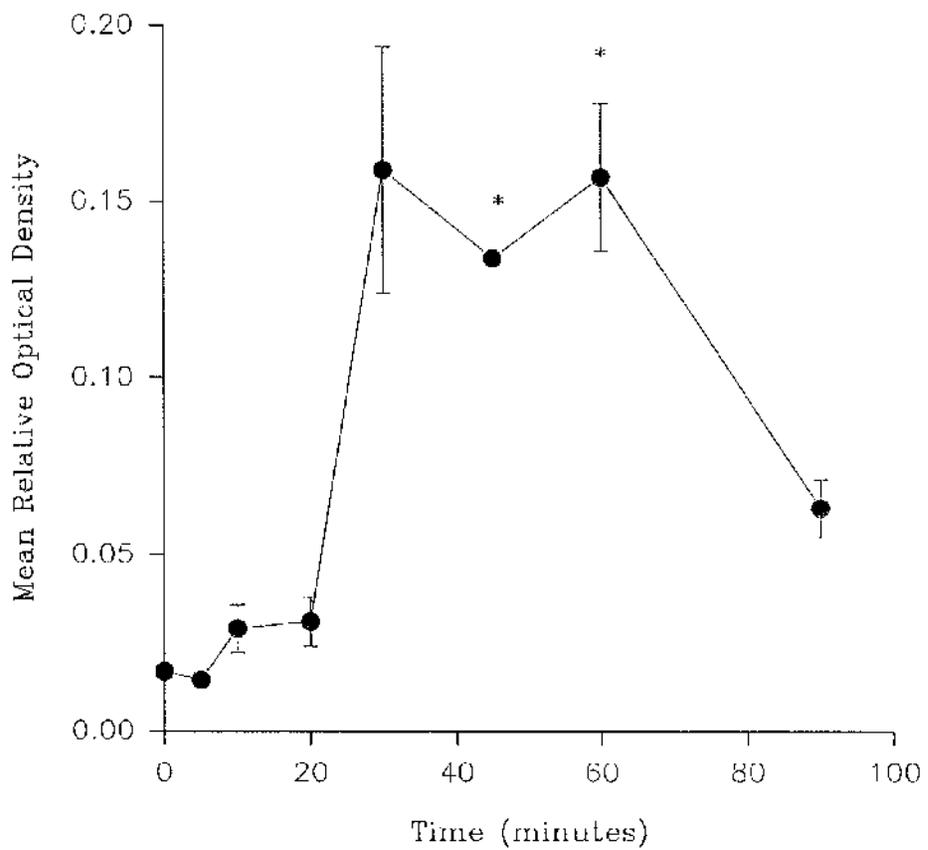
6.1 The time course effect of haloperidol on zif/268, jun-B and c-fos mRNA levels in cultured rat striatal neurons

Many IEGs including zif/268, jun-B and c-fos are known to be induced in vivo by the dopaminergic antagonist drug haloperidol (Nguyen et al, 1992). Therefore we have investigated the effect of haloperidol on these IEGs in a cultured striatal neuron system.

Neuronal cultures were exposed to dopamine ($5\mu\text{M}$) for two days. Haloperidol ($2\mu\text{M}$), when supplemented into the medium, was shown to induce the expression of zif/268 mRNA in vitro in a time-dependent manner. A dramatic increase in the levels of zif/268 mRNA was observed after 30 minutes, although a significant induction of zif/268 mRNA compared to the levels of expression at time zero, occurred only after 45 minutes [$F(2, 16) = 153.22, p < 0.05$] [Figure 6.1.1]. This maximal induction of zif/268 mRNA remained significantly high up to 60 minutes after haloperidol administration, $p < 0.05$ [Figure 6.1.1], and thereafter declined back to basal levels.

A similar, but more gradual induction of jun-B mRNA levels was observed with haloperidol ($2\mu\text{M}$). The induction of jun-B mRNA levels were significantly higher 20 minutes after treatment with haloperidol [$F(2,16) = 27.46, p = 0.006, p < 0.01$], than the levels expressed at time zero. A similar level of induction was observed 30 minutes after drug administration, $p < 0.01$. After 45 minutes the expression of jun-B mRNA remained significantly higher than at time zero, such that $p < 0.05$, after which the levels were not

Figure 6.1.1 : Effect of haloperidol on zif/268 mRNA levels in striatal cultures. Haloperidol ($2\mu\text{M}$) was added to cultures for 90, 60, 45, 30, 20, 10, and 5 minute time intervals before fixation in buffered formalin. Results, obtained from film autoradiographs, are expressed as mean relative optical density \pm s.e.m.; n=3. Significance was determined by one-way ANOVA, followed by post-hoc Dunnett's test. * $p < 0.05$



significantly induced compared to the levels of jun-B mRNA at time zero (not shown).

Haloperidol (2 μ M), also had a tendency to increase the levels of c-fos mRNA in striatal neuronal cultures, although the increase was generally variable, and proved not to be significantly different from the levels expressed at time zero (not shown).

6.2 Effect of dopamine receptor activation on the levels of c-fos and zif/268 mRNA in cultured rat striatal neurons.

The previous experiment established that maximal induction of zif/268 mRNA occurred 45 minutes after drug application. Since the induction of jun-B mRNA and c-fos mRNA followed a similar pattern, this time point was consequently used for all subsequent experiments. The probes used in the following studies have all been extensively characterised under the conditions used (Wisden *et al.*, 1990). Nevertheless, the specificity of each probe used was confirmed in a number of preliminary experiments. Hybridisation of sense probes labelled to the same specific activity, gave no signal, and addition of a 25-fold excess of unlabelled antisense oligonucleotide to the hybridisation buffer completely displaced the hybridisation signal (refer to chapter 5).

The hybridisation signal representing c-fos mRNA was generally low in unstimulated cells. In unstimulated cultures a small percentage of neurons were labelled for c-fos following treatment with vehicle alone (Hank's balanced salts solution). [Figure 6.2.1]. This was assumed similar to the low levels expressed *in vivo* when the cells remain unstimulated. Following treatment with dopamine

(5 μ M) a dramatic increase in the hybridisation signal for c-fos mRNA was observed. [Figure 6.2.1]. This induction was highly significant compared to the low levels of c-fos mRNA expressed in vehicle treated cultures [F(3, 22) = 18.57, p= 0.0007, p< 0.001]. The magnitude of induction was related to the concentration of dopamine applied. The levels of c-fos mRNA were induced by 100nM dopamine (not shown) , but a maximal induction was obtained at 5 μ M dopamine. Coadministration of the selective D1/D5 receptor antagonist, SCH23390 (1 μ M) with dopamine (5 μ M), clearly caused a reduction in the hybridisation signal due to c-fos mRNA.. This decrease was demonstrated to be significantly different from the signal due to dopamine alone [F (3,22) =18.57, p< 0.05] [Figure 6.2.1]. However ,coadministration of eticlopride (1 μ M), the selective dopamine D2 receptor antagonist, with dopamine proved not to be significantly different from dopamine alone, but was significantly different from vehicle alone [F (3,22) =18.57, p< 0.05] [Figure 6.2.1].

A very similar pattern of induction was observed for zif/268 mRNA in cultured striatal neurons. Low levels of zif/268 mRNA were expressed in vehicle treated cultures, which was assumed similar to the low levels expressed in vivo when the cells are unstimulated. Addition of a 25-fold excess of unlabelled probe to the hybridisation buffer, abolished the hybridisation signal completely (not shown), confirming specificity of the probe. Dopamine (5 μ M), clearly increased the hybridisation signal due to zif/268 mRNA , and this increase was significantly different from vehicle alone (n= 8) [F(3,16) = 12.11, p= 0.0007, p< 0.001] [Figure 6.2.2]. By coadministering SCH23390 (1 μ M) along with dopamine (5 μ M) a dramatic reduction in the hybridisation signal was observed, such that the reduction was different from dopamine alone [F(3,16) = 12.11, p< 0.001] [Figure 6.2.2]. Coadministration of eticlopride (1 μ M) with dopamine (5 μ M) clearly did not reduce the signal representing the level of zif/268 mRNA compared to dopamine alone, but was significantly different from vehicle alone [F (3,16) = 12.11, p< 0.001] [Figure 6.2.2].

Figure 6.2.1 : Effect of dopamine receptor stimulation on c-fos mRNA levels in striatal cultures. Cultures were treated with vehicle or dopamine ($5\mu\text{M}$), or treated by coadministration of dopamine / SCH23390 ($1\mu\text{M}$), or coadministration of dopamine / Eticlopride ($1\mu\text{M}$) for 45 minutes. Results, obtained from film autoradiographs, are expressed as mean relative optical density units $\times 100$ \pm s.e.m. from the number of different cultures shown in parentheses. * $p < 0.05$, *** $p < 0.001$, relative to vehicle alone; +++ $p < 0.001$ relative to dopamine alone.

Figure 6.3.1 : Effect of dopamine D1 receptor stimulation on c-fos mRNA levels in striatal cultures. Cultures were treated with vehicle, or SKF38393 ($1\mu\text{M}$), or treated by coadministration of SKF38393 / SCH23390 ($1\mu\text{M}$), or coadministration of SCH23390 / Eticlopride ($1\mu\text{M}$) for 45 minutes. Results, obtained from film autoradiographs, are expressed as mean relative optical density units $\times 100$ \pm s.e.m. from the number of different cultures shown in parentheses. * $p < 0.05$, *** $p < 0.001$, relative to vehicle alone; +++ $p < 0.001$ relative to SKF38393 alone.

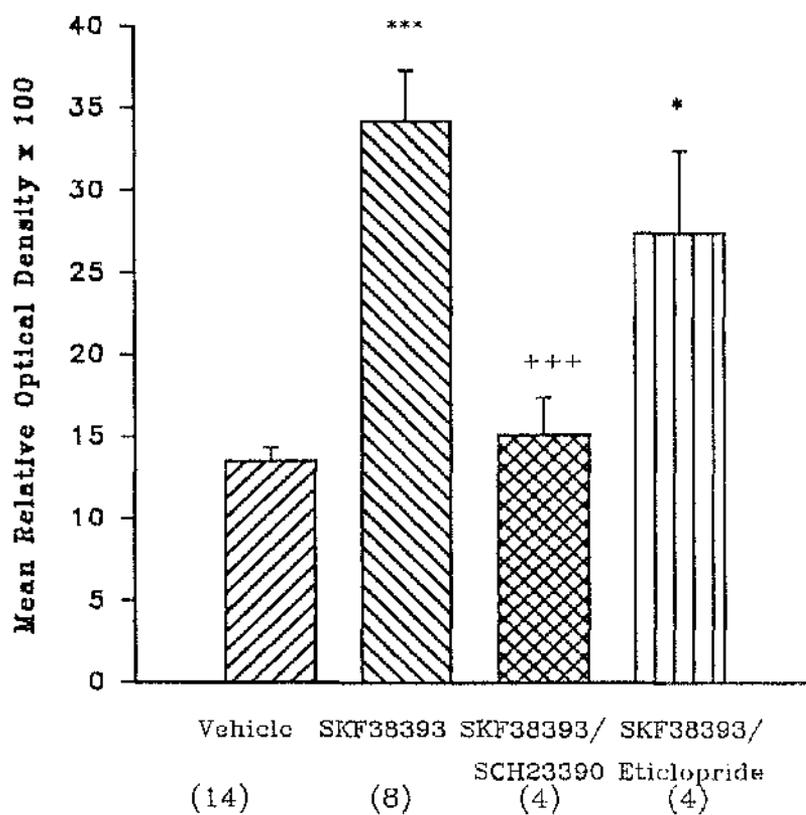
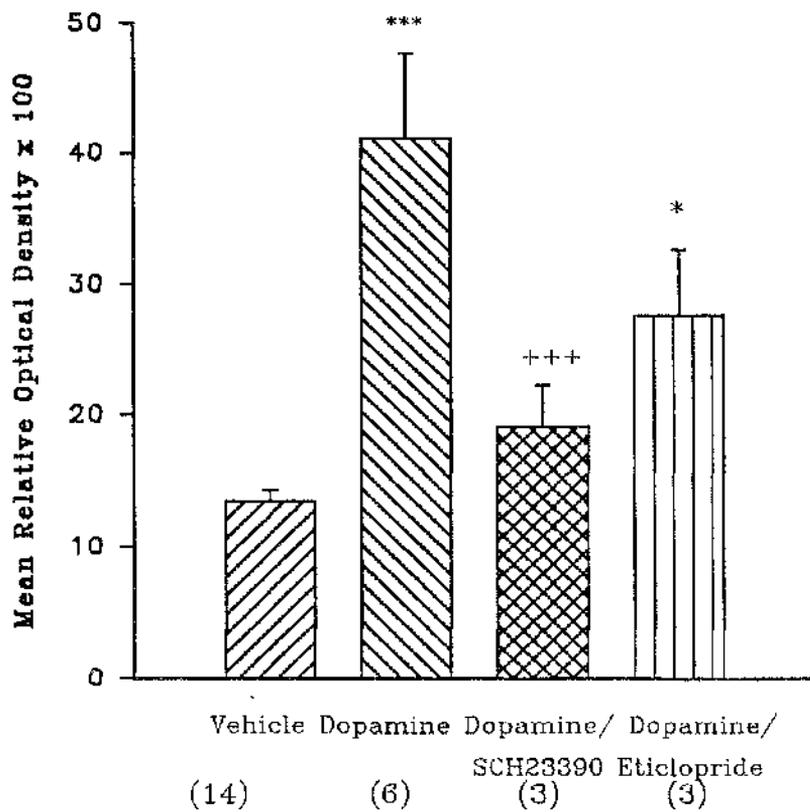
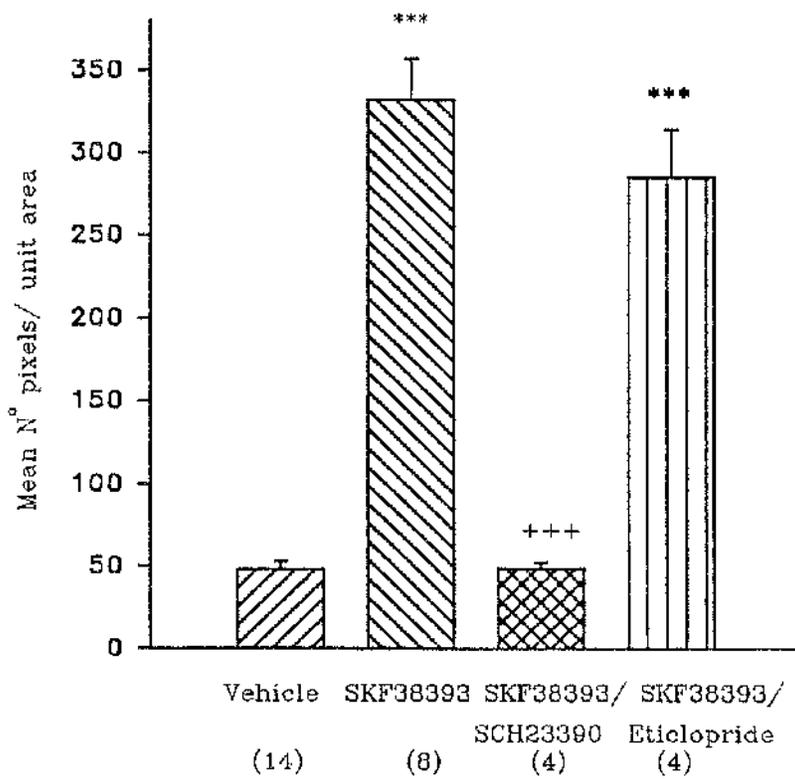
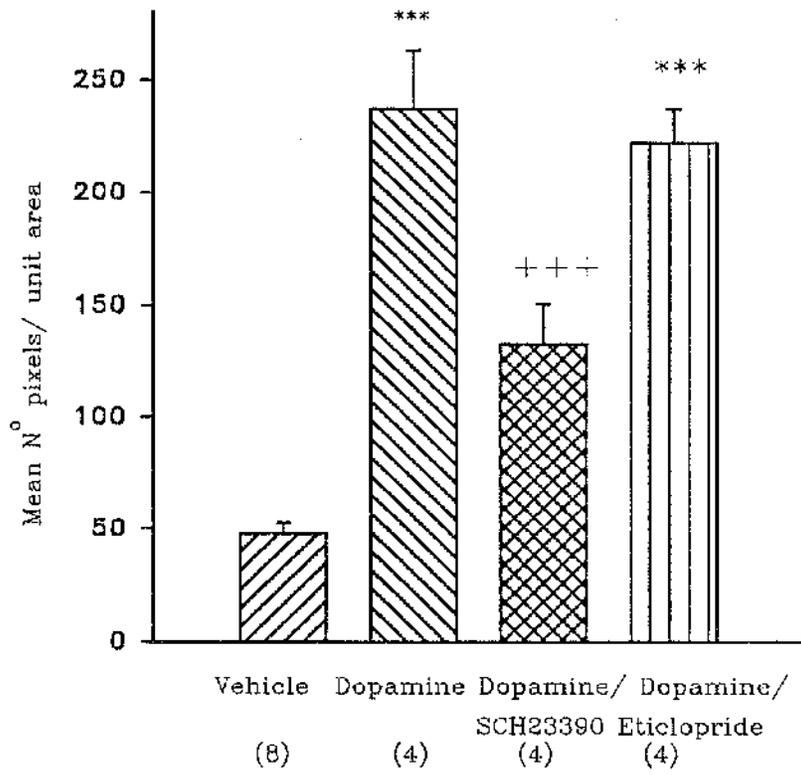


Figure 6.2.2 : Effect of dopamine receptor stimulation on zif/268 mRNA levels in striatal cultures. Cultures were treated with vehicle, or dopamine (5 μ M), or treated by coadministration of dopamine / SCH23390 (1 μ M), or coadministration of dopamine / Eticlopride (1 μ M) for 45 minutes. Results are expressed as mean number of pixels / unit area +s.e.m. from the number of different cultures shown in parentheses. *** p< 0.001, relative to vehicle alone; +++ p< 0.001, relative to dopamine alone.

Figure 6.3.2 : Effect of dopamine D1 receptor stimulation on zif/268 mRNA levels in striatal cultures. Cultures were treated with vehicle, or SKF38393 (1 μ M), or treated by coadministration of SKF38393 / SCH23390 (1 μ M), or by coadministration of SKF38393 / Eticlopride (1 μ M) for 45 minutes. Results are expressed as mean number of pixels / unit area +s.e.m. from the number of different cultures shown in parentheses. *** p< 0.001, relative to vehicle alone; +++ p< 0.001, relative to SKF38393 alone.



6.3 Effect of D1/D5 dopamine receptor stimulation on the levels of c-fos and zif/268 mRNA in cultured striatal neurons

Low levels of both c-fos and zif/268 mRNA were expressed in vehicle treated cultures. Following the addition of the D1/D5 receptor agonist, SKF38393 (1 μ M) a dramatic induction of c-fos mRNA was observed, and this increase was significantly different from vehicle alone [F (3,26) = 33.83, p= 0.0006, p< 0.001] [Figure 6.3.1]. Similarly, treatment with SKF38393 (1 μ M) caused a massive induction of zif/268 mRNA in these cultures [F (3,26) = 33.83, p< 0.001] [Figure 6.3.2, 6.4.3]. SCH23390 (1 μ M) when coadministered with SKF38393, was demonstrated to reduce the hybridisation signal for both c-fos and zif/268 [Figure 6.4.3] respectively. This decrease in the hybridisation signal was significantly different from SKF38393 alone for both c-fos [F (3,26) = 33.83, p< 0.001] [Figure 6.3.1] and zif/268 [F (3,26) = 33.83, p< 0.001] [Figure 6.3.2, 6.4.3]. However, coadministration of eticlopride (1 μ M) with SKF38393 was shown to cause an induction of c-fos mRNA [F (3,26) = 33.83, p< 0.05] [Figure 6.3.1] and zif/268 mRNA [F (3,26) = 33.83, p< 0.001] [Figure 6.3.2, 6.4.3], compared to vehicle alone (n= 14), but there was no significant decrease in the signal when compared to SKF38393 alone (n= 8).

In every case the increase in silver grain density was found to occur over the smaller, more intensely counterstained cells in the cultures. These were identified in preliminary immunocytochemical experiments as neurons. Despite a large percentage of neurons being labelled following application of dopaminergic agonists, there were still many neurons that did not show a

significant accumulation of silver grains. No increases in silver grain density were detected over the larger, more weakly counterstained non-neuronal cells.

6.4 Effect of inhibition of protein kinase A on c-fos and zif/268 mRNA levels in striatal cultures

The second messenger systems involved in the induction of the IEGs c-fos and zif/268 mRNA were investigated using selective inhibitors of serine/threonine protein kinases. The selective protein kinase A inhibitor, KT5720 (2 μ M) (Kase *et al.*, 1987), when applied alone to striatal cultures had no effect on the expression of either IEG when compared to vehicle treated cultures (not shown). However pretreatment of the cultures for 30 minutes with KT5720, prior to the addition of SKF38393 (1 μ M) caused a small but significant reduction in the ability of SKF38393 to induce c-fos mRNA levels [F (3,26) = 14.23, p= 0.0033, p< 0.01] [Figure 6.4.1, 6.3.4]. In the presence of KT5720, the levels of c-fos mRNA due to indirect stimulation via SKF38393, were still significantly higher [F (3,26) = 14.23, p<0.001], than the levels expressed in vehicle alone. Pretreatment with a relatively non-selective inhibitor of all protein kinases, H7 (50 μ M), caused a more dramatic reduction in the hybridisation signal due ^{to} the induction of c-fos mRNA by SKF38393 [F (3,26) = 14.23, p< 0.001] when compared to SKF38393 alone [Figure 6.4.1]. This information suggests the involvement of a protein kinase other than protein kinase A.

Figure 6.4.1 : Effect of inhibition of protein kinase A on c-fos mRNA levels in striatal cultures. KT5720 (2 μ M), or H7 (50 μ M) or vehicle, were added 30 minutes prior to treatment with SKF38393 (1 μ M), for 45 minutes. Results are expressed as mean number of pixels / unit area + s.e.m. from the number of different cultures shown in parentheses. *** p< 0.001, relative to vehicle alone; ++ p< 0.01, +++ p< 0.001, relative to SKF38393 alone.

Figure 6.4.2 : Effect of inhibition of protein kinase A on zif/268 mRNA levels in striatal cultures. KT5720 (2 μ M), or H7 (50 μ M), or vehicle were added 30 minutes prior to treatment with SKF38393 (1 μ M), for 45 minutes. Results are expressed as mean number of pixels / unit area + s.e.m. from the number of different cultures shown in parentheses. * p< 0.05, *** p< 0.001, relative to vehicle alone; +++ p< 0.001, relative to SKF38393 alone.

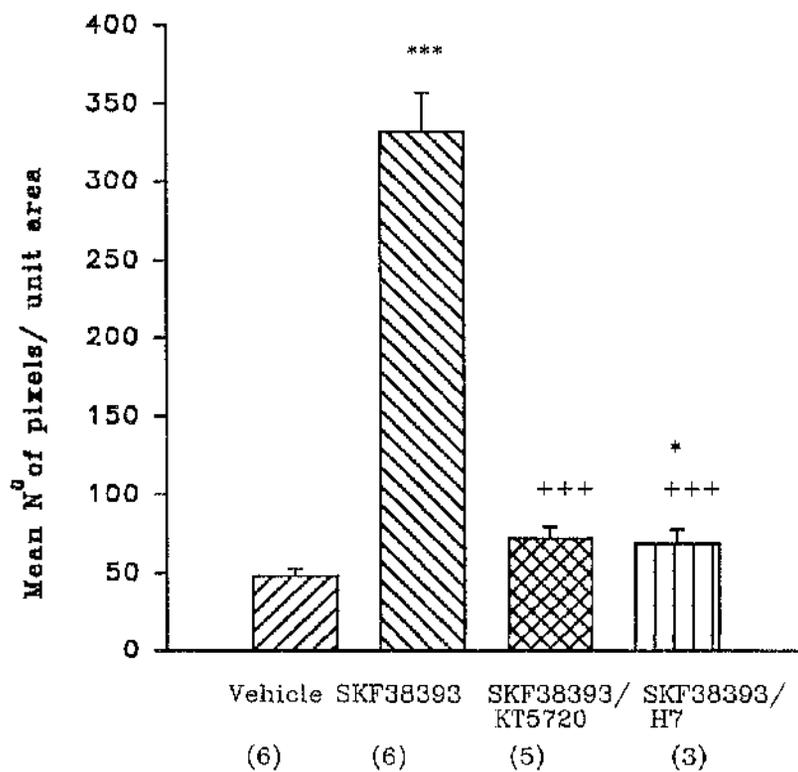
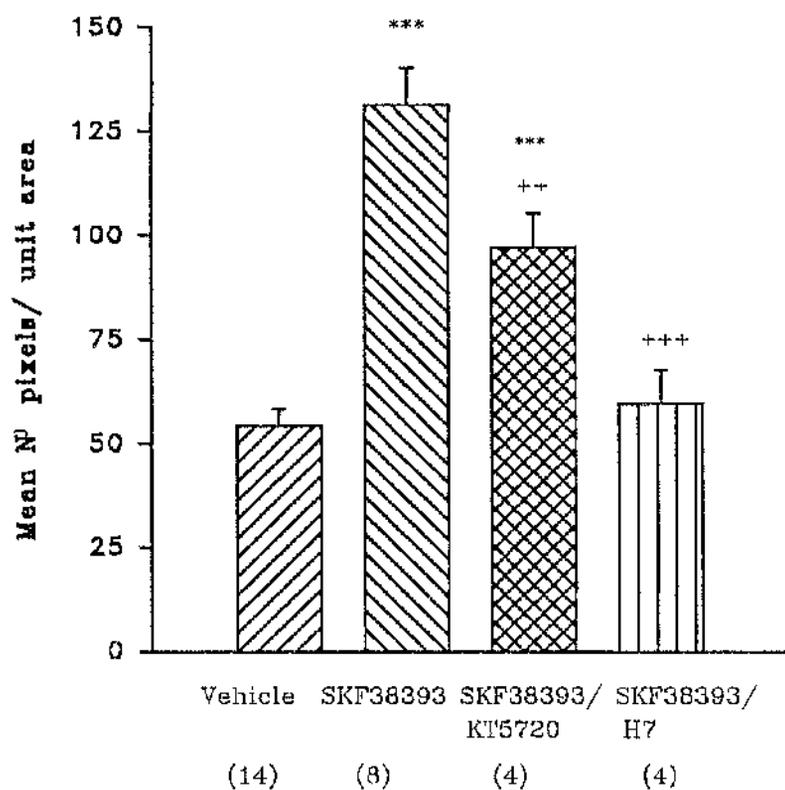
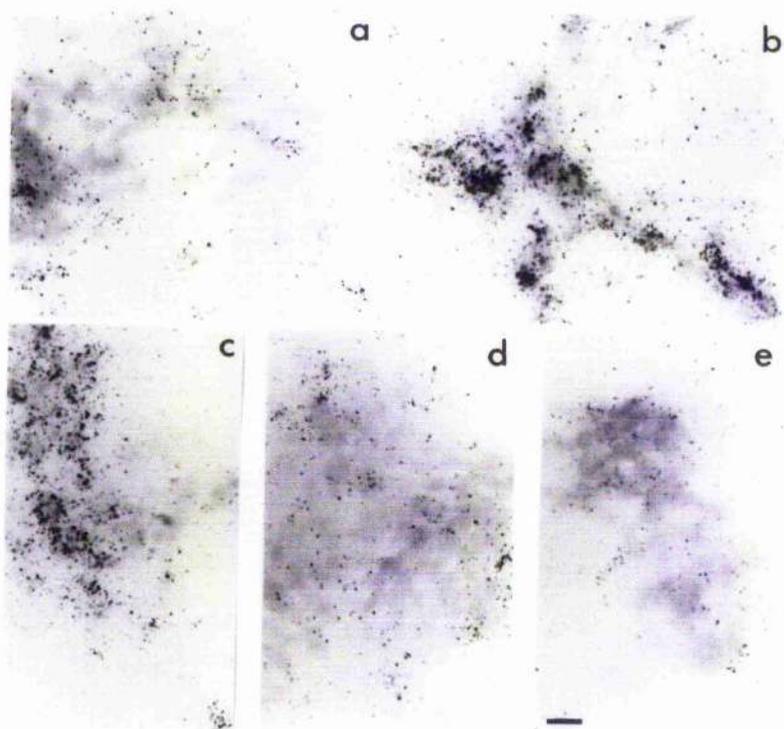


Figure 6.4.3 : Induction of *zif/268* mRNA by D1-like dopaminergic agonists. Cultured rat striatal neurones are shown following hybridisation with the probe to detect *zif/268* mRNA. Black silver grains overlying counterstained cell bodies represent the hybridisation signal. Cells were treated with (a) vehicle (HBSS) (b), SKF38393 (1 μ M) (c) SKF38393 (1 μ M) + eticlopride (1 μ M), (d) SKF38393 (1 μ M) + SCH23390 (1 μ M), (e) SKF38393 (1 μ M) after pretreatment with KT5720 (2 μ M). Scale bar represents 20 μ m.



Pretreatment of the cultures with KT5720 (2 μ M), dramatically inhibited the ability of SKF38393 to induce zif/268 mRNA by approximately 90%. This decrease in signal was shown to be significantly different from SKF38393 treatment alone, such that $F(3,16) = 66.40$, $p = 0.00076$, $p < 0.001$, [Figure 6.4.2, 6.4.3]. Similarly, pretreatment with H7 (50 μ M) caused a significant reduction in the levels of zif/268 mRNA due to stimulation by SKF38393 [$F(3,16) = 66.40$, $p < 0.001$] [Figure 6.4.2].

6.5 Effect of inhibition of protein kinase C on the levels of c-fos and zif/268 mRNA in rat striatal cultures

From the previous experiment, it is clear that the role of protein kinase enzymes in the induction of c-fos and of zif/268 mRNA is relatively complex. Therefore we went on to investigate the role of protein kinase C (PKC) in these responses using the selective PKC inhibitor, calphostin C (Bruns *et al.*, 1991). This PKC inhibitor has been reported to be activated only by exposure to light (Bruns *et al.*, 1991).

Initially the efficiency of PKC inhibition in striatal cultures was monitored by observing the ability of calphostin C to reduce the activation of c-fos and zif/268 gene expression by phorbol-12-myristate-13-acetate (PMA). A significant induction in the levels of c-fos mRNA [$F(2,13) = 23.42$, $p = 0.007$, $p < 0.01$] [Figure 6.5.1] and zif/268 mRNA [$F(2,9) = 36.19$, $p = 0.0006$, $p < 0.001$] [Figure 6.5.3], compared to treatment with vehicle alone, were observed following treatment with 100nM PMA. Pretreatment for 30 minutes with the selective PKC inhibitor, calphostin C, caused a marked reduction in the levels of c-fos and zif/268 mRNA (approximately 70%). The decrease in c-

Figure 6.5.1: Effect of calphostin C on the phorbol ester-induced expression of c-fos mRNA in striatal cultures. Calphostin C (1mM), or vehicle were added 30 minutes prior to treatment with phorbol-12-myristate-13-acetate ($10^{-7}M$) for 45 minutes. Results obtained from film autoradiographs, are expressed as mean relative optical density units $\times 100$ \pm s.e.m. from the number of different cultures shown in parentheses. ** $p < 0.01$, *** $p < 0.001$, relative to vehicle alone; + $p < 0.05$ relative to phorbol-12-myristate-13-acetate alone.

Figure 6.5.2 : Effect of calphostin C on the SKF38393-induced expression of c-fos mRNA in striatal cultures. Calphostin C ($1\mu M$), or vehicle were added 30 minutes prior to treatment with of SKF38393 ($1\mu M$) for 45 minutes. Results, obtained from film autoradiographs, are expressed as mean relative optical density units $\times 100$ \pm s.e.m. from the number of different cultures shown in parentheses. *** $p < 0.001$, relative to vehicle alone; +++ $p < 0.001$ relative to SKF38393 alone.

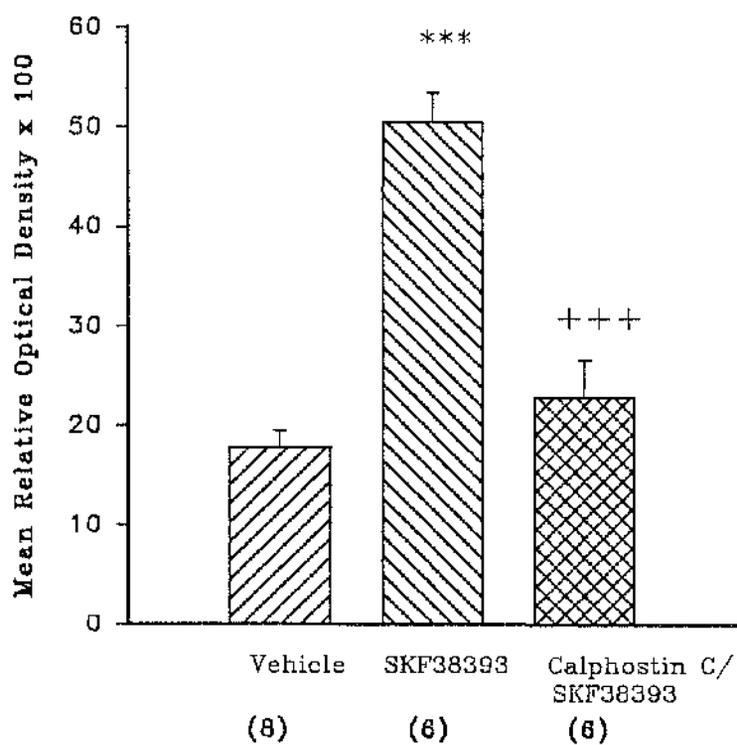
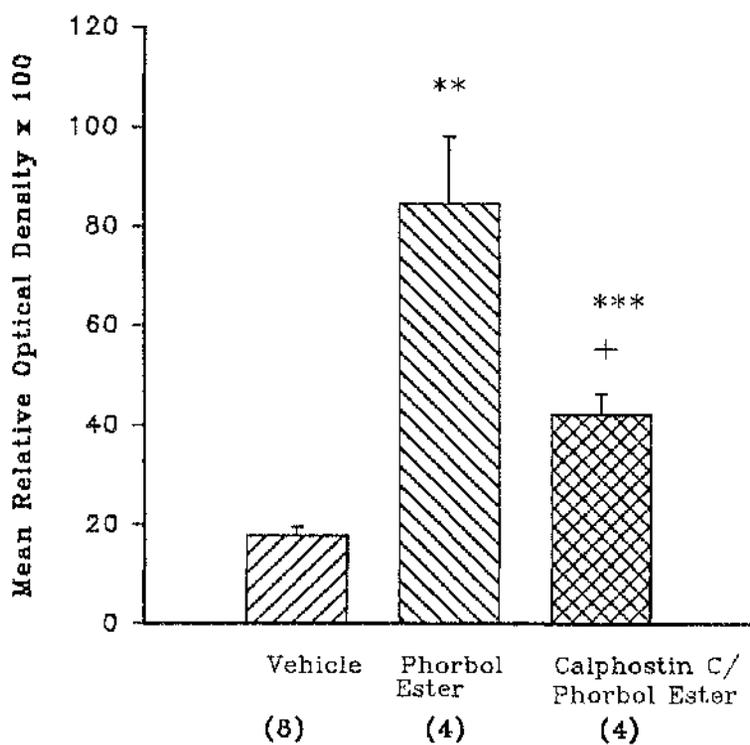
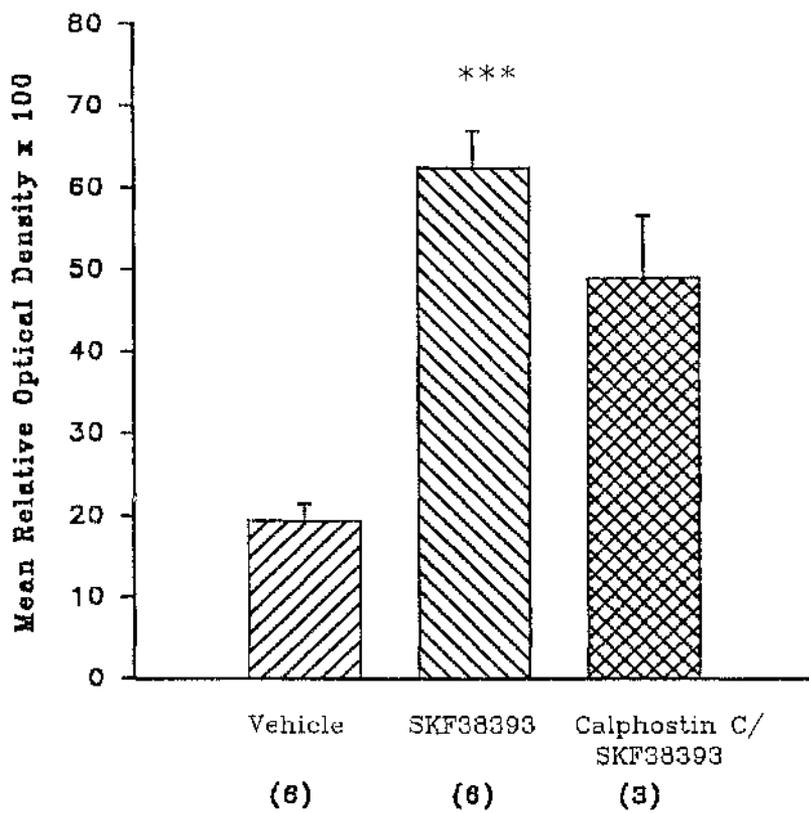
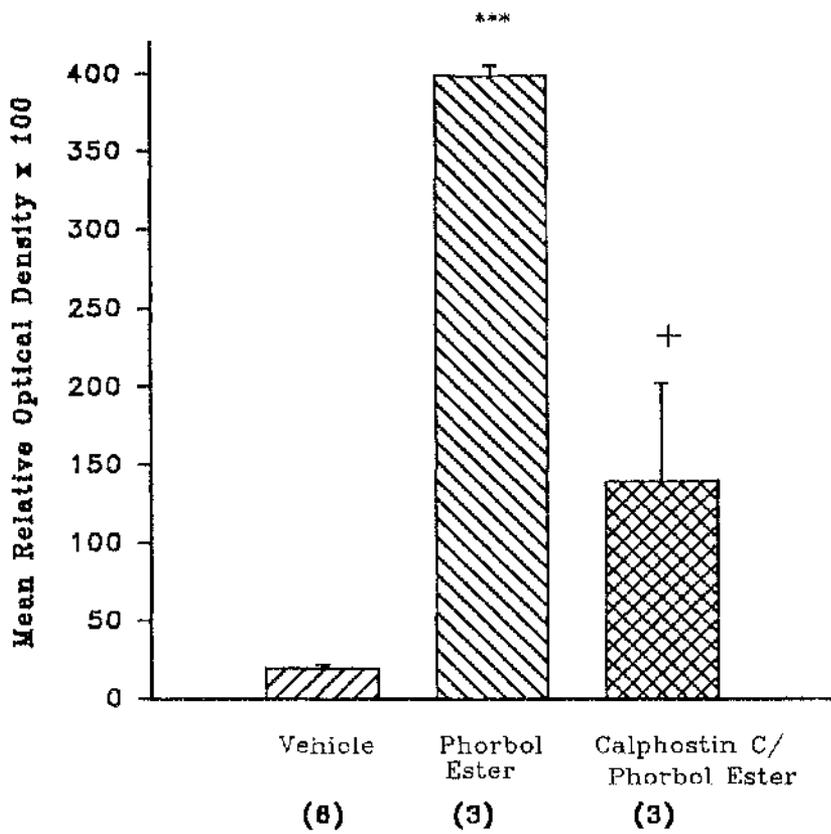


Figure 6.5.3 : Effect of calphostin C on phorbol ester-induced zif/268 mRNA levels in striatal cultures. Calphostin C ($1\mu\text{M}$), or vehicle were added to the cultures 30 minutes prior to treatment with phorbol-12-myristate-13-acetate (10^{-7}M) for 45 minutes. Results, obtained from film autoradiographs, are expressed as mean relative optical density units $\times 100 \pm \text{s.e.m.}$ from the number of different cultures shown in parentheses. *** $p < 0.001$, relative to vehicle alone; + $p < 0.05$, relative to phorbol-12-myristate-13-acetate alone.

Figure 6.5.4 : Effect of calphostin C on the SKF38393-induced expression of zif/268 mRNA levels in striatal cultures. Calphostin C ($1\mu\text{M}$), or vehicle were added 30 minutes prior to treatment with SKF38383 ($1\mu\text{M}$) for 45 minutes. Results, obtained from film autoradiographs, are expressed as mean relative optical density units $\times 100 \pm \text{s.e.m.}$ from the number of different cultures shown in parentheses. *** $p < 0.001$, relative to vehicle alone.



fos mRNA levels was shown to be significantly different from cultures treated with phorbol ester alone [$F(2,13) = 23.42, p < 0.05$] [Figure 6.5.1], and similarly the reduction in zif/268 mRNA levels was also significantly different from phorbol ester stimulated cultures alone [$F(2,9) = 36.19, p < 0.05$] [Figure 6.5.3].

As in previous experiments, SKF38393 ($1\mu\text{M}$) was shown significantly to induce the expression of c-fos mRNA [$F(2,17) = 36.06, p = 0.00033, p < 0.001$] [Figure 6.5.2]. Following pretreatment with calphostin C ($1\mu\text{M}$) for 30 minutes, the ability of SKF38393 ($1\mu\text{M}$) to induce the c-fos gene was dramatically inhibited. This reduction was demonstrated to be significantly different from SKF38393 treatment alone [$F(2,17) = 36.06, p < 0.001$] [Figure 6.5.2], such that the levels of c-fos mRNA were not significantly different from basal levels. The expression of zif/268 mRNA was also shown to be increased to significant levels after SKF38393 ($1\mu\text{M}$) treatment [$F(2,12) = 37.14, p = 0.00086, p < 0.001$] [Figure 6.5.4], as in previous experiments. However, pretreatment with calphostin C ($1\mu\text{M}$) was unable to cause a significant reduction in the increase in zif/268 mRNA after treatment with SKF38393 [Figure 6.5.4].

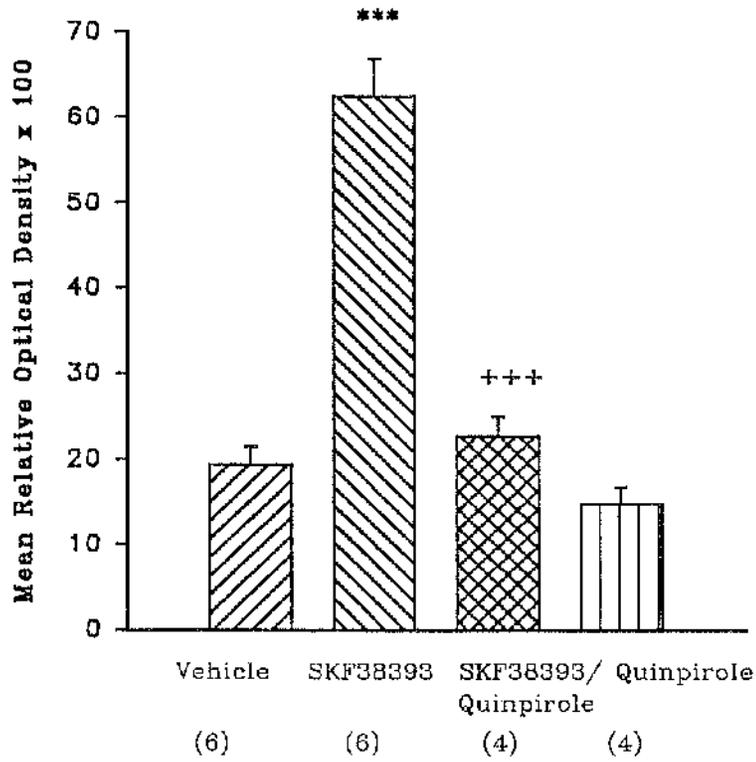
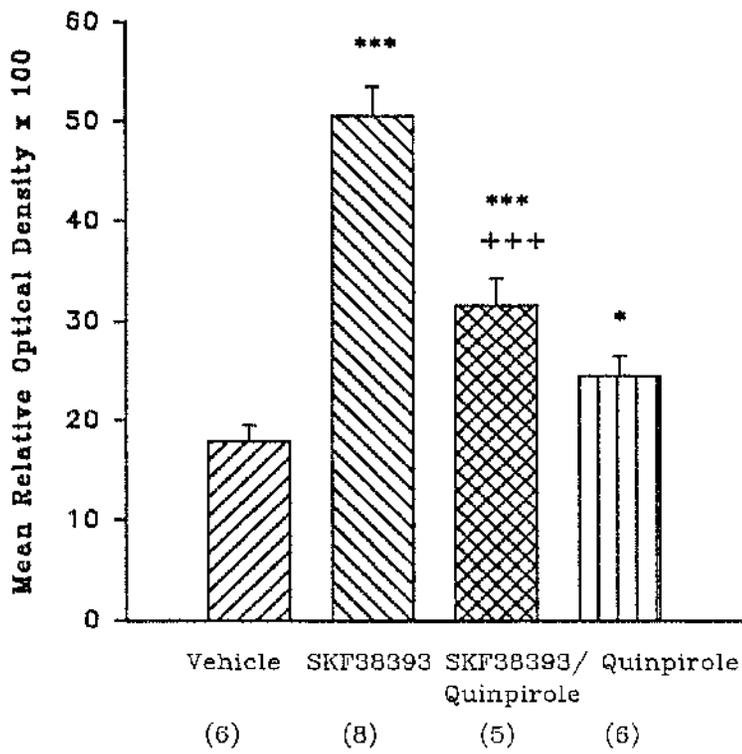
6.6 Effect of quinpirole on c-fos and zif/268 mRNA expression in cultured rat striatal neurons

Low levels of c-fos and zif/268 mRNA were expressed in vehicle treated cultures compared to the highly significant induction of both c-fos mRNA [$F(3,19) = 33.14, p = 0.0006, p < 0.001$] [Figure 6.6.1] and zif/268 mRNA [$F(3,16) = 28.23, p = 0.0009, p < 0.001$] [Figure 6.6.2] after SKF38393

treatment. Coadministration of SKF38393 ($1\mu\text{M}$) together with quinpirole ($1\mu\text{M}$), the selective D2 dopamine receptor agonist, caused a marked fall in the hybridisation signal representing c-fos mRNA, due to SKF38393 treatment. This decrease was significantly different from SKF38393 alone [$F(3,19) = 33.14$, $p < 0.01$] [Figure 6.6.1]. A similar suppression of the hybridisation signal for zif/268 mRNA was observed when quinpirole was coapplied with SKF38393, and this was significantly different from cultures treated with SKF38393 alone [$F(3,16) = 28.23$, $p < 0.001$]. [Figure 6.6.2]. Quinpirole ($1\mu\text{M}$) when applied alone had no effect on the levels of zif/268 mRNA expressed in these cultures [Figure 6.6.2]. However, the levels of c-fos were slightly increased in the presence of quinpirole alone. This increase was shown to be significantly different from vehicle treated cultures [$F(3,19) = 33.14$, $p < 0.05$] [Figure 6.6.1].

Figure 6.6.1: Effect of quinpirole on the SKF38393-induced expression of c-fos mRNA levels in striatal cultures. Quinpirole (1 μ M), or vehicle was added 30 minutes prior to treatment with SKF38393 (1 μ M), for 45 minutes. Results, obtained from film autoradiographs, are expressed as mean relative optical density units \times 100 + s.e.m. from the number of different cultures shown in parentheses. * $p < 0.05$, *** $p < 0.001$, relative to vehicle alone; +++ $p < 0.001$, relative to SKF38393 alone.

Figure 6.6.2 : Effect of quinpirole on the SKF38393-induced expression of zif/268 mRNA levels in striatal cultures. Quinpirole (1mM), or vehicle were added 30 minutes prior to treatment with SKF38393 (1mM), for 45 minutes. Results, obtained from film autoradiographs, are expressed as mean relative optical density units \times 100 + s.e.m. from the number of different cultures shown in parentheses. *** $p < 0.001$, relative to vehicle alone; +++ $p < 0.001$, relative to SKF38393 alone.



DISCUSSION

Initially it was necessary to determine the time course of induction of the immediate early genes under investigation. Previous *in vivo* studies have suggested that peak induction of the immediate early genes *c-fos* and *zif/268* mRNAs occurs thirty to sixty minutes after the initial injection of both psychostimulants or neuroleptic drugs *in vivo* (Robertson *et al.*, 1989; Cole *et al.*, 1992; Nguyen *et al.*, 1992; Bhat *et al.*, 1992). In agreement with this our results show that the immediate early gene *zif/268* is maximally induced in primary culture approximately forty-five minutes after the addition of haloperidol to the culture medium. The rise in *zif/268* mRNA levels *in vivo*, has been shown to be apparent as early as fifteen minutes after the injection (Cole *et al.*, 1992). The lack of an induction of *zif/268* mRNA at early time points *in vitro*, is likely to reflect differences in the neuronal environment such as temperature, and thus may be responsible for delayed signal transduction processes. Similar to the *in vivo* situation (Cole *et al.*, 1992; Nguyen *et al.*, 1992), *zif/268* mRNA levels *in vitro*, decline back to basal levels by approximately three hours.

6.7 Dopaminergic Regulation of Immediate Early Genes In Vitro

In this study we have demonstrated that exposure of rat striatal primary cultures to dopamine resulted in dramatic increases in the levels of *c-fos* and *zif/268* mRNAs. The use of *in situ* hybridisation to monitor these changes has the advantage that the cell types where the changes in gene expression occur can be identified with some confidence. It is worth noting that despite the small

proportion of non-neuronal cells in such cultures, some pharmacological agents can stimulate dramatic increases in immediate early gene expression in the glial cells. The induction of *c-fos* and *zif/268* mRNA levels that we observed in these studies, following dopamine receptor stimulants appears to occur in the same population of cells that exhibit immunoreactivity for neuron-specific enolase. Therefore this induction of immediate early gene expression is occurring in the striatal neurons.

We have previously shown that a high proportion of striatal neurons in primary culture express either the proenkephalin gene or the preprotachykinin gene, and therefore are likely to represent the medium spiny projection neurons characterised in the mature striatum (Graybiel, 1986). Interneurons are thought to constitute only a small percentage of the number of neurons in the normal striatum (Graybiel, 1986), and we have shown earlier that the corresponding neurochemical phenotypes represent a similarly small proportion of striatal neurons in primary striatal culture. The high proportion of neurons containing *c-fos* mRNA or *zif/268* mRNA in these experiments after dopaminergic stimulation suggests that immediate early gene induction is taking place in the culture counterpart of the medium spiny neuron. This is consistent with evidence suggesting that the induction of immediate early gene expression in striatal neurons *in vivo* following manipulation of the level of dopaminergic activity occurs in medium spiny neurons (Dragunow *et al.*, 1990; Graybiel *et al.*, 1990; Robertson *et al.*, 1990; Robertson *et al.*, 1992). The induction of *c-fos* or *zif/268* following dopamine stimulation was observed with a dosage ranging from 100nM to 5 μ M, concentrations which would be expected to stimulate all dopamine receptor subtypes.

In rats with a unilateral 6-hydroxydopamine lesion of the nigrostriatal pathway, there is evidence that the induction of *c-fos* by SKF38393 occurs primarily in the medium spiny neurons expressing the preprotachykinin gene and projecting to the substantia nigra (Robertson *et al.*, 1989; Robertson *et al.*,

1992). The number of neurons showing increased c-fos expression in culture after treatment with the dopaminergic D1 agonist, SKF38393, means that the preprotachykinin-containing neurons must be involved, but we would not rule out the possibility that enkephalinergic neurons in culture also respond to SKF38393 with an increased level of c-fos expression.

Since the addition of SKF38393 to the culture medium was able to reproduce the dopaminergic stimulation of c-fos and zif/268 mRNA, but the addition of quinpirole had no effect on zif/268 and a minor inductive effect on c-fos, it is most likely that this response is mediated by stimulation of a D1/ D5-like receptor. This is further supported by the observation that SCH23390, the D1 selective antagonist, was able to attenuate the inductive effect of SKF38393, but the selective D2 antagonist, eticlopride was unable to affect these responses. This is in accordance with previous work in vivo, which also suggests that the induction of c-fos and zif/268 after administration of dopaminergic agonists is likely to occur via stimulation of D1-like dopamine receptors (Robertson *et al.*, 1989; Robertson *et al.*, 1990; Graybiel *et al.*, 1990; Arnauld *et al.*, 1993; Cole *et al.*, 1992; Mailleux *et al.*, 1992; Moratalla *et al.*, 1992). The effects of dopaminergic agonists on immediate early gene expression in primary cultures reported here therefore appear to reflect the corresponding in vivo actions very closely.

Significant increases in c-fos and zif/268 mRNA levels in striatal neurons were detected following treatment with SKF38393 at concentrations between 30nM and 5 μ M. The EC50 value obtained is very close to the K_D values reported for the binding of SKF38393 to striatal membranes (Seeman and Van Tol, 1993). This suggests that there is a low receptor reserve for SKF38393 in stimulating these changes, consistent with the biochemical evidence that, in striatal tissue, SKF38393 acts as an agonist with only moderate efficacy (Setler *et al.*, 1978).

It is clear that, in a number of different cell types, increased transcription of both the *c-fos* and *zif/268* genes can be observed following activation of either protein kinase A (PKA) or protein kinase C (PKC) (review by Sheng and Greenberg, 1990). This is in line with the presence of cyclic AMP- responsive enhancers and phorbol ester- responsive enhancers in the promoter regions of these genes. The biochemical evidence suggests that D1-like dopamine receptors can be coupled either to stimulation of adenylate cyclase or phospholipase C, with consequent activation of PKA or PKC respectively (Mahan *et al.*, 1990; Monsma *et al.*, 1990; Sunahara *et al.*, 1991; Tiberi *et al.*, 1991; Frail *et al.*, 1993). The dopaminergic stimulation of *c-fos* and *zif/268* gene expression in striatal neurons could therefore potentially be mediated via either PKA or PKC.

Highly selective inhibitors are now available which allow the intracellular actions of different serine / threonine kinases to be distinguished. The highly selective inhibitor of PKA, namely KT5720 (Kase *et al.*, 1987), blocked the ability of SKF38393 to stimulate *zif/268* expression by 90%. The K_i value for KT5720 against PKA is approximately 60nM, whereas its K_i value against other kinases such as PKC is greater than 2 μ M (Kase *et al.*, 1987). The concentration of KT5720 used in this study would therefore be predicted to cause a 95% inhibition of PKA and less than 50% inhibition of PKC. The degree of attenuation of the effect of SKF38393 on *zif/268* induction observed with KT5720 therefore strongly implicates protein kinase A in this effect.

Calphostin C is a highly selective inhibitor of PKC, which shows negligible activity against other serine/ threonine kinases (Kobayashi *et al.*, 1989). The concentration of 1 μ M calphostin C used in these experiments, should consequently inhibit PKC by more than 90%, with only minimal effects on other serine/ threonine kinases. However, an important aspect concerning the use of calphostin C, is that its activity is dependent on exposure to light (Bruns *et al.*, 1991). We observed that treatment of the cultures with the

phorbol ester, phorbol-12-myristate-13 acetate (PMA), resulted in a clear induction of both *c-fos* and *zif/268* mRNA levels compared to controls. Pretreatment with calphostin C caused a 70% decrease in the ability of PMA to induce expression of both *c-fos* and *zif/268* mRNA levels. While this may imply that the activation of calphostin C was not quite maximal in our protocol, it provides a definite indication of the degree of PKC inhibition achieved.

Consistent with the above evidence that the actions of dopaminergic agonists on *zif/268* expression were mediated entirely by PKA, calphostin C pretreatment had no significant effect on the stimulation of *zif/268* mRNA levels by SKF38393. In contrast, calphostin C pretreatment caused a dramatic reduction in the ability of SKF38393 to enhance *c-fos* expression. The relatively greater inhibition by calphostin C of the SKF38393 response as compared to the PMA response may simply reflect a sub-maximal activation of PKC by the endogenous intracellular pathway, subsequent to stimulation of D1-like dopamine receptors. It is of some interest that the SKF38393-induced increase in *c-fos* gene expression was also reduced by inhibition of PKA. This implies that PKA activity also participates in the D1-dopaminergic induction of *c-fos* expression.

These results therefore suggest that stimulation of *c-fos* expression by SKF38393 involves mainly PKC, but with an additional contribution from PKA. Conversely, the stimulation of *zif/268* gene expression by SKF38393 appears to proceed almost entirely via PKA. However, it is important to remember that there may be a further level of complexity involved in these effects, in that phosphorylation of cyclic AMP-response element binding proteins by PKA or PKC may affect their relative affinity for the cyclic AMP-responsive and phorbol ester-responsive enhancers (Merino *et al.*, 1989). In other words, even when only a single kinase is activated, its ability to modulate gene expression may be affected by altering the basal level of activity of another kinase.

Nevertheless, some important conclusions can be drawn from these results. The intracellular pathways coupled to induction of IEGs vary in different cell types. It is clear that in non-neuronal cells and cell lines, both PKA and PKC are powerful stimulants for the induction of both *c-fos* and *zif/268* (reviewed by Sheng and Greenberg, 1989). The information available on neuronal cell responses is rather scarce, but it appears that in primary cultures of cerebellar granule cells, *c-fos* expression can be stimulated by PKC, but not by PKA (Szekely *et al.*, 1989). In contrast, in primary cultures of cerebral cortical neurons, activation of PKA can produce a mild enhancement of expression of the *c-fos* and *zif/268* genes (Vaccarino *et al.*, 1993). Our results represent, to the best of our knowledge, the first indication that both PKA and PKC can be coupled to stimulation of *c-fos* and *zif/268* expression in striatal neurons. Furthermore, our data suggests that the population of D1-like dopamine receptors on striatal neurons includes receptors coupled to both adenylate cyclase and to phospholipase C. Until now, the evidence for coupling of D1-like receptors in the central nervous system to phospholipase C has been derived from *in vitro* experiments using translated endogenous and exogenous nucleic acids (Mahan *et al.*, 1990; Monsma *et al.*, 1990; Frail *et al.*, 1993), or from extrapolation of results from kidney cells (Felder *et al.*, 1989). The evidence reported here therefore provides a functional confirmation of the molecular evidence that D1-like receptors on striatal neurons can couple to both second messenger systems.

The different relative contribution of PKC- and PKA-dependent pathways to the dopaminergic induction of *c-fos* and *zif/268* expression may have implications for the regulation of striatal gene expression by other neurotransmitters. For example a receptor negatively linked to adenylate cyclase might act to suppress *zif/268* expression more than *c-fos* expression. It appears that, when activation of both PKC and PKA occurs in striatal neurons, the rate of *zif/268* transcription may be more sensitive to changes in PKA

activity, while the rate of c-fos transcription may be more sensitive to changes in PKC activity. In the light of our results, investigations of the second messenger systems involved in the regulation of immediate early gene expression by other neurotransmitter receptors are clearly of some importance.

While we have shown that dopamine and SKF38393 are able to exert a powerful stimulatory influence on striatal immediate early gene expression by activating D1-like dopamine receptors, we also detected a relatively small enhancement of c-fos expression following treatment with the selective D2-like agonist quinpirole. This effect was specific for the c-fos gene, in that quinpirole did not affect zif/268 expression. At present the *in vivo* information suggests that D2-like agonists should act to suppress immediate early gene expression. However, there are thought to be a relatively small number of D3 receptors in the rat striatum, and quinpirole is known to be an effective agonist at D3 receptors (Sokoloff *et al.*, 1990). It has recently been reported that stimulation of D3 dopamine receptors can activate c-fos in a transfected cell line (Pilon *et al.*, 1994). Our results raise the interesting possibility that stimulation of D3 receptors may activate c-fos gene expression in a population of striatal neurons. However, further work would be necessary to explore this possibility.

We also report that activation of D2-like dopamine receptors by quinpirole reduces the stimulation of c-fos and zif/268 expression by SKF38393. Since the *in vivo* evidence suggests that tonic activity at D2-like receptors suppresses immediate early gene expression, (Dragunow *et al.*, 1990; Nguyen *et al.*, 1992; Robertson and Fibiger, 1992) this action of quinpirole conforms more closely to expectations. There has been considerable debate as to whether individual striatal neurons express both D1-like and D2-like dopamine receptors. Based on an inability to detect D1 receptor mRNA and D2 receptor mRNA in the same population of neurons by *in situ* hybridisation (Gerfen *et al.*, 1990; Le Moine *et al.*, 1991), it has been argued that these two receptor types are not colocalised in striatal neurons. In contrast functional and immunochemical evidence suggests

that medium spiny neurons contain both D1 and D2 receptors (Lester *et al.*, 1993; Surmeier *et al.*, 1992; Larson and Ariano, 1994).

Our results have demonstrated that quinpirole can act functionally to antagonise the effects of SKF38393 on striatal neurons. If it is assumed that SKF38393 and quinpirole are not acting trans-synaptically in our studies, then the data appears to confirm the coexpression of D1-like and D2-like receptors in the same neuronal population. There is evidence that striatal neurons in primary culture form functional synapses (Pin *et al.*, 1986; Weiss *et al.*, 1986; Dubinsky, 1989; Misgeld and Dietzel, 1989). However, the synapses appear to be predominantly GABAergic (Pin *et al.*, 1986; Misgeld and Dietzel, 1989), so that an indirect action of quinpirole to inhibit synaptic activity would be predicted to enhance immediate early gene induction rather than to suppress it. Furthermore we have found that bicuculline does not influence *c-fos* or *zif/268* mRNA levels (data not shown), suggesting that intrinsic GABAergic activity is not a major factor regulating immediate early gene expression in striatal neurons. Our results therefore lend further weight to the theory that medium spiny neurons express both D1-like and D2-like dopamine receptors, and that these two receptor classes are in functional opposition.

Chapter 7

Antisense Oligonucleotide Strategies

INTRODUCTION

7.0 Antisense Oligonucleotides

Single stranded oligonucleotides, complementary to a specific mRNA are called antisense oligonucleotides. Several investigators have used "antisense strategy" to block the synthesis of specific proteins (review by Dolnick, 1990; Vand der Krol *et al.*, 1988). Although the exact mechanism of action of antisense oligonucleotides is unknown, they can achieve a translational arrest of a target mRNA when hybridisation occurs *in vivo* or in a cell-free translation system. Effective antisense oligonucleotides are generally of short length (15-20 bases). It is believed that this length is short enough to be taken up by cells, but long enough to provide specificity of binding to a unique base sequence of mRNA (Zamecnik and Stephenson, 1978; Zamecnik *et al.*, 1986; Holt *et al.*, 1988; Van der Krol *et al.*, 1988; Dolnick, 1990). There is current evidence suggesting that the mechanism of oligonucleotide uptake is predominantly fluid-phase endocytosis (pinocytosis) (Stein and Cheng., 1993) which may be initiated by receptor-mediated recognition (Yakubov *et al.*, 1989). The mechanisms of potential oligonucleotide interactions with target nucleic acids are very complex. According to the most popular theories, the antisense oligonucleotides exert their effects by inducing translational arrest of protein synthesis either by stimulating ribonuclease H-mediated degradation (Walder and Walder, 1988) or preventing the translocation of the ribosome along the targeted mRNA sequence (Boiziau *et al.*, 1991).

Two types of antisense oligonucleotides are presently in use as experimental tools, either the unmodified phosphodiester oligonucleotides or the nuclease resistant phosphorothioate analogues. Different cell types have been shown to differ greatly with respect to oligonucleotide uptake as well as oligonucleotide

degradation. It has been suggested that many neuronal systems have a very low capacity to degrade oligonucleotides and that they are therefore amenable to treatment with either phosphodiester oligonucleotides or nuclease-stable analogues (Wahlestedt et al., 1993; Standifer et al., 1993; Akabayashi et al., 1994). It has previously been shown that phosphodiester oligonucleotides can enter cultured neurons at a concentration of up to 5% in the media (Caceres and Kosik, 1990). Effective treatment with antisense oligonucleotides is entirely dependent on the gene type under study and the turnover rate of the protein. In general, for receptor synthesis inhibition a minimum treatment time of two to five days has been shown necessary (Wahlestedt et al., 1993; Erlinge et al., 1993; Standifer et al., 1993).

The potential uses of antisense oligonucleotides has gradually become clear and a considerable focus on experimental use is currently in progress. In the past decade antisense oligonucleotides have been effective in inhibiting herpes simplex virus replication (Zamecnik and Stephenson, 1978) and in blocking the synthesis of human immunodeficiency viral proteins (Zamecnik et al., 1986). The antisense strategy has also been used in studying the function of cellular oncogenes and other proteins on cell proliferation and differentiation in vitro (Holt et al., 1988; Wickstrom et al., 1988; Becker et al., 1989; Tortora et al., 1990; Maier et al., 1990; Venturelli et al., 1990; Wang et al., 1992). In clonal cell lines antisense oligonucleotides have been used to target specific proteins involved in second messenger systems (Wagner et al., 1992). An antisense oligonucleotide has also been used against the NMDA receptor complex preventing NMDA receptor channel formation and functionally reducing neurotoxicity elicited by NMDA receptors in vitro (Wahlestedt et al., 1993). Both the muscarinic M2 and GABA_B receptor proteins have been successfully decreased by more than 50% using antisense oligonucleotides and thus has eliminated the ability of these receptors when stimulated to inhibit cyclic AMP formation (Holpainen and Wojcik, 1992). Only a few reports have

shown the effectiveness of antisense oligonucleotides in primary cultured neurons (Caceres and Kosk, 1990; Listerud et al., 1991).

Direct administration of antisense oligonucleotides *in vivo* has recently proved effective against certain IEGs and receptor proteins. Direct infusion into the striatum of an antisense oligonucleotide against the IEG *c-fos*, has been shown to reduce amphetamine-induced production of *fos*-like immunoreactivity, without affecting *zif/268* expression (Chiasson et al., 1992). Similarly, the ability of cocaine to induce *c-fos* protein in striatum and also locomotor stimulation was blocked when an antisense oligonucleotide against *c-fos* was injected into the nucleus accumbens (Heilig et al., 1993). Zhang and Creese (1993) have shown that an intracerebroventricular infusion of D2 antisense selectively decreases striatal D2 receptors by 50% in the rat, while not effecting D1, muscarinic and 5-HT2 receptor proteins. Direct infusion of D2 antisense has also been shown to inhibit rotational behaviour induced by quinpirole, a D2 agonist, but not that induced by the D1 agonist SKF38393, or the muscarinic agonist oxotremorine in the striatum of rats previously treated with 6-hydroxy dopamine (Weiss et al., 1993). The use of antisense strategy has also revealed a possible new subtype of D2 receptor on the basis of functional properties (Valerio et al., 1994).

The aim of these experiments was therefore to attempt to identify the functional dopamine receptor subtype, whose stimulation is involved in the induction of both *c-fos* and *zif/268* mRNA, in striatal neurons, after treatment with SKF38393, a non-selective agonist for both D1 and D5 dopamine receptor subtypes.

RESULTS

7.1 Effect of blockade of dopamine D1 and D5 receptor synthesis on the levels of the IEGs c-fos and zif/268 mRNA using antisense oligonucleotide strategies in rat striatal neuron cultures

The use of antisense oligonucleotides to arrest the translation of receptor mRNAs and thus reduce receptor synthesis has recently been employed to investigate receptor mediated responses. In this study we were interested in the nature of the receptor(s) involved in the induction of c-fos and zif/268mRNA after treatment of cultures with SKF38393, the selective D1/D5 receptor agonist.

Initially the dose range of both the D1 and D5 antisense oligonucleotides was determined. We found that doses of 2 μ M and above were toxic to the neurons in culture, whilst doses of 1 μ M appeared to be effective. Addition of SKF38393 (1nM-5 μ M) to cultured striatal neurons was demonstrated to increase the expression of both c-fos mRNA and zif/268 mRNA in a dose-dependent manner. The EC₅₀ for the SKF38393-induced stimulation of both c-fos and zif/268 mRNA was in the region of 95nM. There was no difference in the basal level of expression of c-fos mRNA in striatal cultures pretreated for 2 days with 15-mer antisense phosphorothioate oligonucleotides (1 μ M) complementary to a region of the dopamine D1 and D5 receptors alone compared to vehicle treated cultures. However, in cultures pretreated for 2 days with D1 antisense, followed by SKF38393 (1nM-5 μ M) treatment for 45 minutes, the slope of the dose response curve representing induction of c-fos mRNA had clearly shifted in a downward direction, implying that the D1 receptor is involved in this response. This shift in the dose response

curve to SKF38393 was demonstrated to be significantly different from the the dose response curve to SKF38393 alone ($F= 118.64$, $p< 0.001$, ANOVA minitab, $n= 3$). [Figure 7.1.1]. Less of a shift was observed in the presence of the dopamine D5 receptor antisense oligonucleotide, however, this overall shift was significantly different from the dose response curve to SKF38393 alone ($F= 18.58$, $p< 0.001$) [Figure 7.1.2]. As an appropriate control for the effects of the D1 and D5 receptor antisense oligonucleotides, a 15-mer D1 'sense' phosphorothioate oligonucleotide was synthesised, having the same sequence as a region of the D1 receptor mRNA. Pretreatment of the cultures with a 15-mer D1 sense oligonucleotide ($1\mu\text{M}$) for two days , followed by acute treatment with SKF38393 (1nM - $5\mu\text{M}$) for 45 minutes had no effects on the induction of c-fos mRNA by SKF38393 [Figure 7.1.3]. However, the dose response curve in the presence of the D1 sense oligonucleotide was significantly different from the dose response curve to SKF38393 in the presence of the D1 antisense oligonucleotide ($F=22.98$, $p< 0.05$) and similarly was significantly different from the curve to SKF38393 in the presence of the D5 antisense ($F= 18.17$, $p< 0.05$) [Figure 7.1.3].

The induction of zif/268 mRNA following SKF38393 treatment showed a similar dose dependent profile, with an EC_{50} of 95nM . However, when cultures were pretreated for two days with the D1 'sense' oligonucleotide, the dose response curve to SKF38393 (1nM - $5\mu\text{M}$) was clearly different from SKF38393 (1nM - $5\mu\text{M}$) alone, such that $F= 37.55$, $p< 0.001$, $n= 3$ [Figure 7.1.4]. This would suggest that the sense oligonucleotide can effect the ability of SKF38393 to induce zif/268 under these conditions. Pretreatment with the D1 antisense oligonucleotide ($1\mu\text{M}$) also caused a significant shift in the SKF38393 dose response curve ($F= 81.57$, $p< 0.001$, $n= 3$) [Figure 7.1.5] , and

similarly pretreatment with the D5 antisense oligonucleotide ($1\mu\text{M}$) resulted in a significant shift in the dose response curve to SKF38393 (1nM - $5\mu\text{M}$) ($F=159.51$, $p<0.001$, $n=3$) [Figure 7.1.6]. The effects observed with the sense oligonucleotide suggests that the reduction in response seen with the two antisense oligonucleotides is likely to be non-specific.

Figure 7.1.1 : Effect of a dopamine D1 antisense oligonucleotide on c-fos mRNA levels in striatal cultures. Cells were pretreated for two days with a 15-mer phosphorothioate oligonucleotide corresponding to a D1 dopamine receptor antisense (1 μ M, n= 3). Cells were then exposed to SKF38393 (1nM-5 μ M, n= 5) alone or in the presence of D1 antisense (n= 3), or vehicle (HBSS) alone or in the presence of D1 antisense. After 45 minutes the cells were fixed in buffered formalin and then processed for in situ hybridisation with a specific radiolabelled probe for c-fos. Results are expressed as mean number of pixels per unit area \pm s.e.m. The dose response curve in the presence of the D1 antisense was significantly different from SKF38393 treatment alone $p < 0.001^{***}$ (two-way ANOVA, Minitab).

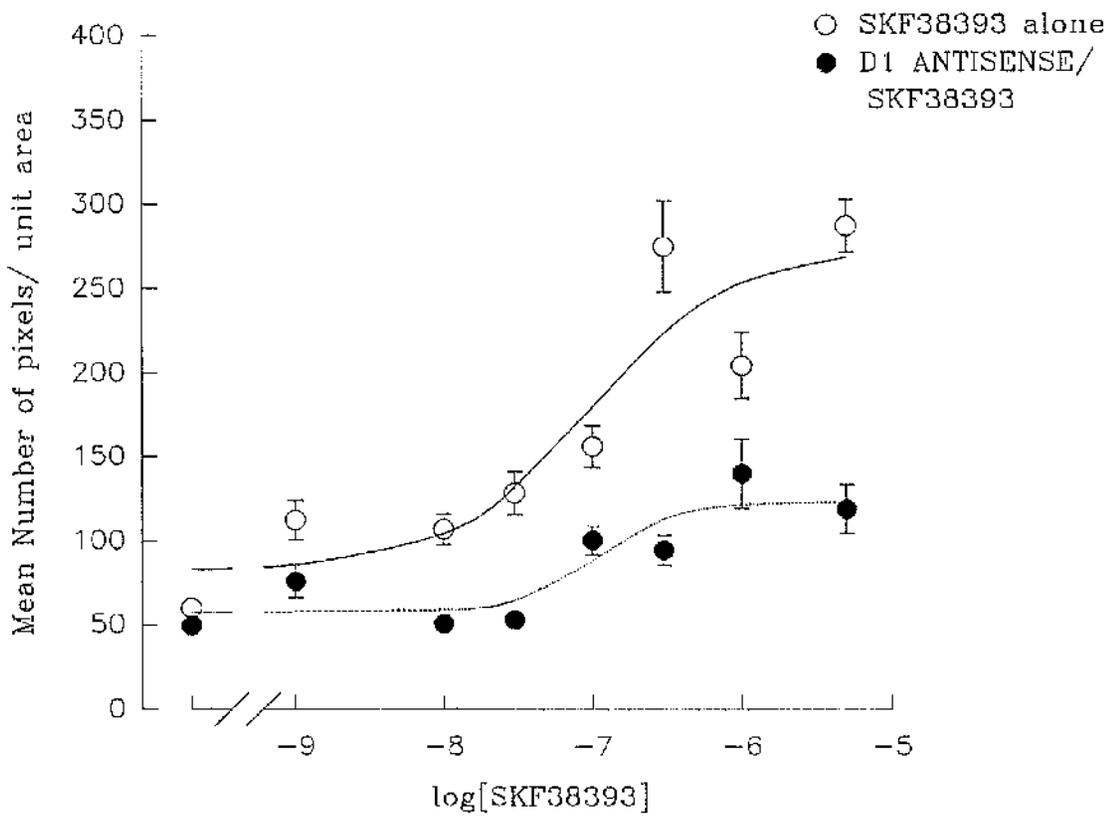


Figure 7.1.2 : Effect of a dopamine D5 antisense oligonucleotide on c-fos mRNA levels in striatal cultures. Cells were pretreated for two days with a 15-mer phosphorothioate oligonucleotide corresponding to a D5 dopamine receptor antisense ($1\mu\text{M}$, $n=3$). Cells were then exposed to SKF38393 (1nM - $5\mu\text{M}$, $n=5$) alone or in the presence of D5 antisense ($n=3$), or vehicle (HBSS) alone or in the presence of D5 antisense. After 45 minutes the cells were fixed in buffered formalin and then processed for in situ hybridisation with a specific radiolabelled probe for c-fos. Results are expressed as mean number of pixels per unit area \pm s.e.m. The dose response curve in the presence of the D5 antisense was significantly different from SKF38393 treatment alone $p < 0.001^{***}$ (two-way ANOVA, Minitab).

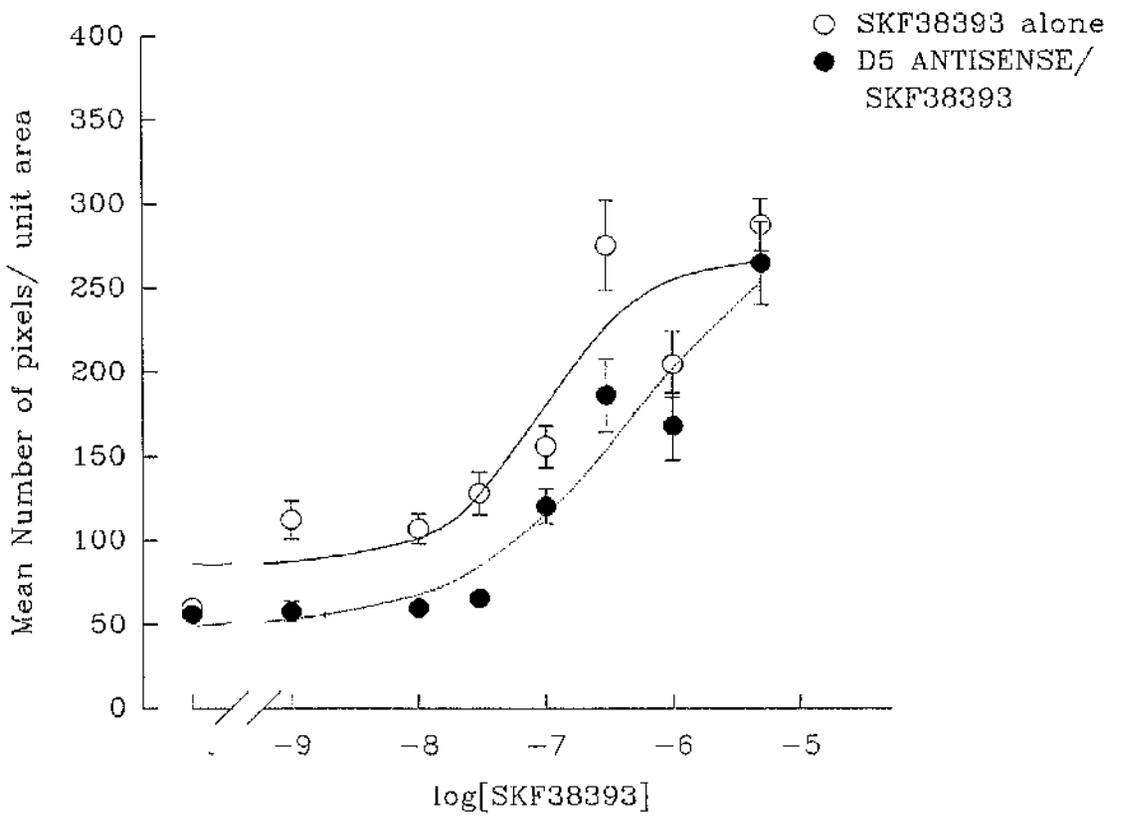


Figure 7.1.3 : Effect of a dopamine D1 sense oligonucleotide on c-fos mRNA levels in striatal cultures. Cells were pretreated for two days with a 15-mer phosphorothioate oligonucleotide corresponding to a D1 dopamine receptor sense probe (1 μ M, n= 3). Cells were then exposed to SKF38393 (1nM-5 μ M, n= 5) alone or in the presence of D1 sense (n= 3), or vehicle (HBSS) alone or in the presence of D1 sense. After 45 minutes the cells were fixed in buffered formalin and then processed for in situ hybridisation with a specific radiolabelled probe for c-fos. Results are expressed as mean number of pixels per unit area \pm s.e.m. The dose response curve in the presence of the D1 sense was not significantly different from SKF38393 treatment alone (two-way ANOVA, Minitab).

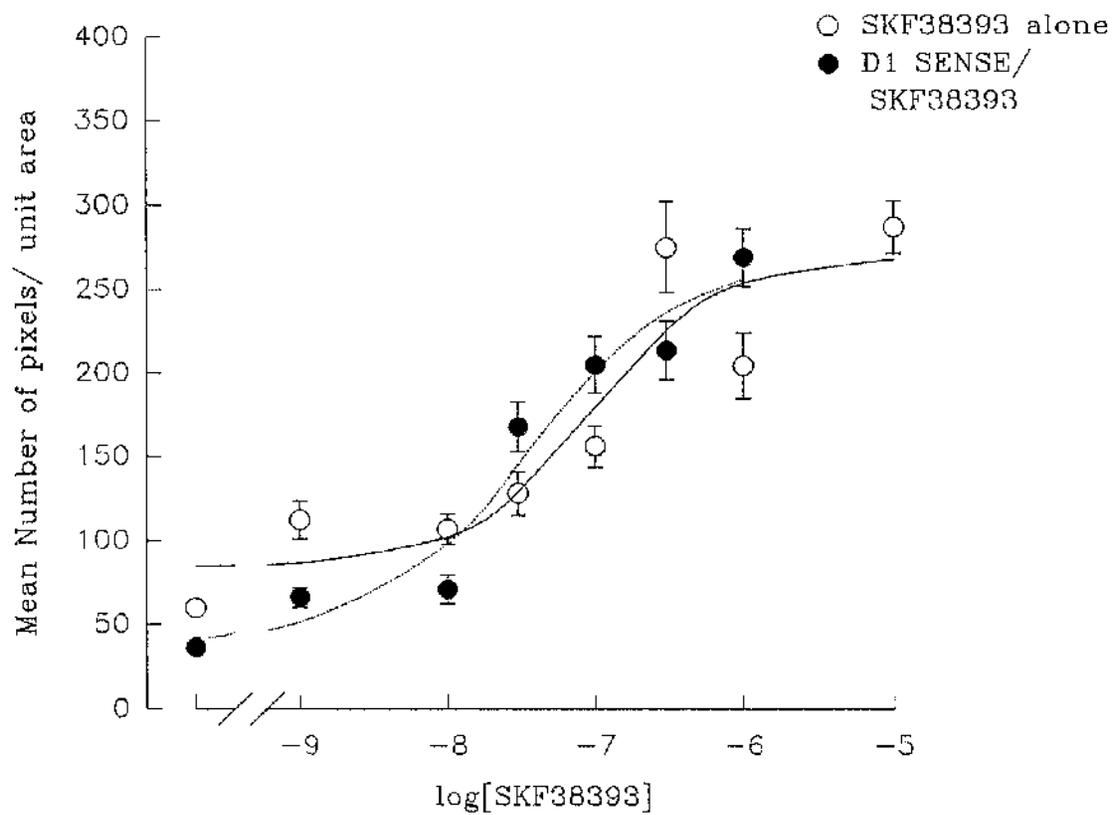


Figure 7.1.4 : Effect of a dopamine D1 sense oligonucleotide on *zif/268* mRNA levels in striatal cultures. Cells were pretreated for two days with a 15-mer phosphorothioate oligonucleotide corresponding to a D1 dopamine receptor sense probe ($1\mu\text{M}$, $n= 3$). Cells were then exposed to SKF38393 (1nM-5 μM , $n= 5$) alone or in the presence of D1 sense ($n= 3$), or vehicle (HBSS) alone or in the presence of D1 sense. After 45 minutes the cells were fixed in buffered formalin and then processed for in situ hybridisation with a specific radiolabelled probe for *zif/268*. Results are expressed as mean number of pixels per unit area \pm s.e.m. The dose response curve in the presence of the D1 sense was significantly different from SKF38393 treatment alone $p < 0.001$ (two-way ANOVA, Minitab).

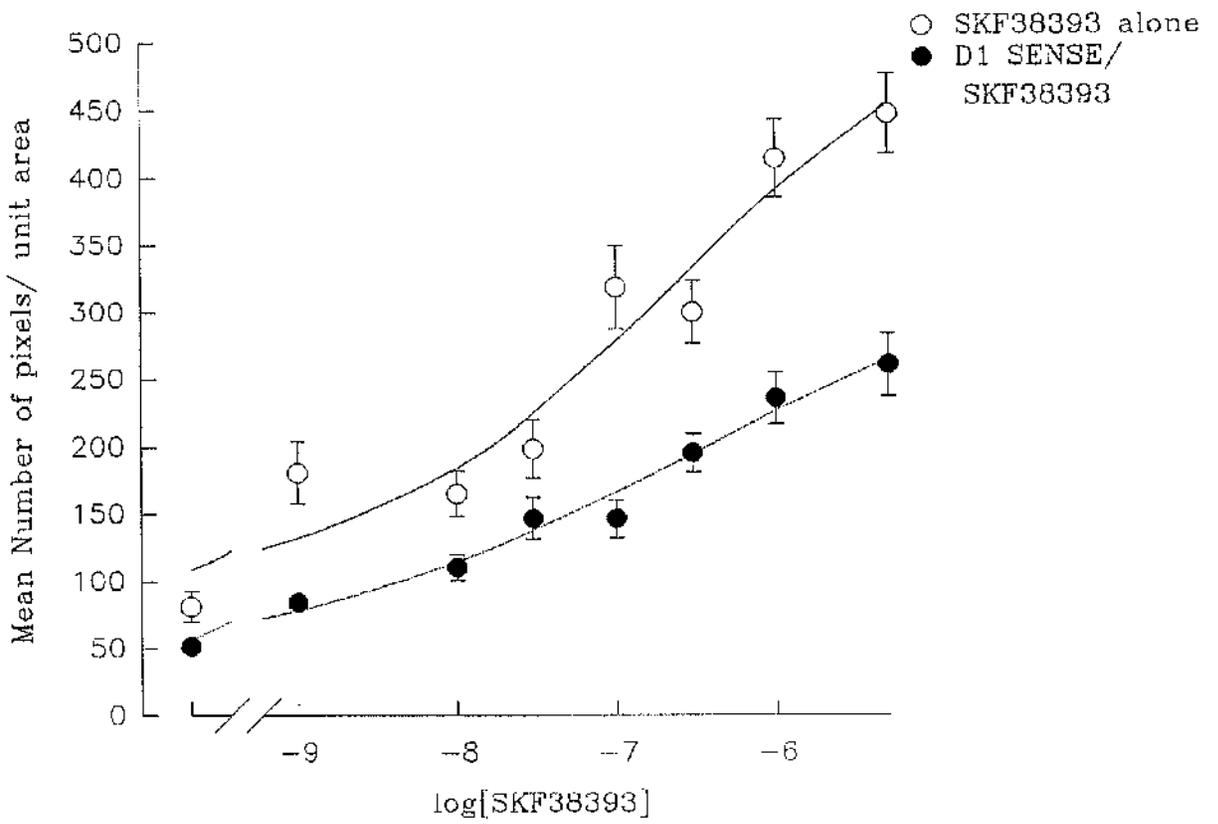


Figure 7.1.5 : Effect of a dopamine D1 antisense oligonucleotide on *zif/268* mRNA levels in striatal cultures. Cells were pretreated for two days with a 15-mer phosphorothioate oligonucleotide corresponding to a D1 dopamine receptor antisense (1 μ M, n= 3). Cells were then exposed to SKF38393 (1nM-5 μ M, n= 5) alone or in the presence of D1 antisense (n= 3), or vehicle (HBSS) alone or in the presence of D1 antisense. After 45 minutes the cells were fixed in buffered formalin and then processed for in situ hybridisation with a specific radiolabelled probe for *zif/268*. Results are expressed as mean number of pixels per unit area \pm s.e.m. The dose response curve in the presence of the D1 antisense was significantly different from SKF38393 treatment alone $p < 0.001$ *** (two-way ANOVA, Minitab).

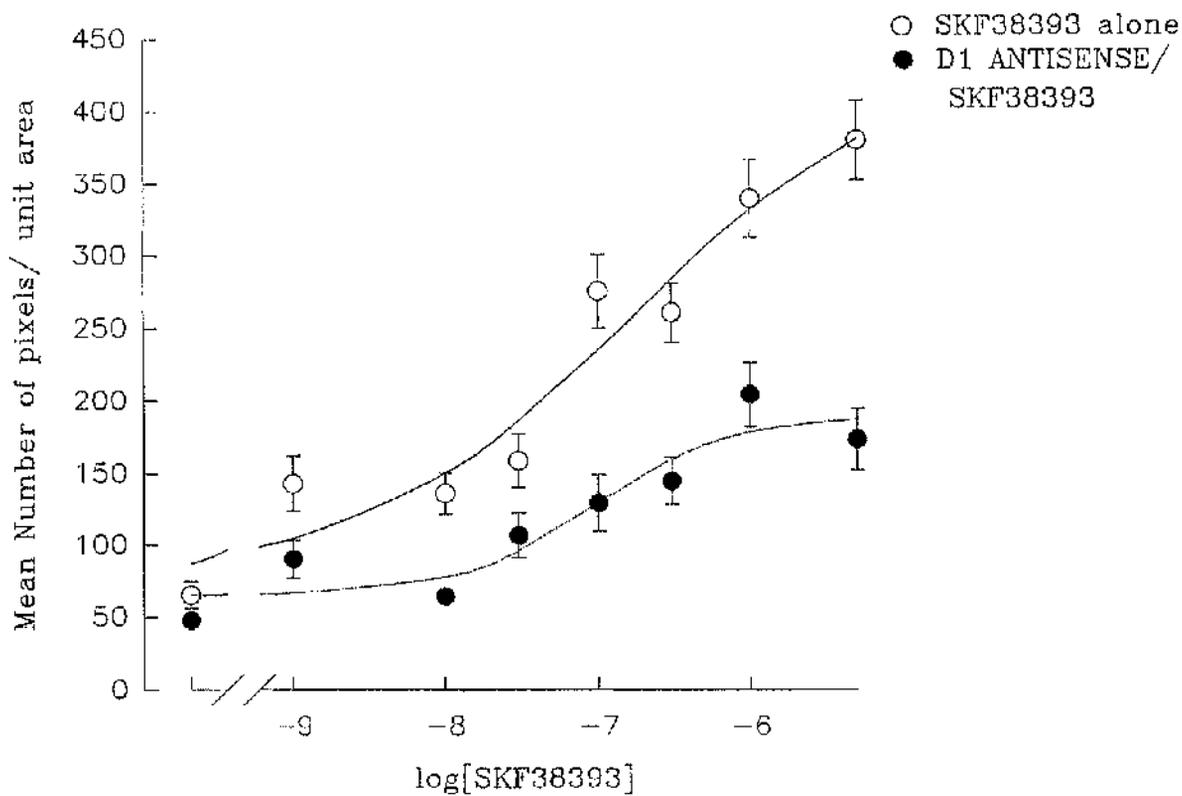
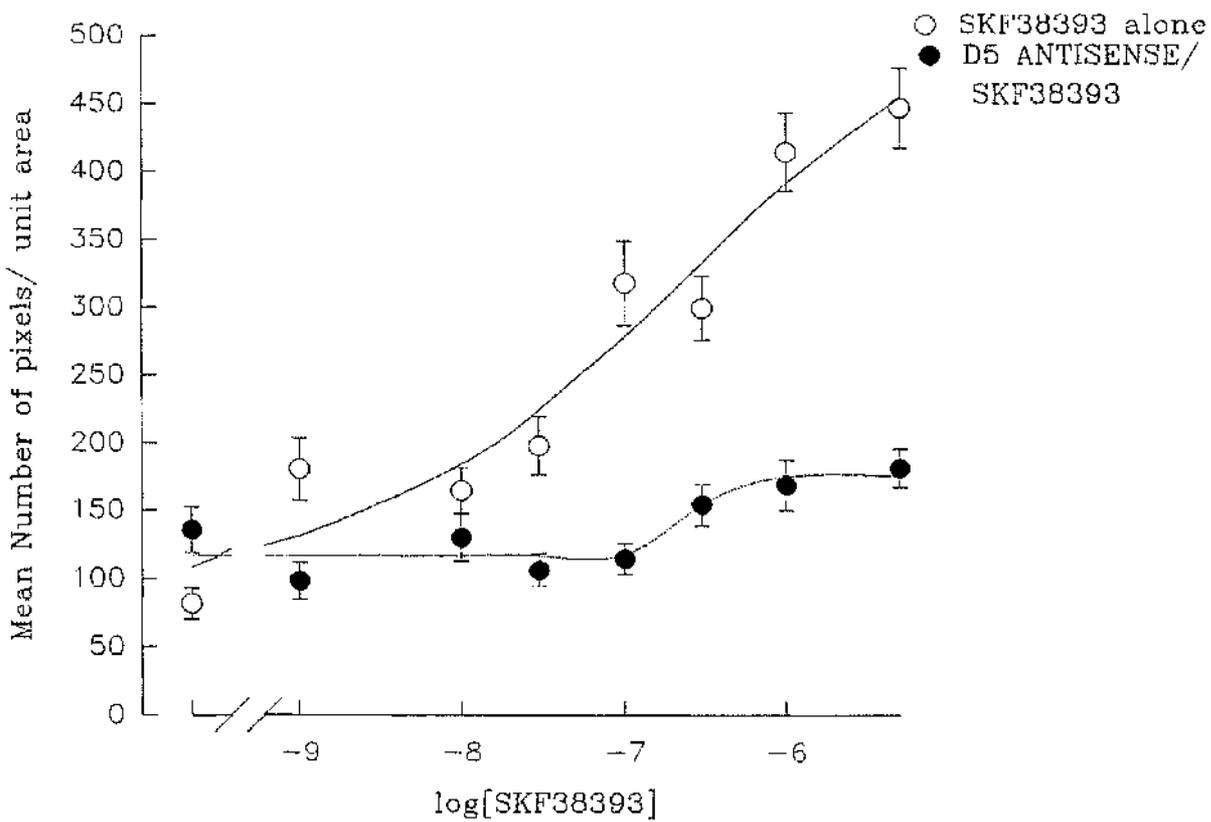


Figure 7.1.6 : Effect of a dopamine D5 antisense oligonucleotide on zif/268 mRNA levels in striatal cultures. Cells were pretreated for two days with a 15-mer phosphorothioate oligonucleotide corresponding to a D5 dopamine receptor antisense (1 μ M, n= 3). Cells were then exposed to SKF38393 (1nM-5 μ M, n= 5) alone or in the presence of D5 antisense (n= 3), or vehicle (HBSS) alone or in the presence of D5 antisense. After 45 minutes the cells were fixed in buffered formalin and then processed for in situ hybridisation with a specific radiolabelled probe for zif/268. Results are expressed as mean number of pixels per unit area \pm s.e.m. The dose response curve in the presence of the D5 antisense was significantly different from SKF38393 treatment alone $p < 0.001$ *** (two-way ANOVA, Minitab).



DISCUSSION

7.2 Effect of dopamine receptor antisense oligonucleotides on immediate early gene expression in striatal cultures

In previous results we have demonstrated that acute treatment with SKF38393 can induce the expression of the immediate early genes *c-fos* and *zif/268* in primary embryonic rat striatal cultures. This is in agreement with *in vivo* work where the induction of *c-fos* by SKF38393 has been shown, following a unilateral 6-hydroxydopamine lesion of the nigrostriatal pathway (Robertson *et al.*, 1989; Robertson *et al.*, 1992) and with *in vitro* research demonstrating the induction of *c-fos* with SKF38393 treatment in cerebral cortical cultures (Vaccarino *et al.*, 1993).

SKF38393, a potent D1/ D5 dopamine receptor agonist, is thought to be producing this stimulatory effect on both *c-fos* and *zif/268* gene expression by activating D1-like receptors *in vivo* and *in vitro*. Until now it has been impossible to distinguish D1-mediated effects from D5-mediated effects, simply because there are no selective ligands to differentiate between these two highly homologous dopamine receptor subtypes. However, with the introduction of antisense oligonucleotides targeted towards particular receptor mRNAs, it is now possible to achieve a substantial degree of protein synthesis inhibition. Two 15-nucleotide-long antisense oligonucleotides were designed to recognize a sequence of the D1 and D5 dopamine receptor genes respectively, in an attempt to answer this question more fully.

In this study we have demonstrated that *c-fos* mRNA can be dose-dependently increased by SKF38393 *in vitro*, with an EC_{50} value of approximately 110nM. This EC_{50} value obtained is very close to the K_D values previously reported for the binding of SKF38393 to striatal membranes

(Seeman and Van Tol, 1993). The levels of c-fos mRNA reached a plateau at around 5 μ M SKF38393. Pretreatment of primary striatal cultures with a D1 sense phosphorothioate oligonucleotide, at a concentration not causing toxic effects, demonstrated by no change in D2 receptor binding and no obvious change in cell numbers, clearly had no effect on the dose-dependent induction of c-fos gene expression by SKF38393, in rat striatal cultures. This would be expected, because the sense oligonucleotide is acting as a control here, in that it has the same sequence as part of the D1 receptor mRNA and therefore no binding interaction to cellular mRNA will occur if conditions are optimal, with non-specific binding reduced to a minimum. In contrast, pretreatment with the D1 phosphorothioate antisense oligonucleotide, at a concentration not causing toxicity, was clearly able to reduce the dose-dependent induction of c-fos mRNA by SKF38393. The D1 antisense caused a greater than 50% reduction of the maximum effect of SKF38393.

Similarly pretreatment with the phosphorothioate antisense oligonucleotide directed towards the D5 dopamine receptor protein reduced the overall effect of SKF38393, particularly at low concentrations of SKF38393. However, the maximum effect of SKF38393 was unaltered by pretreatment with the D5 antisense. In general, these results are suggesting that the D1 dopamine receptor primarily mediates the inductive effect of SKF38393 on c-fos gene expression in the rat striatum. However, there may be a smaller D5 component to this effect. It is known that the density of D5 dopamine receptors in the striatum is much lower than the density of D1 dopamine receptors, and that the affinity for the endogenous ligand dopamine, is ten-fold higher for the D5 receptor (Van Tol *et al.*, 1991). These properties of the D5 receptor may be thought to functionally represent a 'fine control' over the dopaminergic system. Therefore stimulation of D1 and D5 dopamine receptors may contribute to the induction of c-fos by SKF38393. The results also indicate

that in rat primary striatal cultures, in the case of D1 receptors, there appears to be little receptor reserve.

The results obtained for the SKF38393- induced expression of *zif/268* mRNA in the presence of the antisense oligonucleotides are much more difficult to interpret. Similar to the results obtained for *c-fos*, a dose-dependent increase in *zif/268* mRNA was observed following acute treatment with SKF38393 alone, with a very similar EC_{50} value to that obtained for *c-fos* induction, of around 100nM. However, pretreatment with the D1 sense oligonucleotide clearly inhibited the dose-dependent effect of SKF38393 on *zif/268* gene expression. This non-specific inhibitory effect of the D1 sense makes any effect of the D1 or D5 antisense very difficult to interpret as a specific effect. Therefore the effects of the dopamine D1 and D5 antisense on *zif/268* expression after SKF38393 are likely to be non-specific in these experiments.

One of the major disadvantages in the use of 'knockdown' approaches such as antisense oligonucleotide inhibition, is the narrow experimental and therapeutic windows that accompany antisense use. With the D1 and D5 antisense a concentration of 1 μ M proved effective in reducing the SKF38393 effect on *c-fos* gene expression by 50%. However, we found that concentrations of 2 μ M and above proved toxic to the cells in primary striatal culture. Therefore the existence of a narrow window for effective concentration may not always parallel the therapeutic window. In particular, potential problems are associated with an incomplete effect of oligonucleotide treatment. If a biological phenomenon has a great deal of spare capacity or redundancy, for example, then only a fraction of receptors have to be occupied to elicit a full response, and the oligonucleotide treatment may become functionally 'silent'. In this study it would also be necessary to demonstrate reductions in D1-like receptor binding, and to rule out oligonucleotide effects on proteins other than the targeted D1 and D5 receptor proteins, by performing

appropriate ligand-binding experiments, which are at present still in progress in our laboratory. Further appropriate controls for these experiments would involve the use of random sequence oligonucleotides and mismatch oligonucleotides.

However, the results from this study are the first to provide an insight into the specific receptor mechanism involved in the inductive effect of SKF38393 on *c-fos* expression in the rat striatum. Antisense oligonucleotides targeted towards the D1 and D5 dopamine receptor proteins have been used to reduce the respective receptor protein and thus establish which dopamine receptor subtype(s) contribute to the stimulatory effect of SKF38393 on *c-fos* gene expression in rat striatal culture. We have produced evidence to suggest that D1 receptor stimulation is primarily involved in this effect, with a minor D5 component. However, the evidence that non-specific effects occur on another cellular process- the induction of *zif/268* expression, suggests that extreme caution should always be observed when using antisense oligonucleotides.

Chapter 8

Serotonergic Regulation of Immediate Early Gene Expression in vitro

INTRODUCTION

8.0 Serotonergic Modulation of Immediate Early Gene Expression in the Central Nervous System

The serotonergic system is involved in numerous physiological functions. In the central nervous system, serotonin (5-HT) is known to be involved in the homeostatic regulation of fundamental physiological functions such as sleep, appetite, locomotion, nociception, sexual activity and vascular contraction (Wilkinson and Dourish, 1991). Disruption of the serotonergic system has also been implicated as a critical factor in mental disorders such as schizophrenia, anxiety, depression, parkinsonism and obsessive-compulsive disorders (Lopez-Ibor, 1988). This biogenic amine neurotransmitter is synthesised by neurons in the raphe nuclei of the brain stem that project throughout the central nervous system, with the highest density in basal ganglia and limbic structures (Steinbusch, 1984).

At present, at least seven classes of serotonin receptor have been cloned, each class being supported by a number of receptor subtypes. The functional significance, if any, in terms of the physiological role of many of these newly cloned receptors is as yet to be identified and therefore is of great importance in understanding the role of 5-HT as a regulator of many behavioural responses.

The relatively new field of research into immediate early genes is currently expanding. However, there is very little research, as yet, into the serotonergic modulation of immediate early genes in the central nervous system. Recent work has demonstrated, using cultured spinal cord neurons, that the induction of the immediate early gene *c-fos* and subsequent transactivation of the prodynorphin gene, are coupled events triggered by 5-

HT1A receptor agonists (Lucas *et al.*, 1993). Classical 5-HT1A receptor agonists are known to be inhibitory, although this induction of c-fos following a 5-HT1A receptor stimulation also resulted in a simultaneous increase in cAMP levels. This would appear to be an anomaly from the classical cloned 5-HT1A receptor subtype, suggesting the possibility of another subtype of the 5-HT1A receptor.

The serotonergic system has also been implicated in the regulation of immediate early genes in the striatum. Recent *in vivo* research has suggested that the ability of cocaine to induce the expression of c-fos and zif/268 in brain neurons is partly due to its characteristic property of 5-HT uptake inhibition. Using the selective inhibitors of 5-HT uptake, fluoxetine and citalopram, the ability of mazindol, a dopamine and noradrenaline uptake blocker, to induce zif/268 and c-fos was shown to be potentiated (Bhat and Baraban, 1993). More recently, activation of the serotonin 2A/2C receptor has been demonstrated to cause a localised expression of the immediate early gene c-fos in the frontal cortex, amygdala, globus pallidus, nucleus accumbens and dorsomedial striatum, and this induction could be selectively blocked with the 2A/2C antagonist ritanserin *in vivo* (Leslie *et al.*, 1993). There is also some evidence for the serotonergic regulation of tachykinin gene expression in the rat neostriatum. A recent report has shown that by raising extracellular levels of serotonin with zimelidine, a 5-HT uptake inhibitor, the levels of preprotachykinin mRNA were increased (Walker *et al.*, 1991). This suggests that neostriatal tachykinin biosynthesis is sensitive to alterations in 5-HT neurotransmission.

At present there are several suggestions with regard to the involvement of central serotonergic mechanisms in the actions of antipsychotic drugs. The recently cloned 5-HT6 and 5-HT7 receptor subtypes have been demonstrated to exhibit a particularly high affinity for some atypical neuroleptic drugs, including clozapine. The mRNA encoding the 5-HT6 receptor has been shown

to be mainly present in the striatum, olfactory tubercles, nucleus accumbens and in the hippocampus (Monsma *et al.*, 1991). More recently it has been demonstrated that the 5-HT6 receptor is present in primary striatal neuron cultures and it is positively coupled to adenylyl cyclase activity (Sebben *et al.*, 1994). Similarly the 5-HT7 receptor subtype has also been shown to be positively coupled to adenylyl cyclase (Tsou *et al.*, 1994). Therefore the 5-HT6 and 5-HT7 receptor subtypes may have a functional role in the manifestation and clinical treatment of certain disease states such as schizophrenia and parkinsonism.

Therefore, it was decided to investigate the role of serotonin in the induction of the immediate early genes *c-fos* and *zif/268*, and to elucidate the receptor mechanism(s) involved.

RESULTS

Much research has been done on the dopaminergic regulation of IEGs in the striatum. However, at present there is very little information on the serotonergic modulation of IEGs in the striatum. Although there are implications for a serotonergic stimulation of *zif/268* in the striatum (Bhat *et al.*, 1993). Therefore in this study we were interested in the serotonergic modulation of the IEGs *zif/268* and *c-fos* in rat striatal neuronal cultures and the receptor site(s) involved.

8.1 Effect of serotonin (5-HT) on the expression of the IEGs *c-fos* and *zif/268* mRNA

Serotonin (5-HT) (1nM-5 μ M) was demonstrated to clearly increase the expression of both *zif/268* and *c-fos* mRNA in striatal cultures in a dose dependent manner. The induction of *zif/268* mRNA forty-five minutes after 5-HT (300nM) can be seen in figure 8.1.19, as an increase in the number of silver grains over cell bodies in striatal culture, compared to vehicle treated cultures. The EC₅₀ of 5-HT to stimulate both *zif/268* and *c-fos* mRNA in striatal culture was approximately 100nM. The induction occurred in a large proportion of the cultured cells, suggesting that the increase in mRNA was observed mainly in medium spiny neurons [Figure 8.1.19]. Pretreatment of striatal cultures with ondansetron (200nM), the selective 5-HT₃ receptor antagonist, clearly had no effect on the levels of *zif/268* mRNA being expressed in culture after addition of 5-HT (1nM-5 μ M). [Figure 8.1.1]. Although *c-fos* mRNA levels due to stimulation by 5-HT were effected by the presence of ondansetron (200nM)

and this was shown to be significantly different from 5-HT (1nM-5 μ M) treatment alone ($F= 9.73$, $p < 0.01$; $n= 3$). [Figure 8.1.2].

The cultures were also pretreated with a number of 5-HT_{1/2} receptor antagonists including cyproheptadine, methiothepin, mesulergine and spiperone. It was found that cyproheptadine (500nM) caused a dramatic reduction in the hybridisation signal representing the induction of *zif/268* mRNA by 5-HT, and this was demonstrated to be significantly different from the expression of *zif/268* mRNA due to 5-HT (1nM-5 μ M) alone ($F= 29.95$, $p < 0.001$; $n= 3$) [Figure 8.1.3, 8.1.19]. A similar reduction in the signal was observed for *c-fos* mRNA ($F= 19.57$, $p < 0.001$; $n= 3$) [Figure 8.1.4]. Pretreatment with methiothepin (200nM) clearly decreased the hybridisation signal for *zif/268* mRNA in culture and was significantly different from 5-HT alone ($F= 39.95$, $p < 0.001$; $n= 3$) [Figure 8.1.5]. A reduction in the hybridisation signal for *c-fos* mRNA was also observed ($F= 33.89$, $p < 0.001$; $n= 3$) [Figure 8.1.6]. In cultures treated with mesulergine (100nM) prior to the addition of 5-HT (1nM-5 μ M) a marked decrease in *zif/268* mRNA was observed and this was significantly different from 5-HT (1nM-5 μ M) alone ($F= 28.71$, $p < 0.001$; $n= 3$) [Figure 8.1.7]. Pretreatment with the lower concentration of 30nM mesulergine also resulted in a highly significant decrease in the levels of *zif/268* mRNA in culture ($F= 17.36$, $p < 0.001$; $n= 3$) following a 5-HT-induced stimulation [Figure 8.1.8]. Mesulergine (100nM) caused a reduction in the levels of *c-fos* mRNA as a result of a 5-HT receptor stimulation. This decrease proved to be significantly different ($F= 7.10$, $p < 0.01$; $n= 3$) from 5-HT alone (1nM-5 μ M) [Figure 8.1.9]. Spiperone (100nM) pretreatment was also demonstrated to cause a massive inhibition of the 5-HT-induced stimulation of *zif/268* mRNA in striatal cultures, and this decrease was shown to be significant ($F= 59.36$, $p < 0.001$; $n= 3$) compared to 5-HT (1nM-5 μ M) alone [Figure 8.1.10], as did the lower dose of 30nM spiperone ($F= 31.96$, $p < 0.001$; $n= 3$) [Figure 8.1.11, 8.1.19]. As spiperone has very low

affinity for the 5-HT_{1C} receptor, the low dose used in this study would suggest that the increase in zif/268 mRNA observed, is not due to stimulation of the 5-HT_{2C} receptor. Pretreatment with spiperone (30nM) also resulted in a dramatic and highly significant inhibition of the hybridisation signal for c-fos mRNA ($F= 40.86$, $p<0.001$; $n= 3$) compared to 5-HT treatment alone [Figure 8.1.12].

The effects of a number of 5-HT receptor agonists were investigated in striatal culture. Addition of the 5-HT_{2C} receptor agonist mCPP (1nM-5 μ M) had no overall effects on the levels of both zif/268 mRNA [Figure 8.1.13] and c-fos mRNA [Figure 8.1.14]. Similarly the selective 5-HT_{1A} receptor agonist 8OH-DPAT (1nM-5 μ M) had no significant effect on the expression of the zif/268 [Figure 8.1.15] and c-fos genes [Figure 8.1.16]. Finally the selective 5-HT_{2A/2C} receptor agonist, \pm - DOI (1nM-5 μ M) was shown to have a slight effect on the levels of zif/268 and c-fos mRNA in striatal culture, but this induction did not prove to be significant [Figures 8.1.17 and 8.1.18 respectively].

Figure 8.1.1 : Effect of selective inhibition of the serotonergic 5-HT₃ receptor on the levels of expression of zif/268 mRNA in rat striatal neuron cultures. Cultures were pretreated for 30 minutes with ondansetron (200nM) prior to the addition of vehicle or 5-HT (1nM-5 μ M) (n= 3) or treated with vehicle or 5-HT (1nM-5 μ M) alone (n= 8). Results, obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m. Significance was determined by ANOVA, General Linear Model, Minitab. The two dose response curves were not significantly different.

Figure 8.1.2 : Effect of selective inhibition of the serotonergic 5-HT₃ receptor on the levels of expression of c-fos mRNA in rat striatal neuron cultures. Cultures were pretreated for 30 minutes with ondansetron (200nM) prior to the addition of vehicle or 5-HT (1nM-5 μ M) (n= 3) or treated with vehicle or 5-HT (1nM-5 μ M) alone (n= 7). Results, obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m. Significance was determined by ANOVA, General Linear Model, Minitab. The two dose response curves were significantly different, $p < 0.01$ **.

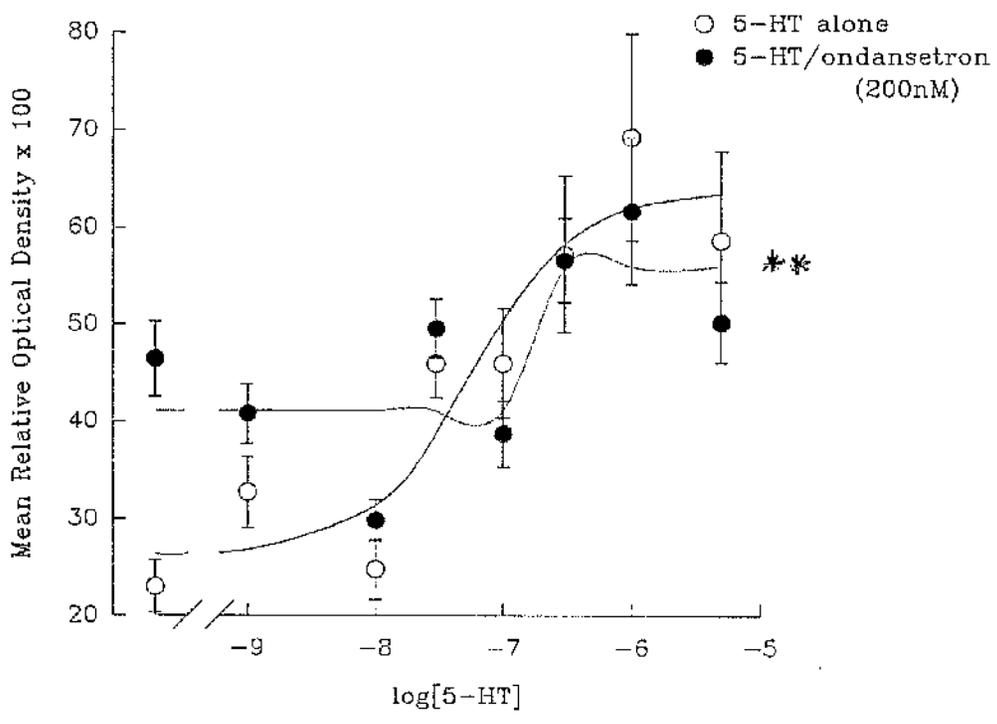
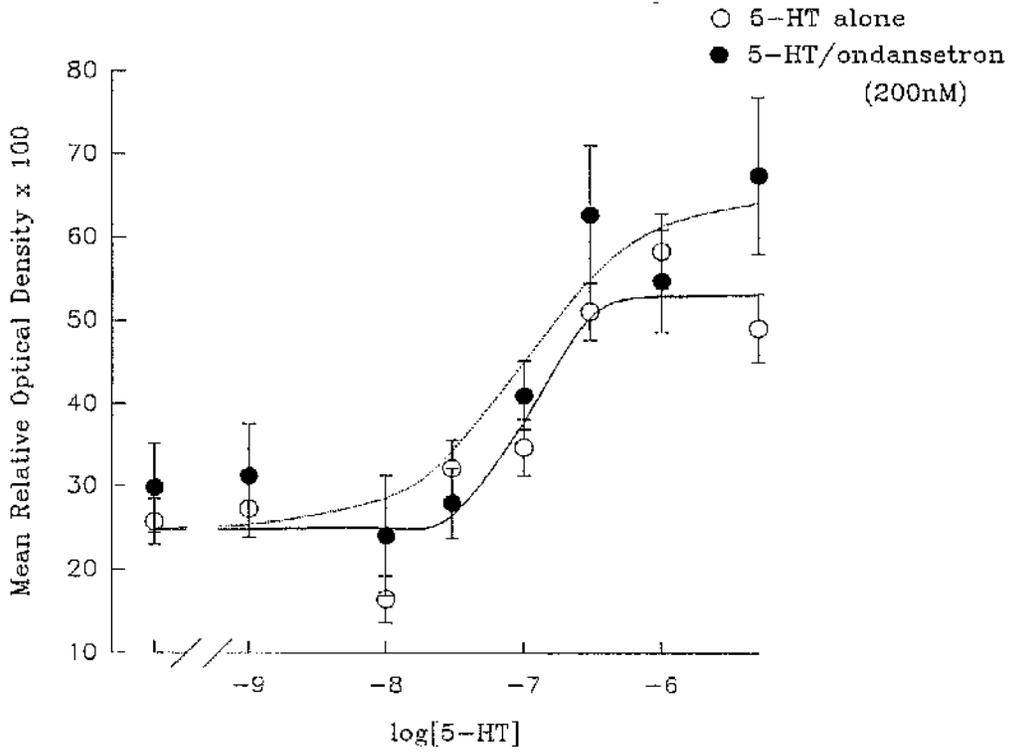


Figure 8.1.3 : Effect of inhibition of serotonergic 5-HT_{1/2} receptors on the levels of expression of zif/268 mRNA in rat striatal neuron cultures. Cultures were pretreated for 30 minutes with cyproheptadine (500nM), prior to the addition of vehicle or 5-HT (1nM-5 μ M) (n= 3) or treated with vehicle or 5-HT (1nM-5 μ M) alone (n= 8). Results obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m. Significance was determined by ANOVA, General Linear Model, Minitab. The two dose response curves were significantly different, $p < 0.001$ ***.

Figure 8.1.4 : Effect of inhibition of serotonergic 5-HT_{1/2} receptors on the levels of expression of c-fos mRNA in rat striatal neuron cultures. Cultures were pretreated for 30 minutes with cyproheptadine (500nM), prior to the addition of vehicle or 5-HT (1nM-5 μ M) (n= 3) or treated with vehicle or 5-HT (1nM-5 μ M) alone (n= 7). Results obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m. Significance was determined by ANOVA, General Linear Model, Minitab. The two dose response curves were significantly different, $p < 0.001$ ***.

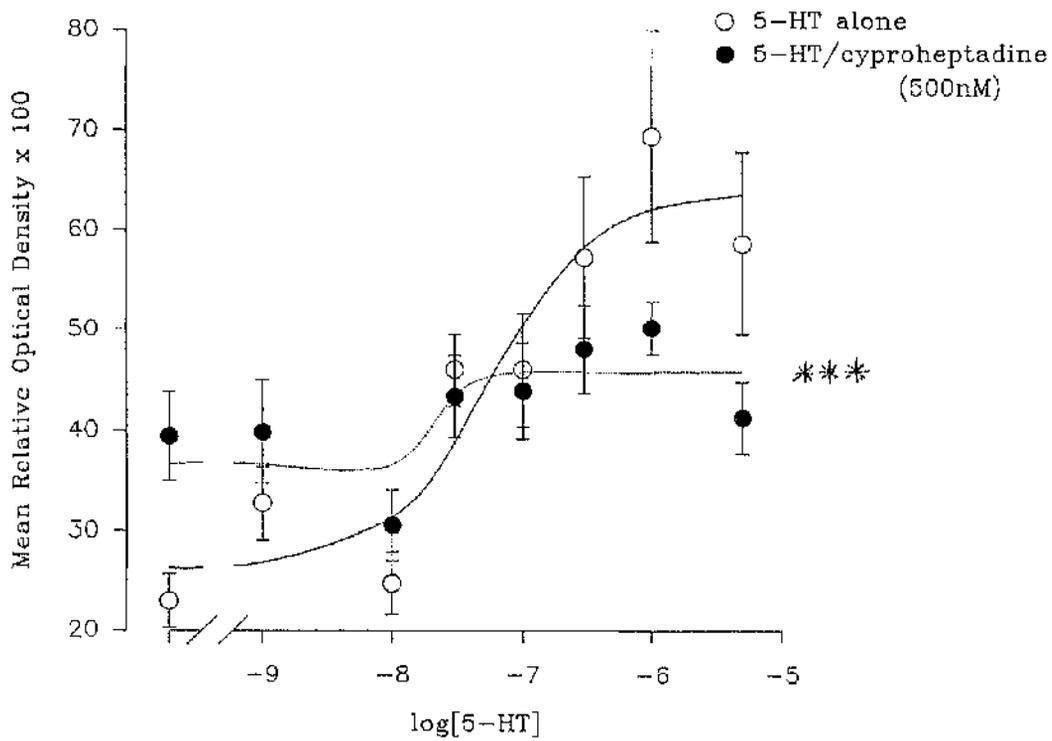
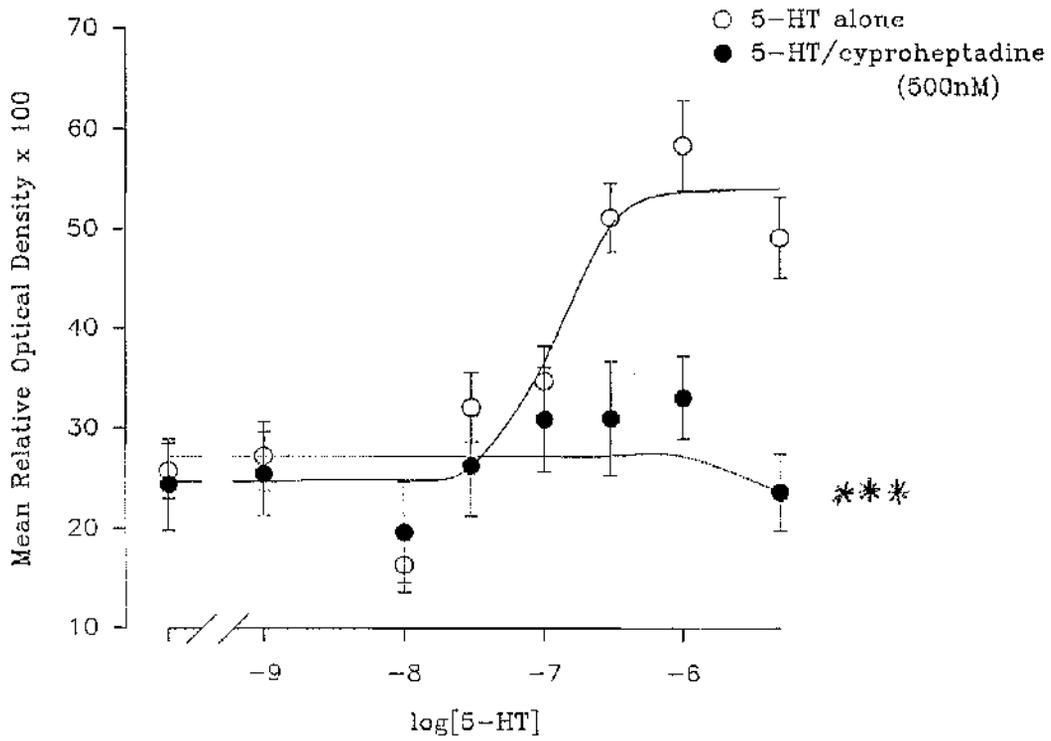


Figure 8.1.5 : Effect of inhibition of serotonergic 5-HT_{1/2/6} receptors on the levels of expression of zif/268 mRNA in rat striatal neuron cultures. Cultures were pretreated for 30 minutes with methiothepin (200nM), prior to the addition of vehicle or 5-HT (1nM-5 μ M) (n= 3) or treated with vehicle or 5-HT (1nM-5 μ M) alone (n= 8). Results obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m. Significance was determined by ANOVA, General Linear Model, Minitab. The two dose response curves were significantly different, $p < 0.001$ ***.

Figure 8.1.6 : Effect of inhibition of serotonergic 5-HT_{1/2/6} receptors on the levels of expression of c-fos mRNA in rat striatal neuron cultures. Cultures were pretreated for 30 minutes with methiothepin (200nM), prior to the addition of vehicle or 5-HT (1nM-5 μ M) (n= 3) or treated with vehicle or 5-HT (1nM-5 μ M) alone (n= 7). Results obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m. Significance was determined by ANOVA, General Linear Model, Minitab. The two dose response curves were significantly different, $p < 0.001$ ***.

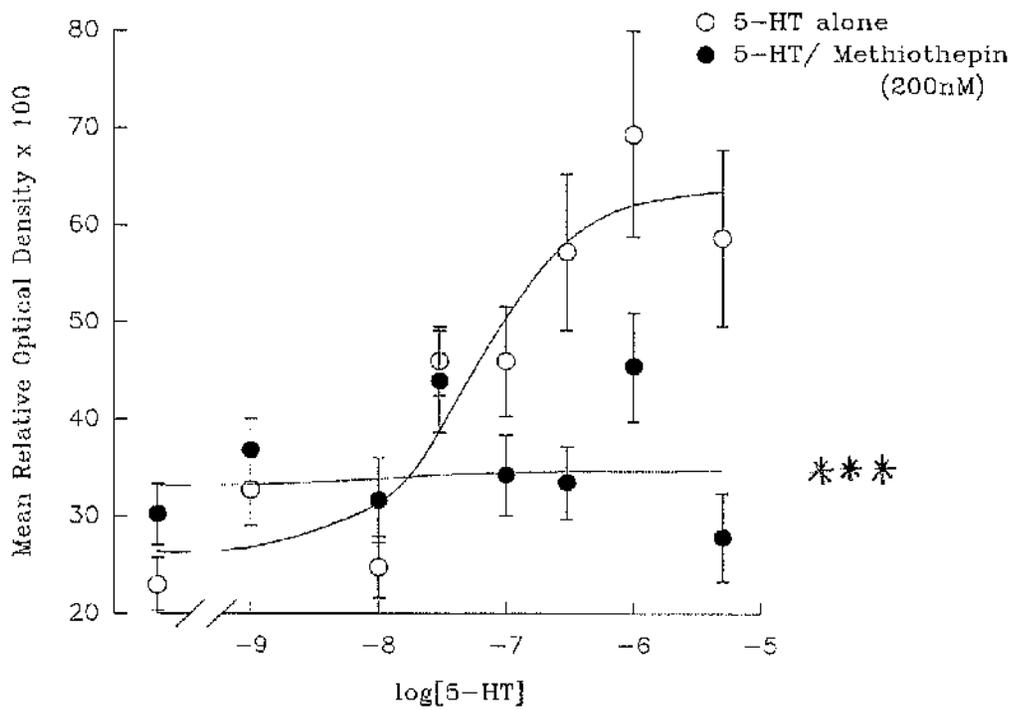
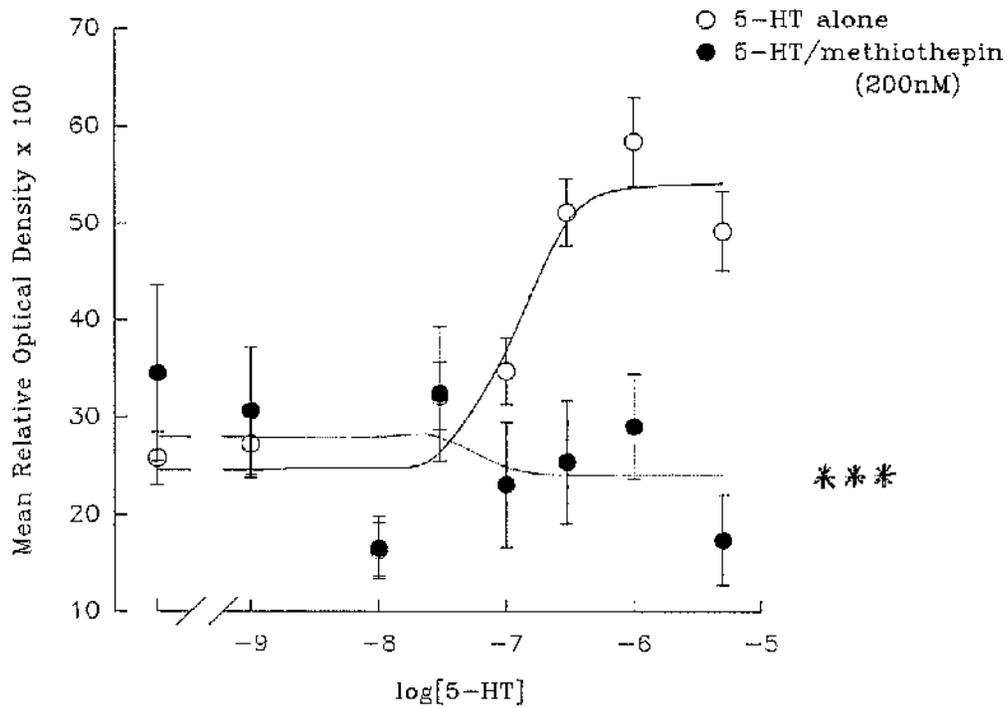


Figure 8.1.7 : Effect of inhibition of serotonergic 5-HT_{1A} /_{2A} /_{2C} receptors on the levels of expression of zif/268 mRNA in rat striatal neuron cultures. Cultures were pretreated for 30 minutes with mesulergine (100nM), prior to the addition of vehicle or 5-HT (1nM-5 μ M) (n= 3) or treated with vehicle or 5-HT (1nM-5 μ M) alone (n= 8). Results obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m. Significance was determined by ANOVA, General Linear Model, Minitab. The two dose response curves were significantly different, $p < 0.001$ ***.

Figure 8.1.8 : Effect of inhibition of serotonergic 5-HT_{1A} /_{2A} /_{2C} receptors on the levels of expression of zif/268 mRNA in rat striatal neuron cultures. Cultures were pretreated for 30 minutes with mesulergine (30nM), prior to the addition of vehicle or 5-HT (1nM-5 μ M) (n= 3) or treated with vehicle or 5-HT (1nM-5 μ M) alone (n= 8). Results obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m. Significance was determined by ANOVA, General Linear Model, Minitab. The two dose response curves were significantly different, $p < 0.001$ ***.

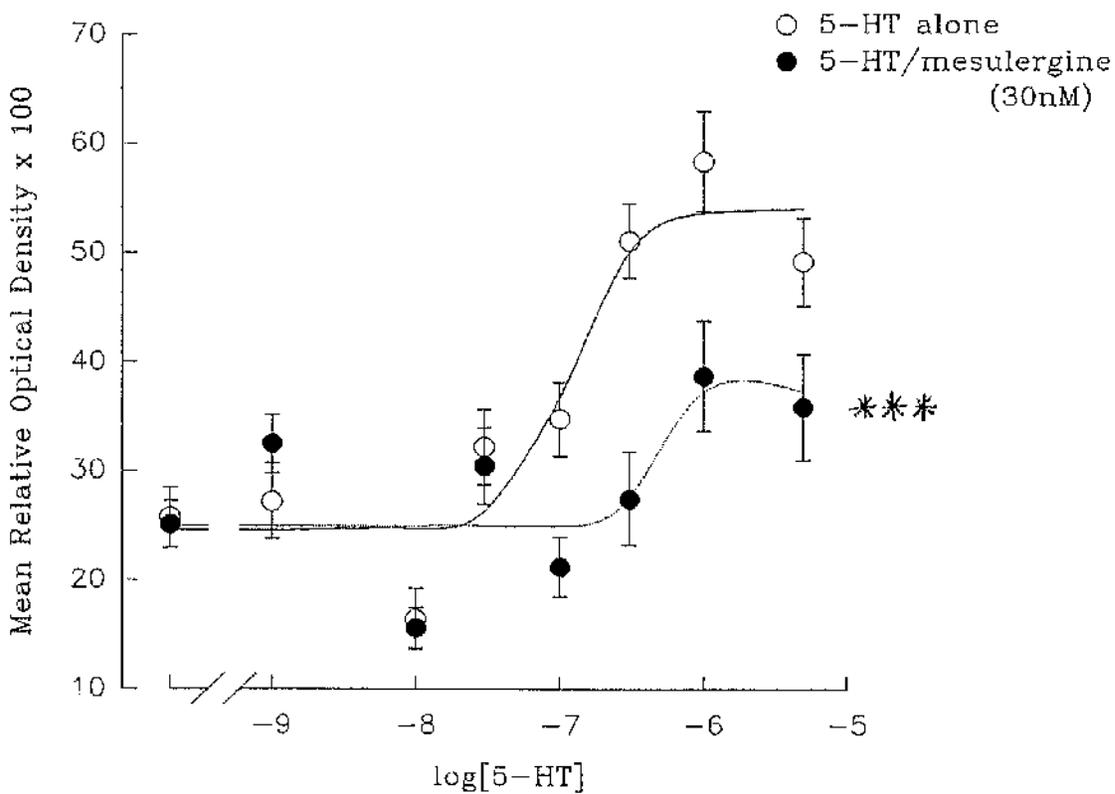
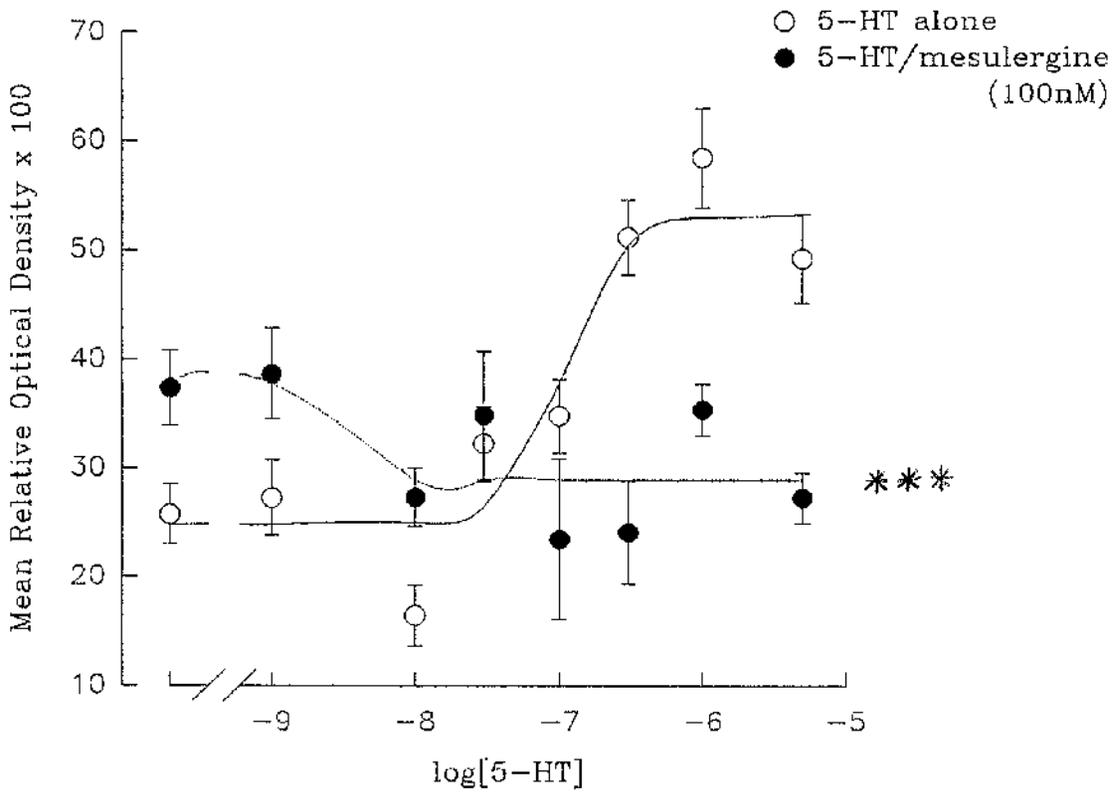


Figure 8.1.9 : Effect of inhibition of serotonergic 5-HT_{1A} /_{2A} /_{2C} receptors on the levels of expression of c-fos mRNA in rat striatal neuron cultures. Cultures were pretreated for 30 minutes with mesulergine (100nM), prior to the addition of vehicle or 5-HT (1nM-5 μ M) (n= 4) or treated with vehicle or 5-HT (1nM-5 μ M) alone (n= 7). Results obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m. Significance was determined by ANOVA, General Linear Model, Minitab. $p < 0.01$ ***.

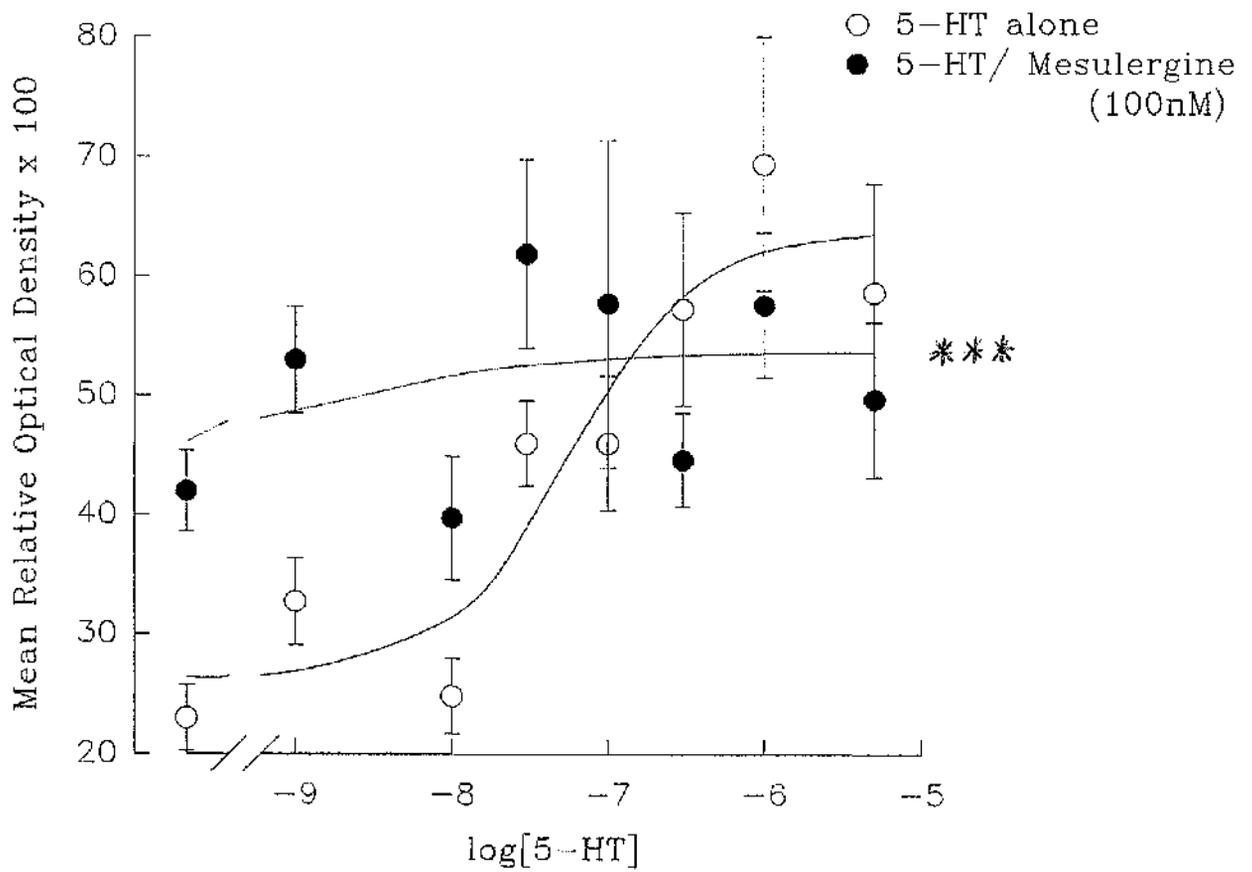


Figure 8.1.10: Effect of inhibition of serotonergic 5-HT_{1A} /2A /2C / 7 receptors on the levels of expression of zif/268 mRNA in rat striatal neuron cultures. Cultures were pretreated for 30 minutes with spiperone (100nM), prior to the addition of vehicle or 5-HT (1nM-5 μ M) (n= 3) or treated with vehicle or 5-HT (1nM-5 μ M) alone (n= 8). Results obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m. Significance was determined by ANOVA, General Linear Model, Minitab. The two dose response curves were significantly different, $p < 0.001$ ***.

Figure 8.1.11 : Effect of inhibition of serotonergic 5-HT_{1A} /2A /2C / 7 receptors on the levels of expression of zif/268 mRNA in rat striatal neuron cultures. Cultures were pretreated for 30 minutes with spiperone (30nM), prior to the addition of vehicle or 5-HT (1nM-5 μ M) (n= 3) or treated with vehicle or 5-HT (1nM-5 μ M) alone (n= 8). Results obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m. Significance was determined by ANOVA, General Linear Model, Minitab. The two dose response curves were significantly different, $p < 0.001$ ***.

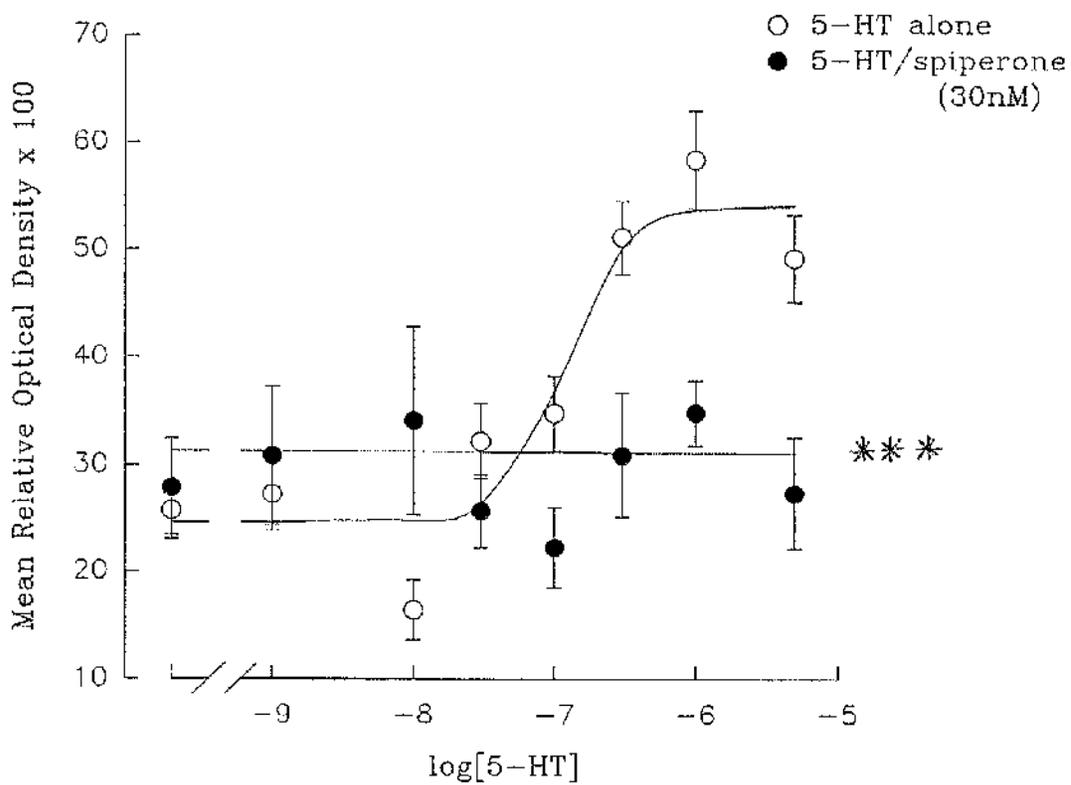
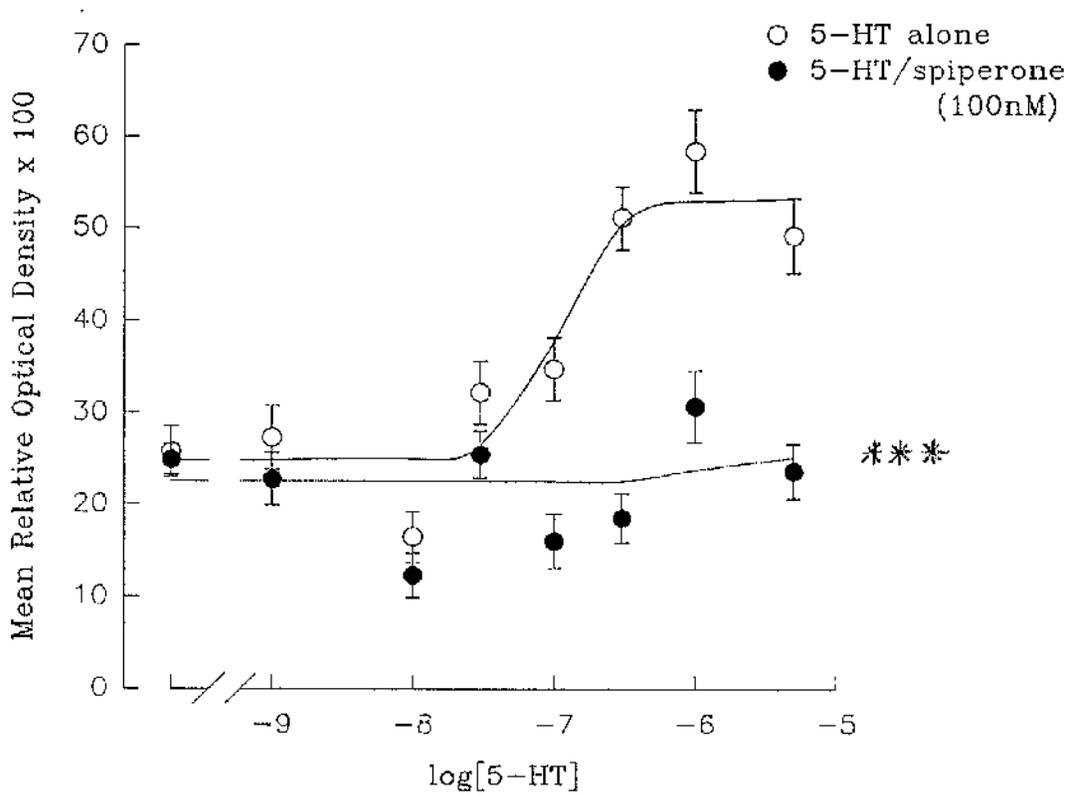


Figure 8.1.12 : Effect of inhibition of serotonergic 5-HT_{1A} /_{2A} /_{2C} / ₇ receptors on the levels of expression of c-fos mRNA in rat striatal neuron cultures. Cultures were pretreated for 30 minutes with spiperone (30nM), prior to the addition of vehicle or 5-HT (1nM-5 μ M) (n= 3) or treated with vehicle or 5-HT (1nM-5 μ M) alone (n= 7). Results obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m. Significance was determined by ANOVA, General Linear Model, Minitab. The two dose response curves were significantly different, $p < 0.001$ ***.

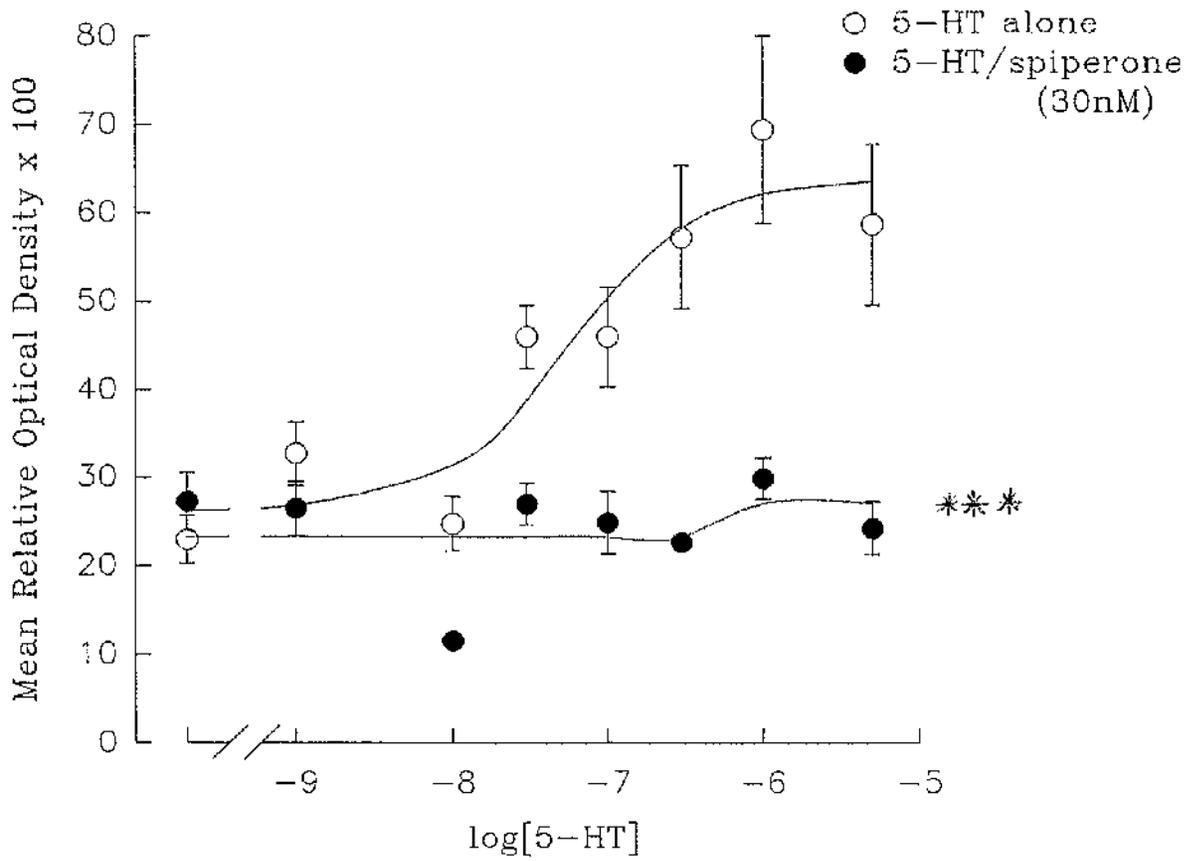


Figure 8.1.13 : Effect of a 5-HT_{2C} receptor stimulation on the level of expression of zif/268 mRNA in rat striatal neuron cultures. Cultures were treated for 45 minutes with vehicle (HBSS), mCPP (1nM-5 μ M, n= 5) alone or with 5-HT (1nM-5 μ M, n= 8) alone for comparison. Result, obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m.

Figure 8.1.14 : Effect of a 5-HT_{2C} receptor stimulation on the level of expression of c-fos mRNA in rat striatal neuron cultures. Cultures were treated for 45 minutes with vehicle (HBSS), mCPP (1nM-5 μ M, n= 3) alone or with 5-HT (1nM-5 μ M, n= 7) alone for comparison. Results, obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m.

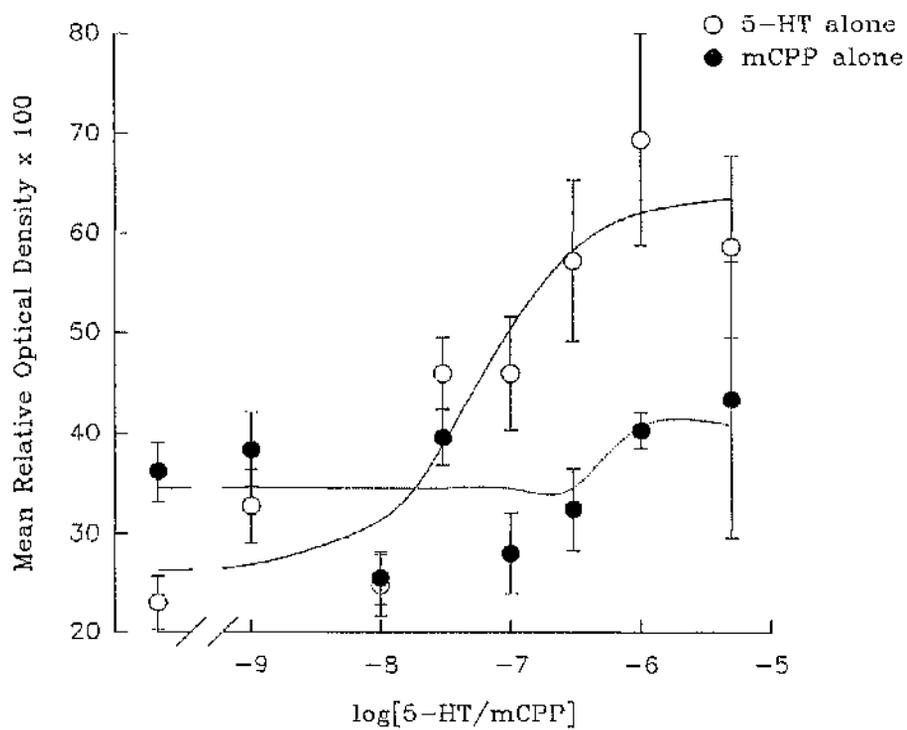
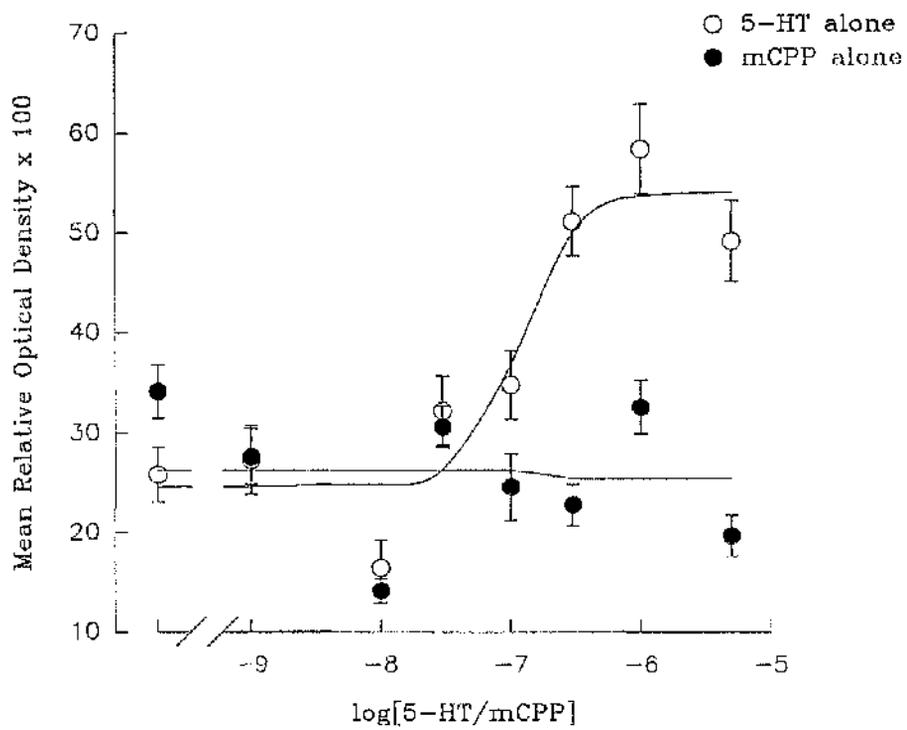


Figure 8.1.15 : Effect of a 5-HT_{1A} receptor stimulation on the level of expression of zif/268 mRNA in rat striatal neuron cultures. Cultures were treated for 45 minutes with vehicle (HBSS), 8OHDPAT (1nM-5 μ M, n= 4) alone or with 5-HT (1nM-5 μ M, n= 8) alone for comparison. Results, obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m. No significant induction of zif/268 mRNA was observed.

Figure 8.1.16 : Effect of a 5-HT_{1A} receptor stimulation on the level of expression of c-fos mRNA in rat striatal neuron cultures. Cultures were treated for 45 minutes with vehicle (HBSS), 8OHDPAT (1nM-5 μ M, n= 4) alone or with 5-HT (1nM-5 μ M, n= 7) alone for comparison. Results, obtained from film autoradiographs, are expressed as mean relative optical density $\times 10^2 \pm$ s.e.m. No significant induction of c-fos mRNA was observed.

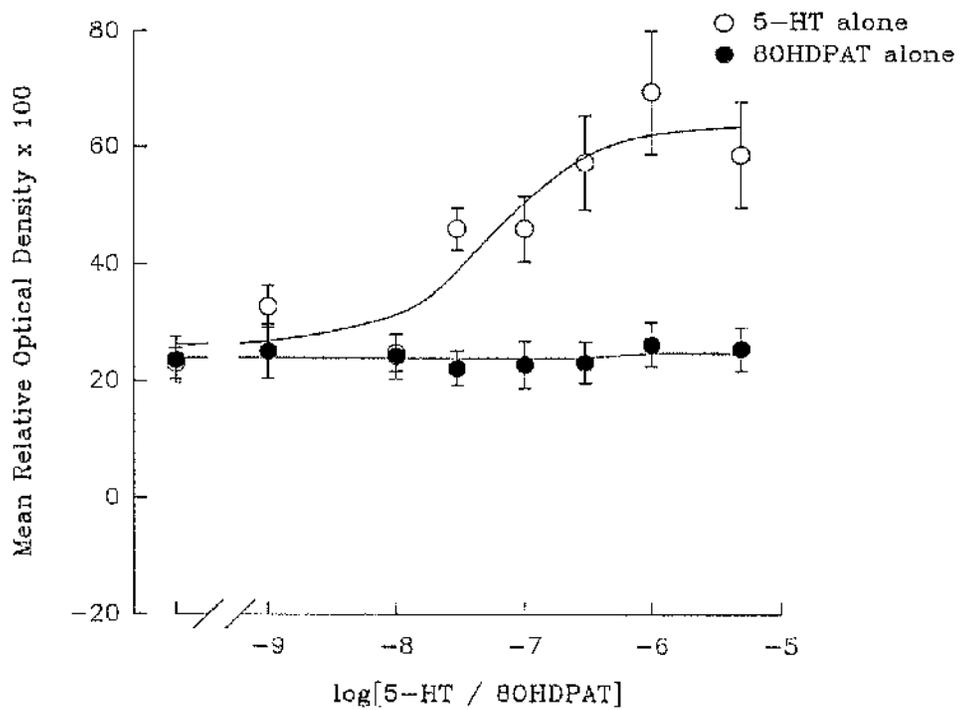
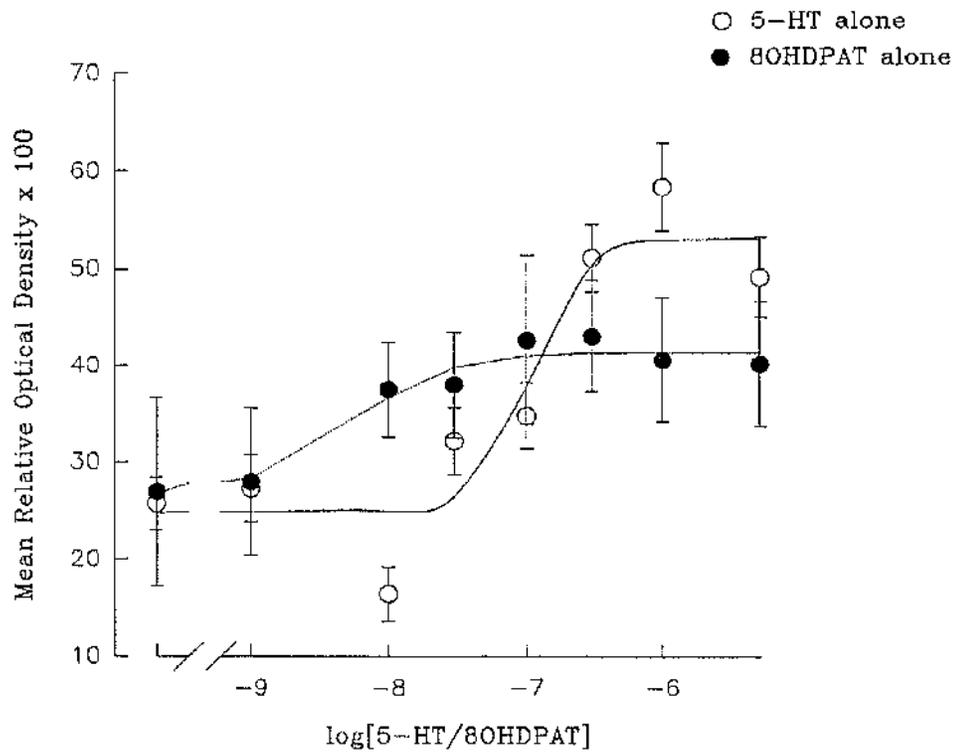


Figure 8.1.17 : Effect of a 5-HT_{2A/2C} receptor stimulation on the level of expression of zif/268 mRNA in rat striatal neuron cultures. Cultures were treated for 45 minutes with vehicle (HBSS), ± DOI (1nM-5μM, n= 3) alone or with 5-HT (1nM-5μM, n= 8) alone for comparison. Results, obtained from film autoradiographs, are expressed as mean relative optical density x 10² ± s.e.m. No significant induction of zif/268 mRNA was observed.

Figure 8.1.18 : Effect of a 5-HT_{2A/2C} receptor stimulation on the level of expression of c-fos mRNA in rat striatal neuron cultures. Cultures were treated for 45 minutes with vehicle (HBSS), ± DOI (1nM-5μM, n= 3) alone or with 5-HT (1nM-5μM, n= 7) alone for comparison. Results, obtained from film autoradiographs, are expressed as mean relative optical density x 10² ± s.e.m. No significant induction of c-fos mRNA was observed.

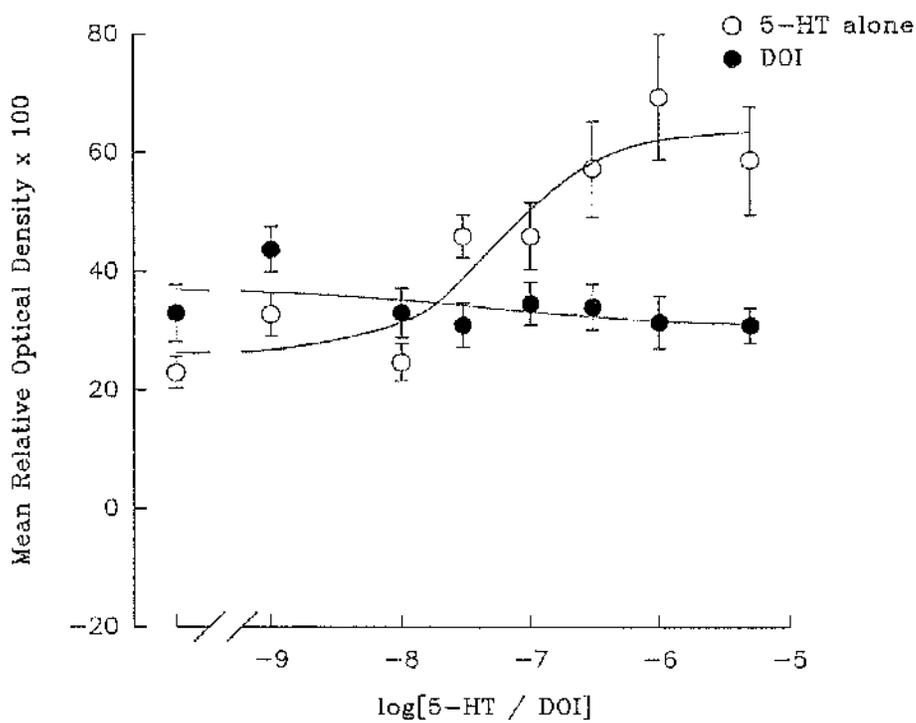
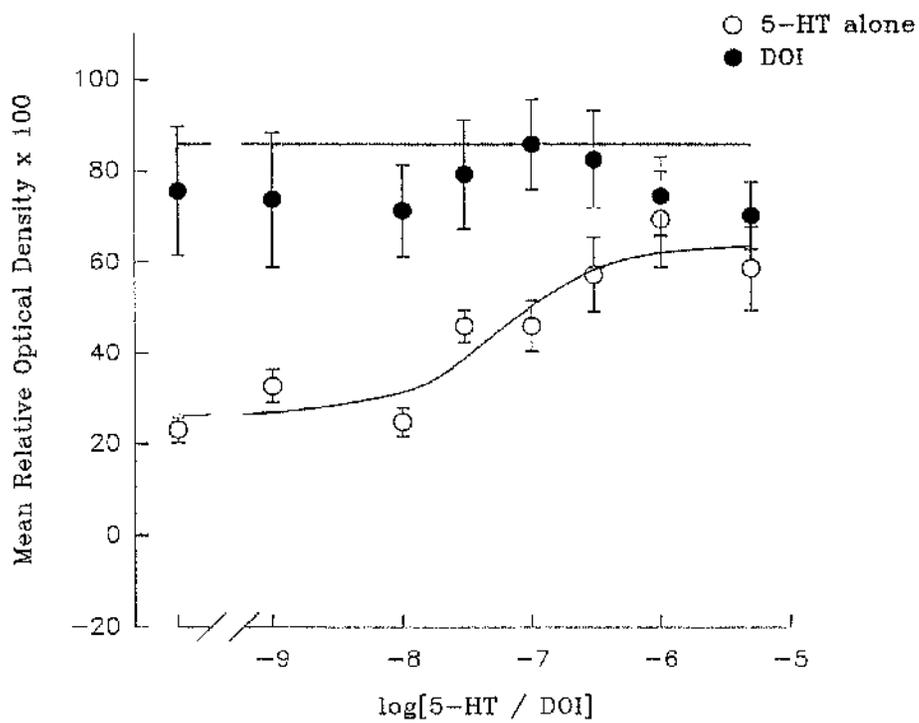
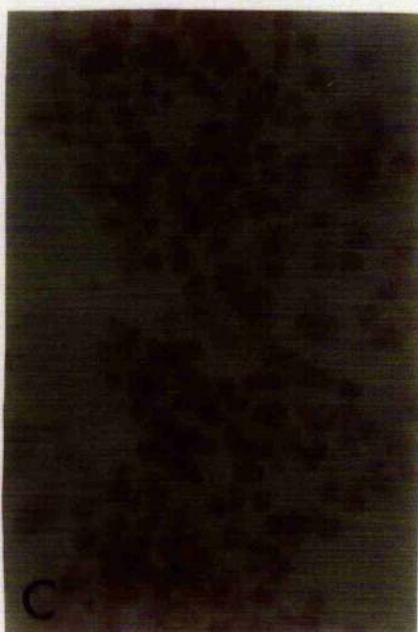
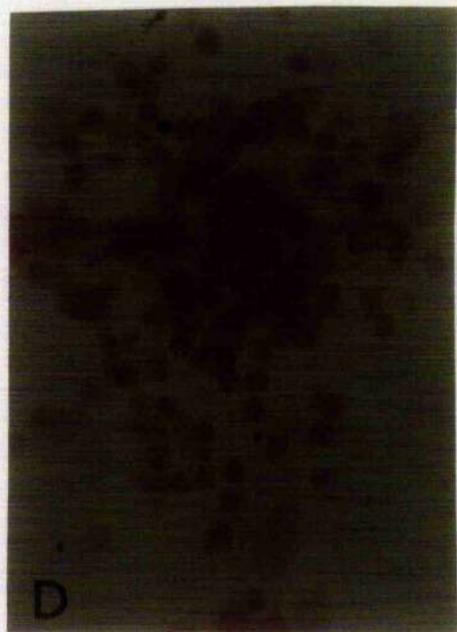


Figure 8.1.19 : Effect of serotonergic antagonists on the level of expression of zif/268 mRNA in rat striatal neuron cultures. Photomicrographs show neuronal cultures treated with (A) vehicle, (B) 5-HT ($1\mu\text{M}$), (C) cyproheptadine (500nM)/ 5-HT ($1\mu\text{M}$) and (D) spiperone (30nM)/ 5-HT ($1\mu\text{M}$). An induction of zif/268 can be seen as an increase in the number of silver grains over the cell bodies. Scale bar represents $25\mu\text{m}$.



DISCUSSION

8.2 Regulation of immediate early genes by serotonin in vitro

It has been known for some time that serotonin may play an important role in the manifestation of certain pathological disorders, including pain, anxiety, schizophrenia and parkinsonism. With recent evidence suggesting that serotonin may be involved in the modulation of immediate early gene expression both in vivo and in vitro, this study was undertaken in order to examine the pattern of immediate early gene expression in striatal neurons grown in culture, and to elucidate a possible receptor mechanism(s) involved.

The results show for the first time that serotonin can induce the expression of both *c-fos* and *zif/268* mRNA in rat primary striatal culture when administered alone. This inductive effect of serotonin on *c-fos* and *zif/268* gene expression proved to be dose-dependent. The EC_{50} value for both *c-fos* and *zif/268* was very similar, and in the range of 100nM. This value is similar to an EC_{50} value of 125 nM, recently reported for the 5-HT stimulation of cyclic AMP production, in striatal neurons in culture (Sebben *et al.*, 1994). Previous research indicates a balance between the dopaminergic and serotonergic systems, and that the central serotonergic system exerts an inhibitory influence on the central dopaminergic system. In particular this has been evident when looking at behavioural effects in rats. Serotonin agonists have been shown to potentiate the cataleptic effect of haloperidol in a dose dependent manner in rats, and pretreatment with serotonin antagonists was shown to reduce this effect (Balsara *et al.*, 1979). It appears that the effect of neuroleptics apparently depends on the balance between the two systems (Kostowski *et al.*, 1972; Balsara *et al.*, 1979). Recent research has suggested that serotonin may act synergistically in the striatum with the neurotransmitter, dopamine, in order to activate transcription factor genes, including *c-fos* and

zif/268 in vivo (Bhat and Baraban, 1993). Our results demonstrate that serotonin may also have a solitary role in activating these transcription factor genes.

A variety of pharmacological agents known to be active at various subtypes of 5-HT receptors were used to characterize the subtype of receptor responsible for this stimulation of *c-fos* and *zif/268* mRNA by serotonin. It was thought to be unlikely that this response was mediated by a 5-HT₃ subtype of receptor, because pretreatment with the selective 5-HT₃ receptor antagonist, ondansetron, had no effect on the expression of *zif/268* mRNA and little effect on *c-fos* mRNA levels after 5-HT stimulation. Receptors of the 5-HT₁ subclass generally couple to inhibition of adenylyl cyclase, although there are reports of the 5-HT_{1A} receptor coupling to stimulation of this second messenger (Shenker *et al.*, 1987). The inactivity of 8-hydroxy-DPAT, a potent 5-HT_{1A} agonist (Gozlan *et al.*, 1983; Dourish *et al.*, 1987), indicates that the 5-HT_{1A} receptor is not responsible for the stimulation of *zif/268* or *c-fos* mRNA. The involvement of a 5-HT_{1B} or 5-HT_{1D} receptor was eliminated by the finding that the drugs spiperone, mesulergine and cyproheptadine, all having very low affinity for these receptor subtypes (Peroutka, 1986; Van Wijngaarden *et al.*, 1990), were clearly effective antagonists of the inductive response of *c-fos* and *zif/268* mRNA, to serotonin stimulation. The lack of agonist activity for mCPP at high doses, also suggests that the response is not mediated by the 5-HT_{1B} receptor subtype (Hoyer and Schoeffter, 1991). A 5-HT_{1E}-like receptor is known to be highly expressed in the rat striatum, although the primary structure of this receptor suggests it may be coupled to an inhibition of adenylyl cyclase through an inhibitory G_i-like protein (Lovenberg *et al.*, 1992). It is unlikely that the induction of *zif/268* and *c-fos* is due to stimulation of a receptor negatively coupled to adenylyl cyclase. This, together with the reported low affinity of mesulergine at this receptor subtype makes it an unlikely candidate. The 5-HT₄ subtype was also eliminated as a possible

candidate, because the drugs spiperone and mesulergine have been reported to be inactive at this receptor (Schiavi *et al.*, 1993). The 5-HT_{5A} and 5-HT_{5B} receptors can also be eliminated, because of their moderate affinity for methiothepin and the knowledge that these receptors are unlikely to be coupled to either adenylate cyclase or phospholipase C (Matthes *et al.*, 1992). Similarly stimulation of the newly cloned 5-HT₆ receptor is unlikely to be responsible for the inductive effect on *c-fos* and *zif/268* gene expression, because spiperone is known to have a very low affinity at this receptor site (Roth *et al.*, 1993), which does not correspond to its action as a highly potent antagonist seen in our experiments. However, the affinity of spiperone at the 5-HT₇ receptor is much higher than at the 5-HT₆ receptor subtype. The 5-HT₇ or 5-HT_x receptor as it is also known, is expressed in the forebrain region in rat brain (Plassat *et al.*, 1993) and has a high affinity for atypical antipsychotic drugs such as clozapine (Roth *et al.*, 1993). This receptor subtype can not be ruled out totally, in view of the current evidence.

The presence of a 5-HT-stimulated increase in *zif/268* and *c-fos* mRNA initially suggested the involvement of a 5-HT₂-like receptor. It is known that this receptor subtype is coupled to phospholipase C and the consequent cascade leading to diacylglycerol and inositol triphosphate production (Conn and Sanders-Bush, 1986; Conn *et al.*, 1986). In previous results we have suggested that both *c-fos* and *zif/268* stimulation may be linked to protein kinase C, an enzyme responsible for the phosphorylation of cellular proteins. This further suggests that the response to 5-HT may involve a 5-HT₂-like receptor subtype. However, the 5-HT_{2C} (formerly known as 5-HT_{1C}) subtype is unlikely to be responsible for mediating the induction of *c-fos* and *zif/268* mRNA *in vitro*, due to the low affinity of spiperone at this subtype of 5-HT₂ receptor (Hoyer *et al.*, 1985). The lack of an agonist effect of mCPP and \pm -DOI also suggests that the 5-HT_{2C} receptor is not involved in mediating this effect. Similarly the 5-HT_{2B} receptor (formerly known as 5-HT_{2F}) is most unlikely to

be mediating the induction of *c-fos* and *zif/268* mRNA in culture. This is due to the low affinity of spiperone for this receptor subtype (Wainscott *et al.*, 1992), together with the low abundance of this receptor subtype in the brain (Pompeiano *et al.*, 1993). The 5-HT_{2A} subtype is possibly the most feasible candidate, however there are some discrepancies which do not coincide with the pharmacologically described 5-HT_{2A} receptor. The 5-HT_{2A} has high affinity for spiperone, mesulergine and cyproheptadine (Zifa and Fillion, 1992), which is in agreement with our results describing potent antagonism of the 5-HT-stimulated increase in *c-fos* and *zif/268* mRNA with these three drugs. The 5-HT_{2A} subtype has also been shown to be expressed at moderate levels in the caudate-putamen in the adult rat (Pompeiano *et al.*, 1993). In humans, 5-HT₂ binding sites have been predominantly located in striosomes in the basal ganglia (Waeber and Palacios, 1994). However, we found that the selective agonist at 5-HT_{2A} and 5-HT_{2C} receptor subtypes, \pm -DOI, was unable to stimulate the induction of *zif/268* or *c-fos* mRNA in striatal culture. In previous reports \pm -DOI has been suggested to be a mixed agonist-antagonist (Dabire *et al.*, 1989) and also be able to inhibit 5-HT-induced neuronal activity in the dorsal raphe, an effect not blocked by classical 5-HT₂ receptor antagonists (Garratt *et al.*, 1991). This suggests that there are properties of \pm -DOI aside from the known agonist activity at 5-HT₂ receptors, that as yet, have not been fully investigated.

These results indicate that the serotonin receptor responsible for the stimulation of both *c-fos* and *zif/268* mRNA in rat striatal culture may be a 5-HT_{2A}-like receptor. However, there are still open avenues of investigation concerning the 5-HT₇ receptor, which has recently been reported to have a high affinity for antipsychotic drugs such as clozapine. As yet, we have been unable to obtain a result with clozapine in this experimental model.

Ultimately the physiological significance of our results is suggestive that serotonin can modulate cellular gene expression which in turn may affect

behavioural responses. Accumulated experimental evidence implies that the immediate early genes are 'third messengers' in signal transduction pathways (Goelet *et al.*, 1986; Curran and Morgan, 1987). In the central nervous system, certain late response genes have been identified as targets for transactivation based on the presence of the appropriate regulatory sequences in their promoters (Comb *et al.*, 1988; Sonnenberg *et al.*, 1989; Naranjo *et al.*, 1991).

It is well characterised that neuroleptics, the major treatment for diseases such as schizophrenia, can cause the induction of a number of immediate early genes in the brain, in addition to affecting the levels of many late response genes. Therefore a functional link has been implied between the induction of immediate early genes and the consequent increase or decrease in late response genes such as enkephalin (Sonnenberg *et al.*, 1989). In view of the affinity of many neuroleptic drugs for 5-HT receptors, the current awareness is suggesting that neuroleptics may be affecting the serotonergic system as easily as the dopaminergic system, and in fact the atypical drug clozapine has a particularly high affinity for some serotonin receptor subtypes. The properties of clozapine and similar atypical compounds result in fewer extrapyramidal side effects when given clinically, and it has been postulated that clozapine's actions on the serotonergic system may be an important contributing factor to its atypical drug profile.

General Discussion

As a whole, this work has not only uncovered some novel findings, but demonstrated the complexity of the functioning of the basal ganglia. In addition to demonstrating an induction of *c-fos* and *zif/268* gene expression in the rat striatal model *in vitro* following a dopaminergic D1/D5 receptor stimulation, the expression of these immediate early genes and others, was also increased by dopaminergic D2-like antagonists *in vivo*. Dopamine D1-like-induced

expression of *c-fos* in the caudate *in vivo* has been shown to occur only after a 6-hydroxy-dopamine-induced lesion (Robertson *et al.*, 1989; 1990), although the D1/D5 receptor antagonist SCH23390 has been demonstrated to reduce the mRNA levels of the transcription factor *zif/268* in the adult rat intact striatum (Mailleux *et al.*, 1992). A possible explanation for the D2-like mediated mRNA induction of these immediate early genes, may be that blockade of D2-like receptors results in an unopposed D1 stimulation.

We have also demonstrated that a further complexity may exist concerning the second messenger pathway prior to immediate early gene induction after a D1-like receptor stimulation. The results shown provide the first evidence that both PKA and PKC participate in the regulation of *c-fos* and *zif/268* gene expression by dopamine in striatal neurones. Further work is clearly required to discover the functional significance of these control mechanisms.

The serotonergic regulation of *c-fos* and *zif/268* gene expression in the striatal model *in vitro* is the first evidence to suggest that serotonin alone can induce these immediate early genes. In addition, we have attempted to reveal a possible receptor mechanism for this induction. Our results are suggestive that the induction may be mediated by a 5-HT_{2A}-like receptor, although conclusive evidence is hampered by the few selective drugs available for serotonin receptor subtypes. Perhaps an improved receptor antisense technology may provide the answer in the future.

We have also shown that the recently cloned transcription factor gene BF-1, present in the striatum, can be regulated by the atypical neuroleptic, clozapine, *in vivo*. This is a novel finding, which may be functionally important considering clozapine has a unique drug profile that provides clinical advantages in schizophrenia.

An important consideration when attempting to explain many of the findings in this research work, is the possibility of interaction occurring between different neuronal systems. The therapeutic efficacy of atypical antipsychotic agents such as clozapine may be attributed to the ability of the drug to act on different transmitter/receptor systems in vivo. From this, it may be suggested that drug efficacy may rely on a complex compensatory mechanism between neuronal systems. Indeed, one explanation for the selective behavioural and physiological effects of atypical agents on the function of the A10 dopamine system is an interaction with other neuronal systems which may compensate for dopamine receptor antagonism in the nigrostriatal system. Since clozapine is known to have a high affinity for a number of 5-HT receptor subtypes in vitro, of which several are known to be expressed in the caudate-putamen, tonic 5-HT activity may be crucial for the resultant physiological effects of this drug. Wilmot and Szczepanik (1989), have previously demonstrated that both acute and chronic intraperitoneal administration of clozapine, at relatively high doses, can decrease the maximal number of cortical 5-HT₂ receptors by greater than 50% of control, whereas typical drugs such as haloperidol, had no effect. The functional significance of 5-HT₂ receptor down-regulation following chronic treatment with an atypical antipsychotic remains to be determined.

The existence of a relationship between the differential effects of clozapine and haloperidol on subcortical D₂ dopamine receptors and cortical or subcortical 5-HT₂ serotonin receptors is not presently known. However, the cortical control of striatal function is regionally specific. The lateral striatum, which shows the greatest changes in D₂ receptors following chronic treatment with haloperidol, receives afferents from the somatosensory frontoparietal cortex. It has been suggested that clozapine may indirectly affect the development of striatal D₂ receptor up-regulation via its effects on cortical somatosensory neurons. Another explanation is that clozapine, as a 5-HT₂ antagonist, directly affects raphe neurons which then influence dopamine neuronal function (Wilmot and Szczepanik, 1989). It has previously been

shown that acute treatment with 5-HT₂ antagonists can antagonize the haloperidol-induced increase in levels of striatal dihydroxyphenylacetic acid (DOPAC) and haloperidol-induced catalepsy (Balsara *et al.*, 1979). Lesions of the dorsal raphe are also known to prevent haloperidol-induced catalepsy (Kostowski *et al.*, 1972). Therefore it would seem that there is a definite interaction between the dopamine and serotonin transmitter systems in the central nervous system.

Regulation of tachykinin biosynthesis by serotonin has recently been shown in the neostriatum (Walker *et al.*, 1991). Depletion of 5-HT with p-chlorophenylalanine treatment decreased preprotachykinin (the prohormone precursor to substance P) mRNA levels in the neostriatum. By contrast, raising extracellular levels of 5-HT with zimelidine, a 5-HT uptake inhibitor, or clorgyline, a monoamine oxidase inhibitor resulted in increased levels of preprotachykinin mRNA. Therefore it may be suggested that certain peptides in the striatum may be regulated by serotonin neurotransmission. Typical neuroleptics such as haloperidol given acutely or chronically, have been shown to decrease the levels of preprotachykinin (PPT) mRNA in the striatum (Bannon *et al.*, 1986). The proposed mechanism for this biochemical change is a decrease in dopamine neurotransmission. In addition to the effect on the dopamine system, haloperidol may also be affecting the serotonin system. Haloperidol has been shown to have an affinity at a number of serotonin receptor subtypes, including the 5-HT₂ receptor (Canton *et al.*, 1993), therefore the decrease in PPT mRNA seen after haloperidol may be due to a decrease in serotonin neurotransmission in addition to a decrease in dopamine neurotransmission. Tonic serotonin levels may therefore have a regulatory role in controlling the expression of neuropeptides in the basal ganglia. The increase in the immediate early genes *c-fos* and *zif/268* which we observed after treatment with 5-HT could also be blocked by 5-HT₂-like receptor antagonist compounds. As yet there is no clear evidence for a functional role of immediate early genes in the brain, although it has been suggested that they may play a part in the regulation of gene expression. Similarly glutamate must also be considered to be a possible interacting component in the

basal ganglia. A glutamatergic projection from the cortex to the striatum suggests that tonic glutamate may also have a homeostatic function in controlling the level of neurotransmitters /peptides in the striatum.

This research has also demonstrated an induction of proenkephalin mRNA after acute typical neuroleptic drug treatment. This is in agreement with previous studies that have shown an induction of proenkephalin mRNA and enkephalin protein in the rat striatum (Hong *et al.*, 1978; Sabol *et al.*, 1983; Tang *et al.*, 1983; Angulo *et al.*, 1986; Sivam *et al.*, 1986; Romano *et al.*, 1987; Morris *et al.*, 1988). We suggest that the increase of proenkephalin mRNA after 24 hours may in fact be related to the induction of a number of immediate early genes prior to this time point. This would be in agreement with the idea that immediate early genes act as "third messengers" in the sequence of events leading to changes in gene expression. However, this is only indirect evidence, and therefore a relationship between immediate early genes and "late response genes", such as enkephalin, is only circumstantial.

Tonic dopamine is thought to have an inhibitory control on enkephalin gene expression. Enkephalin peptide is thought to be located predominantly in striatopallidal neurons (Gerfen and Young, 1988), and striatopallidal neurons have been shown to be a distinct population of neurons expressing a relatively high proportion of D2-like receptors (Le Moine *et al.*, 1990). Blockade of D2-like dopamine receptors with typical neuroleptic drugs such as haloperidol cause an increase in proenkephalin mRNA, as the tonic inhibition is removed. Lesioning the nigrostriatal system, which effectively removes the tonic dopamine acting predominantly through D2-like receptors to maintain an inhibitory control on enkephalin gene expression, also causes an induction of enkephalin mRNA. Therefore it would seem that the D2-mediated inhibitory effect of tonic dopamine on the level of enkephalin gene expression predominates in the 'normal' brain.

Despite the evidence previously described suggesting that D1-like and D2-like dopamine receptors are expressed on different populations of cells *in vivo*

(Gerfen et al., 1990; Le Moine et al., 1990), our results indicate that both D1-like and D2-like receptors are expressed on the same cell in vitro. Administration of the D2 agonist quinpirole has been shown to antagonize the SKF38393-induced expression of both *c-fos* and *zif/268* mRNA. If it is assumed that our model of striatal neurons in culture closely resembles the phenotype of striatal neurons in vivo, then this suggests that a population of neurons in vivo may express both D1-like and D2-like dopamine receptors. There is already a large body of evidence in agreement with this idea (Seeman et al., 1989; Bertorello et al., 1990; Piomelli et al., 1991; Surmeier et al., 1993). Dopaminergic D1-like receptors, when stimulated, appear to be in opposition to dopaminergic D2-like receptors in the striatum. However, it must be remembered that the population size which possess both D1-like and D2-like dopamine receptor subtypes is unknown. It is possible that a relatively small population of neurons possess both phenotypes. Perhaps the proportion of D1:D2 phenotype on neuronal populations within the basal ganglia differ such that one phenotype predominates over the other in vivo. The tonic activity of dopamine on different neuronal populations within the caudate may therefore have localized effects in terms of gene regulation, dependent on the predominant phenotype of dopamine receptor. Several serotonin receptor subtypes are known to be present in the striatum. The physiological significance of these receptors has yet to be determined. Similar to the dopaminergic system, it is not unlikely that these receptor subtypes may be localised to particular neuronal cell populations and in turn regulate gene expression.

The relatively new concept of 'antisense strategy' to block the synthesis of specific proteins, has been employed in this research. The phosphorothioate antisense oligonucleotide directed towards the D1 dopamine receptor clearly reduced *c-fos* mRNA expression in striatal neurons after treatment with a range of doses of the D1-like receptor agonist SKF38393. The effect on *c-fos* mRNA expression after treatment with the D5 antisense oligonucleotide was less robust than that observed after treatment with the D1 antisense, although a significant

decline in mRNA levels were still observed. We have suggested that the physiological effect of c-fos induction following SKF38393, is predominantly mediated through stimulation of the D1 dopamine receptor subtype. However, as the D5 antisense was also effective, it may not be dismissed that the induction of c-fos mRNA in striatal neurons may also be mediated, but to a lesser extent, by the D5 dopamine receptor subtype. We have proposed that the D5 subtype may represent the 'fine control' over the dopaminergic system, due to only a small percentage of D5 receptors being expressed in the striatum, and dopamine, the endogenous ligand *in vivo*, having a higher affinity for the D5 dopamine receptor subtype than for D1 subtype.

At present there are a number of disadvantages with this method of protein 'knockdown'. Clearly, from the data on zif/268 mRNA expression after antisense use, non-specific effects can be a major problem. A number of separate controls, including the use of random sequence oligonucleotides, 'mismatch' oligonucleotides, radioligand binding studies and toxicity testing, should be performed in addition to the 'sense' control oligonucleotide that was used in our study in an attempt to clarify these results.

This research as a whole, has provided an important insight into the changes in gene expression in the basal ganglia. We have investigated the transient changes in immediate early gene expression, together with the more sustained changes in peptide gene expression in the striatum, in response to neuroleptic drug treatments, the clinical therapy for many neurodegenerative diseases. We have uncovered a complex cascade of events prior to the change in expression of the immediate early genes c-fos and zif/268 in the striatum after a dopaminergic stimulus. In addition we have also investigated another major transmitter system in the striatum, namely the serotonergic system, and attempted to reveal the receptor(s) involved in the induction of the immediate early genes c-fos and zif/268 after serotonin treatment.

The complexity of the basal ganglia has undoubtedly left several questions unanswered until a future date.

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