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**AN INVESTIGATION INTO THE RELATIONSHIP BETWEEN THE SYSTEMIC
INFLAMMATORY RESPONSE AND SURVIVAL: A GLASGOW INFLAMMATION
OUTCOME STUDY**

by

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MBChB, MRCS (Glas)

**SUBMITTED IN THE FULFILMENT OF THE REQUIREMENTS FOR THE
DEGREE OF MEDICAL DOCTORATE (MD)**

To

THE UNIVERSITY OF GLASGOW

**FROM RESEARCH CONDUCTED IN THE UNIVERSITY DEPARTMENT OF
SURGERY, GLASGOW ROYAL INFIRMARY, FACULTY OF MEDICINE,
UNIVERSITY OF GLASGOW**

ABSTRACT

Inflammation has been shown to play an integral role in a number of different disease processes. There is evidence that measurements of the systemic immune/inflammatory response are associated with mortality in patients with certain malignancies, as well as those with atherosclerotic disease. A variety of readily available inflammation-based prognostic scores, combining serum constituents of the systemic immune/inflammatory response, have been suggested for use in these patient groups. However, it is unclear whether systemic inflammation-based scores are universally associated with mortality across all tumour types and whether this relationship persists in atherosclerotic conditions including cardiovascular and cerebrovascular disease. The optimal constituents of such a score are also still to be determined. The present thesis further examines these topics with specific reference to:

1. The relationship between the presence of cancer, an inflammation-based prognostic score and biochemical parameters in a large group of unselected patients.
2. The relationship between an inflammation-based prognostic score and cancer survival in patients with malignancies at a number of different sites.
3. A comparison of commonly available inflammation-based prognostic scores in patients with cancer.
4. The prognostic value of a derived neutrophil to lymphocyte ratio, commonly available in patients with cancer undergoing chemotherapy.
5. The optimal serum constituents of the systemic inflammation-based Glasgow Prognostic Score (optimised) in patients with cancer.
6. The optimal constituents and prognostic value of a systemic inflammation-based score for predicting cardiovascular, cerebrovascular and all-cause mortality.

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Glasgow Royal Infirmary

DECLARATION

The work presented in this thesis was undertaken during a period of research between 2009 and 2011 in the University Departments of Surgery and Public Health at Glasgow Royal Infirmary. The introduction (Chapter 1) and literature search (Chapter 2) were written prior to the start of data collection and these sections are therefore based on current knowledge at the time (December 2009). Individual chapters contain more recent references as the study progressed and the knowledge base grew. The work has been completed whilst working as a Speciality Registrar in General Surgery in the South East of Scotland deanery. The study was approved by the Research Ethics Committee, North Glasgow NHS Trust. I declare the work presented in this thesis was solely undertaken by me with the exception of the following:

- Writing of the computer code used to extract biochemical variables (Chapters 2-7) and pathological details (Chapter 2) for patients from the North of Glasgow biochemical and pathological database was performed with assistance from Mr Stephen Balmer, laboratory information management and technology, NHS Greater Glasgow and Clyde.
- Writing of the computer code used to extract haematological variables (Chapters 4-7) for patients from the North of Glasgow haematological database was performed with assistance from Mr Colin Fletcher, consultant clinical scientist, Greater Glasgow and Clyde.
- Extraction of cancer follow-up data from the Scottish Cancer Registry (Chapter 3-7) was performed with assistance from Mr Billy Sloan.

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PUBLICATIONS

1. The relationship between the presence and site of cancer, an inflammation-based prognostic score and biochemical parameters. Initial results of the Glasgow Inflammation Outcome Study.
Proctor MJ, Morrison DS, Talwar D, Balmer SM, Fletcher CD, O'Reilly DS, Foulis AK, Horgan PG, McMillan DC.
BJC 2010; **103**: 870-876.
2. An inflammation-based prognostic score (mGPS) predicts cancer survival independent of tumour site: a Glasgow Inflammation Outcome Study.
Proctor MJ, Morrison DS, Talwar D, Balmer SM, Fletcher CD, O'Reilly DS, Foulis AK, Horgan PG, McMillan DC.
BJC 2011; **104**: 160-161.
3. A comparison of inflammation-based prognostic scores in patients with cancer. A Glasgow Inflammation Outcome Study.
Proctor MJ, Morrison DS, Talwar D, Balmer SM, Fletcher CD, O'Reilly DS, Foulis AK, Horgan PG, McMillan DC.
Eur J Cancer 2011; **47**: 2633-41.
4. A derived Neutrophil to Lymphocyte Ratio (dNLR) predicts survival in patients with cancer.
Proctor MJ, McMillan DC, Morrison DS, Fletcher CD, Horgan PG, Clarke SJ.
BJC 2012. **107**: 695-9.
5. Optimisation of the systemic inflammation-based Glasgow prognostic score: a Glasgow Inflammation Outcome Study.
Proctor MJ, Horgan PG, Talwar D, Fletcher CD, Morrison DS, McMillan DC.
Cancer 2013. **119**. 2325-32.

6. Systemic inflammation predicts all-cause mortality. A Glasgow Inflammation Outcome Study.

Proctor MJ, McMillan DC, Horgan PG, Fletcher CD, Talwar D, Morrison DS.

Submitted to Plos One.

PRESENTATIONS

1. The relationship between the systemic inflammatory response and tumour type in a large hospital based cohort.
The Association of Surgeons of Great Britain and Ireland, Glasgow, 2009 (Poster)
2. The relationship between deprivation and the systemic inflammatory response in cancer: a large hospital based cohort study.
The Association of Surgeons of Great Britain and Ireland, Glasgow, 2009 (Oral)
3. Inflammation-based prognostic score are predictive of outcome in patients with cancer.
The Association of Surgeons of Great Britain and Ireland, Bournemouth 2011 (Oral)

DEDICATION

To my father John, my mother Helen and my sister Lucy for their love and support through thick and thin.

1.0. INTRODUCTION

1.1. THE IMMUNE/INFLAMMATORY RESPONSE

1.1.1. The Local Immune/Inflammatory Response

The immune/inflammatory response, as a reaction to injury, infection, inflammatory conditions or malignancy, occurs at both a local and systemic level. The local immune/inflammatory reaction is characterised by the four clinical signs of inflammation; erythema, heat, swelling and pain. Erythema and increased tissue temperature occur as a result of mast cell degradation and cytokine mediated vasodilatation. Similarly, swelling occurs as a result of cytokine mediated increased capillary permeability and subsequent accumulation of extravascular exudate and inflammatory cells. During the local immune/inflammatory response, pain evoking proinflammatory mediators are also released [Chandrasoma and Taylor, 2005]. The local immune/inflammatory response, through complex immunological processes detailed below, can in turn initiate a global, systemic immune/inflammatory response.

1.1.2. The Innate Immune System

Immunity can be broadly divided into the innate, humoral and adaptive systems [Abbas et al., 2005]. It is however worth noting that these definitions are arbitrary as the processes involved are interrelated and the distinctions between each system blurred at both the local and systemic levels (Figure 1.1). The innate immune system is a non-specific, first line defence against insult or attack and aims to promote clearance of pathogens, wound healing and tissue repair. Cells of the innate immune system have germline encoded recognition with the ability to detect non-self and are therefore integral in local and systemic

inflammatory processes [Janeway and Medzhitov, 2002]. At a local level, activation of the complement system stimulates mast cell release of histamine and protease, resulting in clinical signs of local inflammation [Reid and Porter, 1981; Galli et al., 1993]. As illustrated in Figure 1.1, phagocytic cells initiate inflammatory pathways by releasing pro-inflammatory cytokines and activating the complement system, termed the humoral response [Laskin and Pendino, 1995; Koj, 1996].

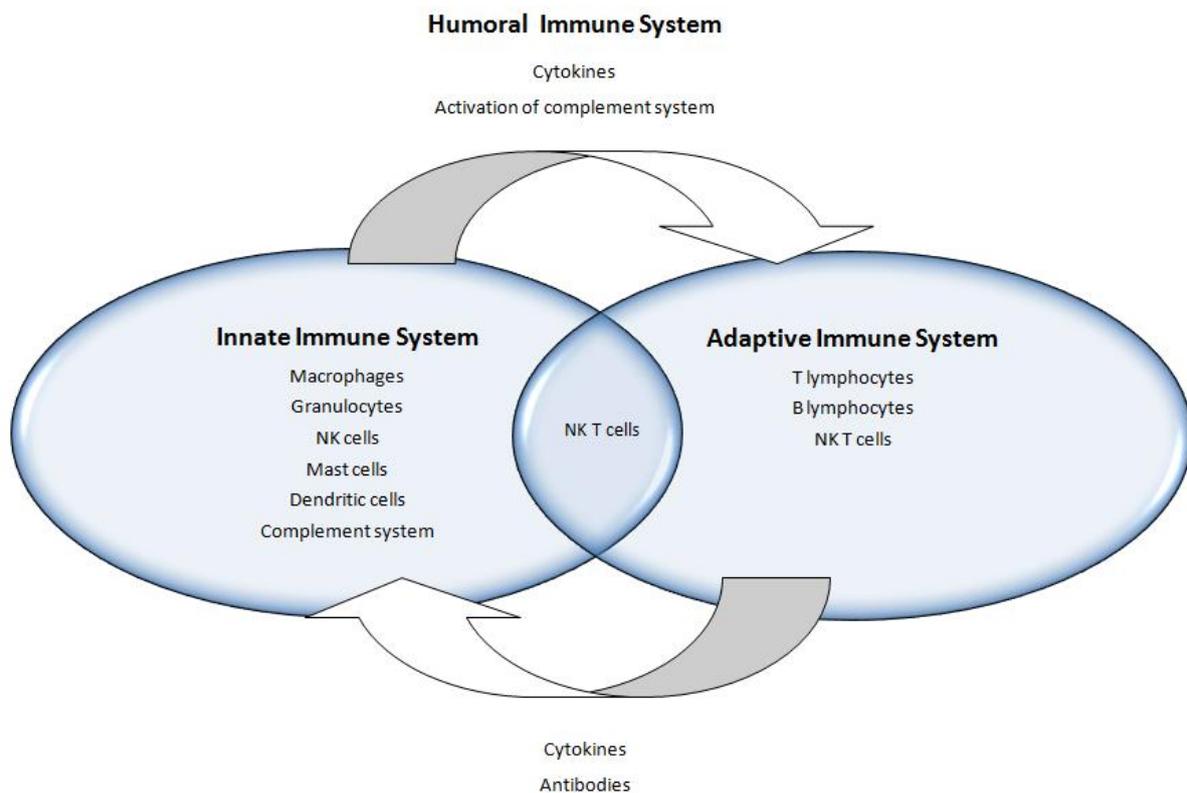


Figure 1.1. Interactions between the innate, humoral and adaptive immune systems [de Visser and Coussens, 2005]

1.1.3. The Humoral Immune System

The humoral system refers to the non-cellular components of the immune system found in extracellular fluid, in particular the serum. It is normally initiated by the innate immune-based local inflammatory response and triggers a cascade of immunological

processes that help link and co-ordinate the innate and adaptive immune/inflammatory responses (Figure 1.1). It is composed of a number of blood based proteins including antibodies, constituents of the complement pathway, cytokines and opsonins that promote pathogen cell lysis and phagocytosis [Janeway et al., 2001]. Indeed, following tissue damage from infection, chemical injury, trauma, malignancy or immune based destruction, a wide variety of growth factors including colony stimulating, as well as a variety of chemotactic cytokines that attract monocytes and stimulate them to transform into macrophages, are released. As well as the recruitment of innate and adaptive immune cells and promotion of pathogen death, humoral immunity also helps to drive the resulting systemic immune/inflammatory response [Baumann and Gauldie, 1994; Janeway and Medzhitov, 2002].

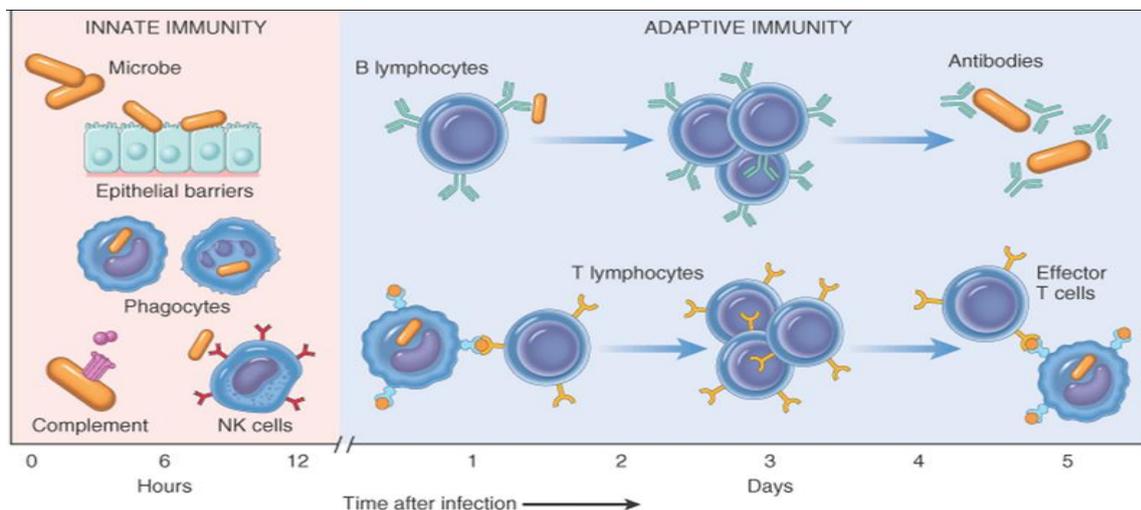


Figure 1.2. The temporal relationship between the innate and adaptive immune system [Abbas et al., 2005]

1.1.4. The Adaptive Immune System

The adaptive immune system is predominantly composed of lymphocytes, and through the recognition of foreign antigens, aims to develop immunological memory and

speed the elimination of future pathogen attacks [Janeway and Medzhitov, 2002]. The adaptive immune system can be classified as antibody mediated (part of the humoral response) or cell mediated, although many of the processes are interrelated. As illustrated in Figure 1.2, the antibody mediated response relies on phagocytic cells of the innate immune system, such as macrophages and neutrophils, as well as dendritic cells and B-lymphocytes, to present non-self peptides to T-lymphocytes and produce circulating antibodies [Unanue, 2002]. Cell mediated immunity results in antigen-specific cytotoxic T-lymphocytes recognition and phagocytosis of previously exposed pathogens [Herberman and Holden, 1978; Santoni et al., 1979].

1.1.5. The Systemic Immune/Inflammatory Response

Following a local immune/inflammatory reaction, and activation of the innate, humoral and adaptive immune systems, circulating pro-inflammatory mediator can result in a subsequent systemic immune/inflammatory response. Indeed, a correlation has been demonstrated between local cytokine production and systemic inflammation in patients with a variety of inflammatory processes [Vernooy et al., 2002; Grigoryev et al., 2008]. While the immune system aims to promote healing and prevent further injury by destroying harmful or infectious agents, the systemic immune/inflammatory response, also referred to as the acute phase reaction, can influence neuroendocrine, haematopoietic, metabolic and hepatic regulation with a number of associated negative physiological outcomes (Table 1.1.) [Baumann and Gauldie, 1994; Koj, 1996]. As part of the systemic immune/inflammatory response, circulating serum constituents of the fore-mentioned innate, humoral and adaptive immune systems, as well as a number of other physiological markers are altered. The systemic immune/inflammatory response has often been overlooked during the study of inflammation, disease pathogenesis and progression, however it is becoming increasingly

evident that this process is closely associated with outcome. This thesis will therefore concentrate on the systemic, opposed to the local immune/inflammatory response.

Table 1.1. The acute phase phenomena

Neuroendocrine
Fever, somnolence, and anorexia
Increased secretion of corticotropin-releasing hormone, corticotropin, and cortisol
Increased secretion of arginine vasopressin
Decreased production of insulin-like growth factor I
Increased adrenal secretion of catecholamines
Hematopoietic changes
Anaemia of chronic disease
Leukocytosis
Thrombocytosis
Metabolic changes
Loss of muscle and negative nitrogen balance
Decreased gluconeogenesis
Cachexia and osteoporosis
Increased hepatic lipogenesis
Increased lipolysis in adipose tissue and decreased lipoprotein lipase activity in muscle and adipose tissue
Hepatic changes
Increased metallothionein, inducible nitric oxide synthase, heme oxygenase, manganese superoxide dismutase, and tissue inhibitor of metalloproteinase-1
Decreased phosphoenolpyruvate carboxykinase activity
Changes in non-protein plasma constituent
Hypo zincemia, hypoferrremia, and hypercupremia
Increased plasma retinol and glutathione concentrations
[Gabay and Kushner, 1999]

1.1.6. Acute Phase Proteins

During the systemic immune/inflammatory response, as a result of physiological changes in protein metabolism and distribution, serum protein levels fluctuate [Fleck, 1989]. Acute phase protein have been defined as those demonstrating a plasma concentration change of at least 25% during an immune/inflammatory reaction [Kushner, 1982]. There are around forty acute phase proteins and these can be positive, (increasing in concentration) or negative (reducing in concentration) [Gabay and Kushner, 1999]. Immunomodulating cytokines, including IL-1, IL-6, IL8 and tumour necrosis factor α , are released from activated cells of

the innate immune system and, under glucocorticoid and growth factor regulatory control, act predominantly on the liver to synthesise positive acute phase proteins [Balkwill and Burke, 1989; Kulkarni and Karlsson, 1993; Wigmore et al., 1997], some of which have immunological roles and function as part of the humoral immune system.

As shown in Figure 1.3, the majority of acute phase proteins are positive and increase in serum concentration during an immune/inflammatory response. Examples of positive proteins include those of the complement system (C3, C4 and factor B), the coagulation and fibrinolytic system (fibrinogen, plasminogen, urokinase and protein S), antiproteases (α 1-

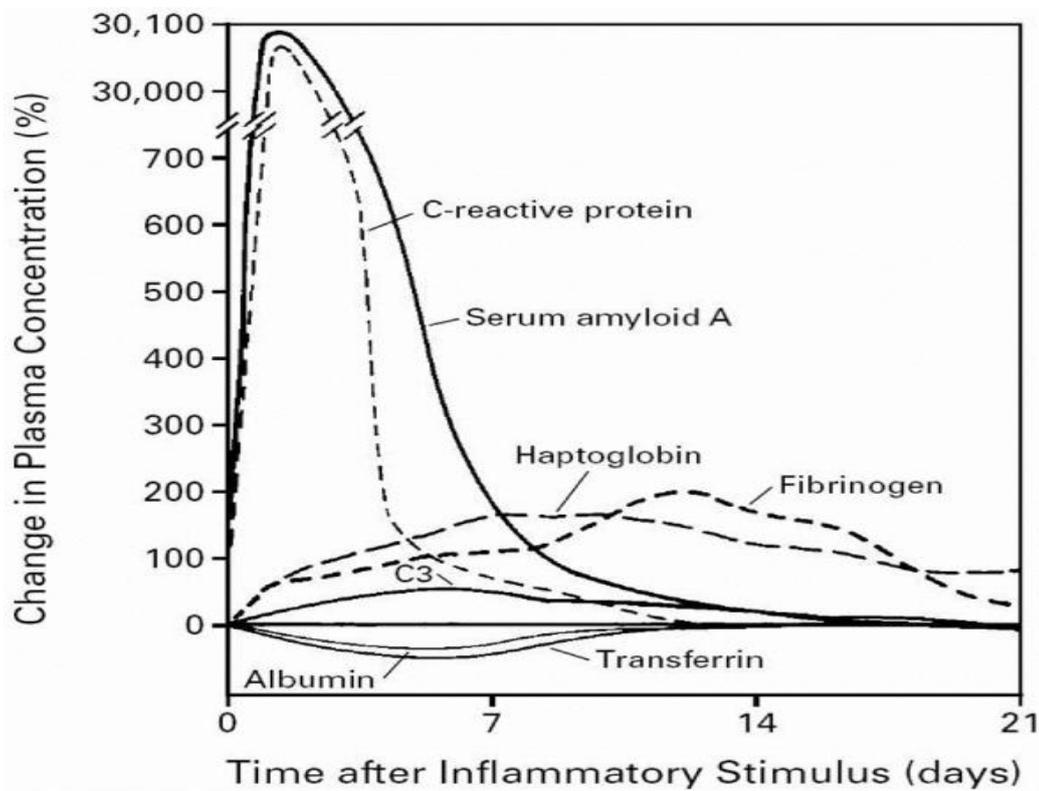


Figure 1.3. The temporal relationship of acute phase proteins [Gabay and Kushner, 1999]

Antichymotrypsin and α 1-Protease inhibitor), transport proteins (Haptoglobin) as well as many others including C-reactive protein and serum amyloid A [Gabay and Kushner, 1999]. Examples of negative proteins include albumin, transferrin, alpha-fetoprotein and insulin-like growth factor I [Gabay and Kushner, 1999].

In 1930, C-reactive protein was the first acute phase protein to be discovered [Tillett and Francis, 1930] and is now one of the most readily available markers of the systemic immune/inflammatory response. C-reactive protein belongs to the pentraxin family of calcium-dependent ligand-binding plasma proteins. It predominantly binds to phosphocholine residues found in necrotic and apoptotic cell walls [Thompson et al., 1999] but has also been shown to have an affinity for extrinsic ligands on bacterial, fungal and parasitic microorganism [Du Clos, 1996]. Binding of these ligands results in the aggregation and precipitation of such structures, initiation of the complement system and subsequent opsonisation and phagocytosis [Mold et al., 1999].

C-reactive protein is currently one of the most commonly used markers of systemic inflammation in clinical practice with the median concentration of C-reactive protein having been found to be 0.8mg/l in healthy individuals. It is also of interest that 90% of healthy people have a concentration of less than 3.0mg/l and 99% a concentration of less than 10mg/l [Shine et al., 1981]. As illustrated in Figure 1.3, serum concentrations peak around 48 hours following insult with a half life of around 19 hours [Vigushin et al., 1993]. It is also known to be one of the most sensitive acute phase proteins and can vary in magnitude by up to 1000 times [Gabay and Kushner, 1999]. C-reactive protein levels in the general population tend to be stable with no seasonal change and only minor yearly variations [Macy et al., 1997; Pepys and Hirschfield, 2003]. However, C-reactive protein concentrations are known to vary across

a number of parameters including body mass index (BMI) [Ford, 1999] and smoking [Wong et al., 2001].

Serum albumin is also commonly sampled in clinical practice and is known to reduce in serum concentration during the acute phase response [Ritchie et al., 1999; McMillan et al., 2001]. Serum albumin is produced in the liver and accounts for the majority of hepatic protein synthesis [Williams and Scott, 1998] with serum levels being influenced by inflammatory cytokine regulation of hepatic production, as well as protein degradation and redistribution [Goldwasser and Feldman, 1997]. A number of conditions, including sepsis, malignancy and chronic disease, can result in increased vascular permeability, albumin redistribution into extra-vascular tissues and subsequent reduction in serum albumin levels [Fleck et al., 1985]. Factors known to influence serum albumin levels include gender [Ritchie et al., 1999], daily activity, muscle mass [Baumgartner et al., 1996] and smoking [Salive et al., 1992].

1.1.7. Liver Function Tests, Calcium and Inflammation

Commonly sampled serum biochemical parameters also affected during the systemic immune/inflammatory response include liver function tests and serum calcium levels. A number of liver function tests have been shown to be associated with an inflammation associated C-reactive protein rise [Kerner et al., 2005; Cheung et al., 2008] and alkaline phosphatase has been associated with altered levels of other acute phase proteins including albumin [Blayney et al., 2008]. Similarly, gamma-glutamyl transferase, thought to be an early marker of oxidative stress, has been associated with raised C-reactive protein levels [Lee and Jacobs, 2005]. Bilirubin, known to have endogenous antioxidant properties, has been found to have lower serum concentrations in the presence of an active leukocytosis

[Stocker et al., 1987; Greabu et al., 2001]. There is also evidence that liver function tests are altered as part of the systemic immune/inflammatory response in conditions including lung and gastrointestinal cancer [Brown et al., 2007; Roxburgh et al., 2009a]. Furthermore, it has been suggested that the systemic immune/inflammatory response and C-reactive protein have an effect on cellular calcium metabolism [Foldes-Filep et al., 1992].

1.1.8. The Chronic Systemic Immune/Inflammatory Response

If the precipitating factor leading to an acute phase response is persistent, this process can be prolonged and a chronic systemic immune/inflammatory state will occur [Gabay and Kushner, 1999]. Examples of chronic local immune/inflammatory conditions that may lead to a chronic systemic immune/inflammatory response include atherosclerotic lesions [Ross, 1999], chronic infection [Cassell, 1998; Moutsopoulos and Madianos, 2006], chronic pancreatitis [Nair et al., 2007] and malignancy [Rakoff-Nahoum, 2006]. This ongoing process is sometimes referred to as the "Systemic Inflammatory Response" that, in part, involves alterations of the pre-mentioned circulating constituents of the immune system and associated serum biochemical variables. It is this phenomenon and its associations with outcome, predominantly in cancer but also in arteriosclerotic disease, that will be focused on in this thesis.

1.2. INFLAMMATION AND CANCER

1.2.1. Cancer Incidence

In 2008, it was estimated that there were 12.4 million new cancer cases diagnosed worldwide. Traditionally cancer has been a disease of Western, high income countries, however, this is changing. As shown in Figure 1.4, over half of all newly detected cancers are now occurring in developing countries [Boyle and Levin, 2008].

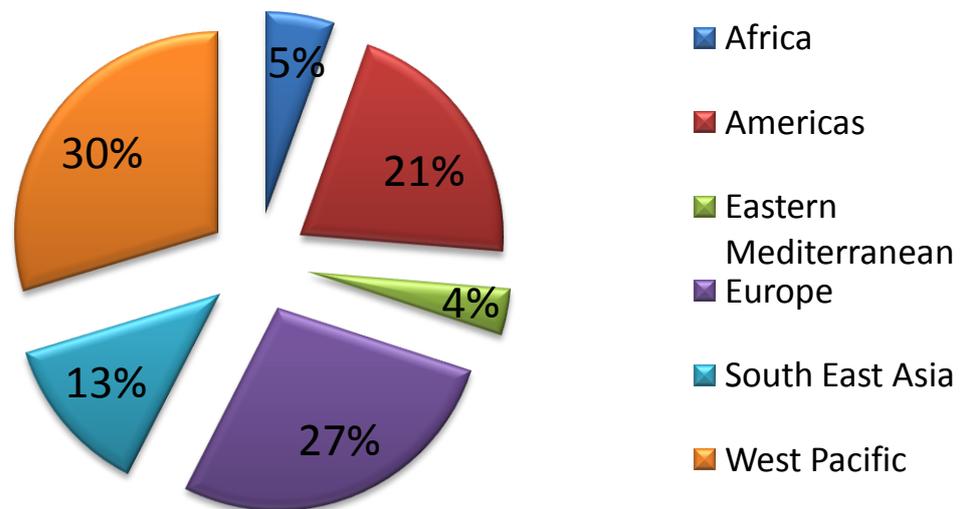


Figure 1.4. Worldwide cancer incidence by region in 2008 [Boyle and Levin, 2008]

With a growing global population and increasing life expectancy, the current incidence figures are set to rise to at least 20 million by 2030 [Boyle and Levin, 2008]. This is expected to be a global phenomenon with a projected increase in European regions from 4.1 million cases per annum in 2008 to at least 5.5 million per annum in 2030 [Boyle and Levin, 2008]. In the United Kingdom cancer incidence has also risen by approximately 25% between 1975 and 2004. As a consequence, more than one in three people will develop

cancer during their lifetime [Cancer Research UK, 2008] with similar projected increases in the number of cases forecasted [Moller et al., 2007].

Worldwide, lung cancer is the most commonly diagnosed tumour with approximately one million new cases in 2002. The incidence of lung cancer varies from approximately 2 per 100 000 in African women to 61 per 100 000 in North American men [Kamangar et al., 2006], a difference largely attributable to cigarette smoking [Ezzati and Lopez, 2003]. Colon cancer is approximately 25 times more common in developed countries, a difference partially attributable to differences in meat, fat and fibre consumption [Trock et al., 1990; Willett et al., 1990]. Female breast cancer is also more common in developed populations, a difference possibly due to an older population and increasing detection since the introduction of screening [Alberg and Singh, 2001].

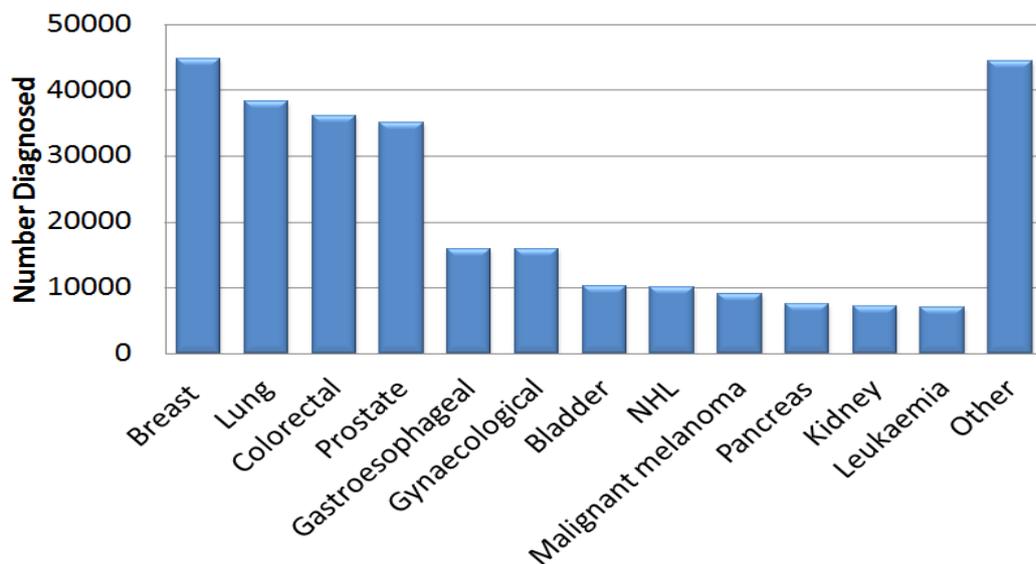


Figure 1.5. Number of cancers diagnosed in the UK per year. [Cancer Research UK, 2008]

Of the 280 000 new cancers diagnosed each year in the United Kingdom (Figure 1.5), the majority occur in people over 60 years of age [Cancer Research UK, 2008]. During 2004

the commonest cancers diagnosed in males were prostate (34 986), lung (22 495), colorectal (19 657), gastroesophageal (10 100) and bladder (7168). During the same time period, in females, the most commonly diagnosed malignancies were breast (44 335) followed by colorectal (16 452), lung (15 818), gynaecological (15 779) and gastroesophageal (5732).

1.2.2. Cancer Mortality

Worldwide, in 2008 there were an estimated 7.6 million deaths attributed to cancer with projected figures to reach at least 12.9 million by 2030 [Boyle and Levin, 2008]. Survival from common cancers including breast, colorectal and prostate, is generally better for those diagnosed in developed countries of western Europe, North America and Australia, when compared to less developed areas including eastern Europe, Algeria and Brazil [Coleman et al., 2008].

In the United Kingdom cancer is the leading cause of death and responsible for approximately thirty percent of all mortality, equating to around 150,000 deaths per year [Cancer Research UK, 2007]. Most deaths occur in males and those over the age of 75 with an increased mortality rate in the socioeconomically deprived [Coleman et al., 2004]. The commonest cancers responsible for death are lung, bowel, breast and prostate with the relative five year survival of these, and other common cancers illustrated in Figure 1.6.

In recent decades there have been improvements of approximately ten percent in cancer mortality rates with these changes largely affecting people under the age of 75 with cancers of the stomach, lung, bladder, bowel and cervix [Cancer Research UK, 2006]. These improvements are likely due to better detection and treatment modalities, however, with increasing population age a rise of 40% in cancer deaths is expected by 2025 [Olsen et al.,

2008]. This clearly has significant financial implications and will require a greater proportion of annual healthcare spending [Bosanquet and Sikora, 2004; Boyle and Levin, 2008].

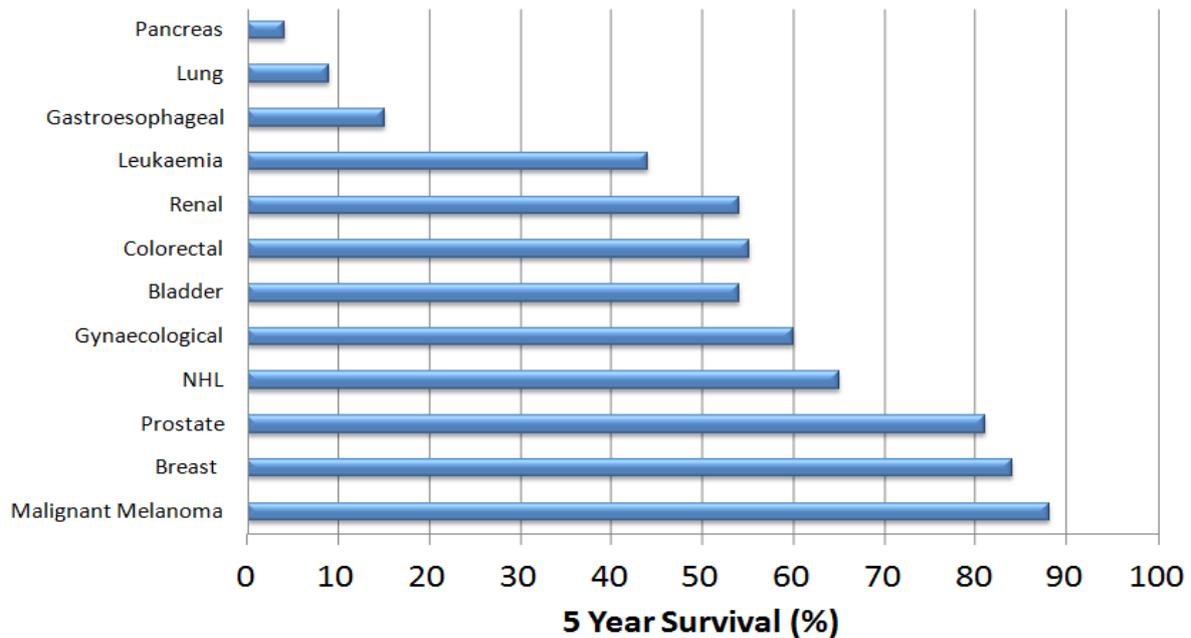


Figure 1.6. Relative 5 year survival in common tumour types in the UK. [Cancer Research UK, 2007]

1.2.3. Patient Demographics and Cancer Survival

A number of patient based factors are known to be related to survival in patients with cancer. Increasing age is associated with reduced survival when adjusted for other known factors in the majority of cancers [Sant et al., 2003; Lam et al., 2007]. In part, this may be explained by inadequate staging and fewer treatment options in the elderly [de Rijke et al., 1996; Fentiman, 1996]. There are however certain tumour types, specifically lung, breast and prostate, that do not conform to this pattern with some evidence that older patients have an improved survival [Romano and Mark, 1992; Merrill and Bird, 2002; Chia et al., 2004; Mathew et al., 2004].

Male gender is also associated with reduced survival in variety of tumour types that can affect both sexes. For example, male gender tends to be associated with a poorer outcome in patients with colorectal and lung cancer, as well as chronic lymphocytic leukaemia [Romano and Mark, 1992; Diehl et al., 1999; Wichmann et al., 2001; McArdle et al., 2003; Sant et al., 2003; Visbal et al., 2004].

Furthermore, increasing socioeconomic deprivation has been associated with reduced survival in a number of different tumour types including lung, breast, colorectal, bladder, prostate, uterine and cervical, independent of other prognostic factors [Schrijvers et al., 1995; Coleman et al., 2004]. In population studies, deprivation has traditionally been measured by the Carstairs index [Sloggett and Joshi, 1998], a postcode based score derived from a number of socioeconomic factors including car ownership, household crowding, head of household social class and male unemployment [Carstairs and Morris, 1991]. Currently, the recommended measurement of deprivation in Scotland is the (SIMD). Similarly it categorises post code areas, however, a more detailed measurement of 37 indicators across seven domains including income, employment, education, housing, health, crime and geological area is used [Bishop et al., 2004]. Scotland is divided into 6,505 data zones with these ranked from 1 (least deprived) to 6,505 (most deprived). These data zones have further been divided into quintiles of the Scottish population, with the present body of work using quintiles that range from 1 (least deprived) to 5 (most deprived). As can be seen in Figure 1.7, an increasing SIMD score has been shown to be associated with an increased cancer incidence as well as an increased cancer associated mortality in the Scottish population [The Scottish Government, 2008].

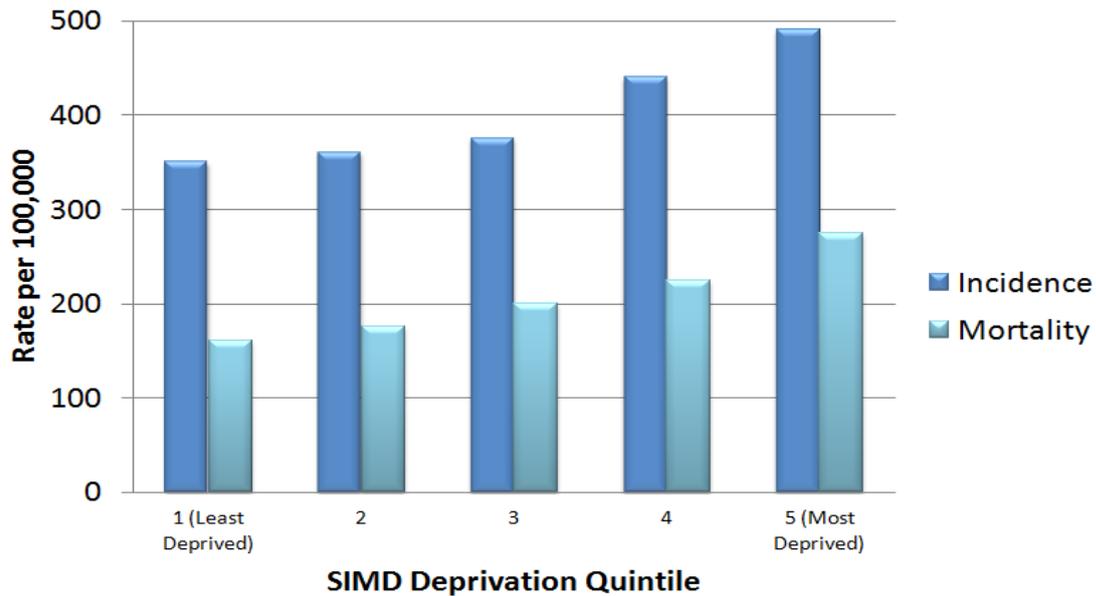


Figure 1.7. Malignant neoplasm age-standardised incidence and mortality by SIMD Quintiles [The Scottish Government, 2008]

1.2.4. Inherited Genetic Mutations and Carcinogenesis

The development of malignancy is a multi-step process depending on the inheritance or development of deoxyribonucleic acid (DNA) mutations that result in the interruption of tumour suppression or activation of pro-oncogenic pathways at a cellular level [Kinzler and Vogelstein, 1996]. Genetic mutations ultimately result in the development of several cellular characteristics, including invasion and the potential to metastasise, that are required for malignant transformation [Hanahan and Weinberg, 2000].

It has long been accepted that cancer has a genetic basis with a variety of tumours noted to have potentially hereditary patterns [Easton, 1994]. In twin studies it has been estimated that genetic predispositions are a contributing factor in approximately 28% of gastric, 35% of colorectal, 27% of breast and 42% of prostate cancers [Lichtenstein et al., 2000]. A number of genes have been identified as increasing cancer risk including BRCA1 and 2 in breast cancer, APC, MSH2, MLH1 in colon cancer and MET in renal cell cancer.

1.2.5. Acquired Genetic Mutations and Carcinogenesis

DNA mutations, whether they result in the activation of oncogenes or the loss of tumour suppressor genes, can occur as a consequence of point mutations, insertions or deletions of base pairs, gain or loss of entire chromosomes, translocations and gene amplifications [Lengauer et al., 1998]. Acquired mutations are thought to be responsible for the majority of DNA mutations and can be secondary to oxidative stress or radiation [Lichtenstein et al., 2000; Hemminki and Jiang, 2002]. Oxidative stress can occur as the result of contact with a number of carcinogens including those found in food [Wong et al., 2001; Straif et al., 2006; Baan et al., 2007], cigarette smoke [Wogan et al., 2004] and other chemical compounds [Baan et al., 2008].

1.2.6. Inflammation and Carcinogenesis

There is growing evidence that a local inflammatory microenvironment encourages the development of malignancy with an increased prevalence of cancer in tissues experiencing chronic inflammation [Balkwill and Coussens, 2004; Vakkila and Lotze, 2004; Lu et al., 2006]. Continued immune/inflammatory stimuli secondary to chemical irritation [Whitcomb, 2004], infection [Pagano et al., 2004] or immunological disorders [Itzkowitz and Yio, 2004] can result in the release of a variety of tumour promoting cytokines [Lin and Karin, 2007]. Increased tissue turnover, loss of tissue architecture and oxidative stress can result in DNA damage with subsequent risk of neoplastic transformation and eventual cancer development [Shacter and Weitzman, 2002; Balkwill et al., 2005; Colotta et al., 2009].

Globally, approximately 17% of all cancers are attributed to infectious aetiology [Kuper et al., 2000; Parkin, 2006]. For example, chronic viral hepatitis has been implicated in the pathogenesis of the majority of hepatocellular cancers [Kew, 1998] and the human

papilloma virus has been clearly associated with the development of cervical cancer [Bosch et al., 2002].

Non-infective, autoimmune inflammatory conditions also predispose individuals to the development of neoplasia. Colorectal cancer incidence increases by approximately 10 fold in patients with chronic inflammatory conditions of the colon, such as ulcerative colitis [Itzkowitz and Yio, 2004]. Fibrotic and inflammatory lung conditions, including idiopathic pulmonary fibrosis and systemic sclerosis, have also been shown to increase risk the of developing cancer [Daniels and Jett, 2005]. Similarly, patients with systemic lupus erythematosus (SLE) have a higher incidence of cervical dysplasia [Blumenfeld et al., 1994].

1.2.7. Inflammation and Cancer Progression

It has been suggested that essential alterations in cell physiology, including self sufficiency in growth, sustained angiogenesis and the potential for tissue invasion and metastasis, are essential if tumours are to succeed in malignant growth [Hanahan and Weinberg, 2000; Colotta et al., 2009]. It is becoming clear, as detailed in Figure 1.8, that the oncogenic process of tumour development and progression is influenced by the interaction of surrounding non-malignant cells [Hanahan and Weinberg, 2000; Coussens and Werb, 2002] and it is therefore of interest that certain malignant tumours may consist of over 90% stromal and inflammatory cells [Mesker et al., 2007]. Moreover, this is consistent with growing evidence that suggests a pro-inflammatory tumour microenvironment is vital for the progression of malignancy [Balkwill and Coussens, 2004; Mantovani et al., 2008; DeNardo et al., 2008].

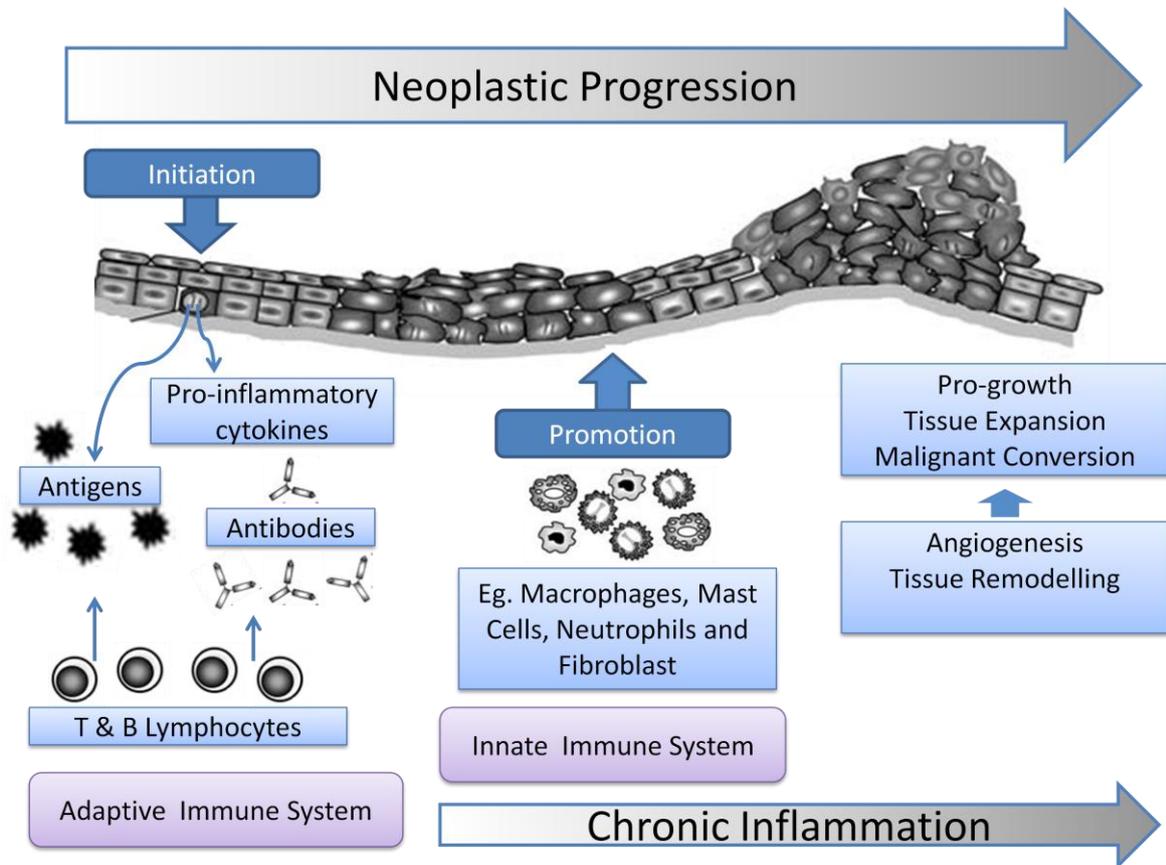


Figure 1.8. Inflammation, carcinogenesis and neoplastic progression [de Visser et al., 2006]

To achieve malignant growth, cancer cells hijack normal immunological and physiological processes to aid angiogenesis and stromal tissue development. For example, infiltrating cells belonging to the innate immune system, including neutrophils, macrophages and mast cells will increase the angiogenic potential of a tumour and allow it to lay down a supportive connective tissue framework [Hanahan and Folkman, 1996; Jackson et al., 1997; Imada et al., 2000; Benitez-Bribiesca et al., 2001; Kalluri and Zeisberg, 2006; Joyce and Pollard, 2009].

Tumour Associated Macrophages (TAMs) are the predominant tumour leukocyte infiltrate and have been implicated in both the destruction of tumour cells as well as the promotion, in part by cytokine production, of cancer progression and metastasis [Jonjic et al.,

1992; Mantovani et al., 2002; Pollard, 2004]. Macrophages have also been shown to have an atrophic, remodelling role that contributes to tumour progression and cancer cell survival by increasing the proliferative potential, increasing cancer cell motility, intravasation and invasiveness as well as increasing angiogenesis, mediating immunosuppression and promoting extracellular matrix formation [Mantovani et al., 2002; Lewis and Pollard, 2006; Nardin and Abastado, 2008]. While it remains a controversial issue, there is some corresponding evidence that TAMs are associated with reduced survival in certain tumour types [Leek et al., 1996; Bingle et al., 2002].

While there is evidence that a number of local inflammatory cells can promote tumour progression, conversely, an abundance of tumour margin lymphocytes has been associated with an improved outcome [Jass et al., 1987]. Indeed, this lymphocytic infiltrate has been, in colorectal, hepatocellular and gastroesophageal cancer, shown to correlate with improved survival [Ma et al., 1994; Wada et al., 1998; Ohashi et al., 2000; Canna et al., 2005]. This is clearly a complex relationship as low intertumoral T regulatory cells, in the presence of high cytotoxic T cells (CD8+), have also been associated with improved survival [Gao et al., 2007]. Moreover, it has been suggested that the inverse relationship between cytotoxic T cell infiltrate and survival may be due to the inhibition of micrometastasis [Chiba et al., 2004] and tumour induced modulation, at the lymph node level, may inhibit or down regulate the action of cytotoxic T-cells and therefore increase the potential for distal spread [Cochran et al., 2006].

Neutrophils produce cytotoxic substances, including proteinases, to defend the host from microorganism attack and promote healing [Smith, 1994]. They can also produce cytokines to drive the immune/inflammatory response with evidence to show that tumour associated neutrophils can play an active part in tumour angiogenesis and proliferation

[Nathan, 2006; Tazzyman et al., 2009]. Neutrophilia is also known to occur as a result of cancer related production of myeloid growth factor, granulocyte colony stimulating factor and cytokine production [Ulich et al., 1987; Lord et al., 1989].

Platelets are also influenced by malignancy and can be activated by direct contact with tumour cells or via release of tumour based cytokines [Zucchella et al., 1989; Grignani et al., 1989]. They have been shown to play a multifactorial role in angiogenesis, the pro-coagulopathy observed in malignancy as well as tumour progression [Poggi et al., 1988; Sierko and Wojtukiewicz, 2004].

1.2.8. Cancer Stage and Indicators of Survival

Cancer staging is important as it currently helps identify those patients who will benefit from potentially curative interventions such as resectional surgery or oncological therapy. It also provides an indication of prognosis and guides the requirement for surveillance following treatment. The most commonly used staging system for malignant tumours is the TNM classification (Table 1.2) which is recommended by the Union for International Cancer Control (UICC), Cancer Research UK as well as the American Joint

Table 1.2. Tumour Node Metastasis classification

Primary Tumour	
T0	No evidence of malignant tumour
Tis	Carcinoma in situ (preinvasive)
T1-T4	Size and/or extent of primary tumour
Regional Lymph Nodes	
N0	No regional lymph node involvement
N1-3	Degree of regional lymph node involvement (number and location)
Distant Metastasis	
M0	No distant metastasis
M1	Distant metastasis present

[National Cancer Institute, 2009]

Committee on Cancer (AJCC). TNM is based on the extent of the primary tumour (T), spread to the local lymph nodes (N) and the presence of distal metastasis (M) [Sobin and Wittekind, 2002]. It can be calculated by clinical means, including examination and radiological investigations, objective assessment of resected pathological specimens or a combination of these techniques. A shortened TNM classification comprising five stages (0-IV) is also commonly used, but differs in detail depending on the specific tumour type (Table 1.3). In colorectal cancer, this broadly corresponds to the Dukes classification where the more advanced the tumour, the shorter the expected survival [National Institute for Health and Care Excellence, 2009]. While the TNM system is used in the majority of tumour types, some central nervous system and haematological cancers do not comply and different systems are used.

Table 1.3. Tumour stage classification

Stage 0	Carcinoma in situ
Stage I-III	Higher number indicates more extensive disease. Larger tumour size and/or spread of the cancer beyond the organ in which it first developed to nearby lymph nodes and/or tissue or organs adjacent to the location of the primary tumour.
Stage IV	Distal metastasis

[National Cancer Institute, 2009]

While being considered the gold standard for staging prognosis, the TNM based classification has potential pitfalls. It is clear that staging by clinical and radiological means may be open to a certain amount of subjective judgment and it has been shown that computerised tomography may provide different results from resected pathological specimens [Schlomer et al., 2006]. There are also concerns that inadequate pathological assessment, with regards to number of lymph nodes examined, may result in under staging and not effectively identify individuals who would benefit from adjuvant therapy [Tepper et

al., 2001; Joseph et al., 2003; Swanson et al., 2003]. These issues have compounded worries that while this system may be adequate for those at the extremes of the stage spectrum (I or IV), those in the middle may be getting inaccurately staged and under or over treated.

A number of other tumour related, pathological features are associated with outcome and survival in a variety of tumour types. Tumour differentiation, as detailed in Table 1.4, is a measure of how abnormal tumour cells appears and a reflection of how rapidly a tumour is likely to grow and disseminate. Tumour differentiation has been associated with outcome in the majority of neoplastic malignancies including gastroesophageal [Khan et al., 2004], breast [Bloom and Richardson, 1957], pancreatic [Lim et al., 2003] and colorectal [Mori et al., 2006] cancers.

Table 1.4. Grade of differentiation classification

Grade 1	Well differentiated (low grade)
Grade 2	Moderately differentiated (intermediate grade)
Grade 3	Poorly differentiated (high grade)
Grade 4	Undifferentiated (high grade)

[National Cancer Institute, 2009]

While these methods are clearly important and provide a wealth of information regarding tumour status, they do little to estimate the host response to the malignant process. Performance status can be used as a potential indicator of host response and aims to measure physical function and gauge how malignancy impacts on a patient's daily life. The Eastern Cooperative Oncology Group (ECOG) performance status score is a 5 point system that ranges from 0 (no symptoms/restriction to activity) to 5 (death) and is commonly used when making decisions as to whether a patient is fit enough to receive chemotherapy [Oken et al.,

1982]. There have however been concerns that these subjective measures of patient function have been shown to have significant inter-observer variability [Ando et al., 2001]. As performance status is a measure of host function, and therefore a surrogate marker of cancer cachexia and muscle mass, it is also of no surprise that an association between the host systemic immune/inflammatory response and performance status has been demonstrated [Brown et al., 2007].

Taken together, it is therefore expected that there is increasing interest in the "seed and soil" hypothesis that suggests patient outcome depends not only on tumour characteristics but also on the host immunological response to a malignant process [Fidler and Poste, 2008]. It therefore stands that a measure of the pre-mentioned host immune/inflammatory response may offer a useful adjunct in the risk stratification of patients with cancer.

1.3. INFLAMMATION AND ATHEROSCLEROSIS

1.3.1. Cerebrovascular and Cardiovascular Atherosclerotic Disease and Mortality

According to the World Health Organisation, cardiovascular and cerebrovascular disease are responsible for the majority of mortality worldwide. In 2004, ischaemic heart disease was responsible for approximately 7.2 million deaths, thus accounting for over 12 percent of all deaths worldwide. The World Health organisation predict that this will increase to approximately 23 million by 2030 and concerns have been raised that cardiovascular disease may become the most common cause of death in the 21st century [Sporn, 1996; Hansson, 2005]. Similarly, cerebrovascular disease was responsible for approximately 10 percent of all deaths worldwide in 2004 with this figure set to increase [Mathers et al., 2004]. In high income countries this pattern persisted with approximately 16 percent of deaths being attributable to ischaemic heart disease and 9 percent as a result of cerebrovascular disease [Mathers et al., 2004]. These reported death rates are consistent with those published for the UK with approximately one third of all deaths being due to circulatory disease [Office for National Statistics, 2008; McNiven D, 2010].

1.3.2. Atherosclerotic Disease and Inflammation

The link between atherosclerotic disease and inflammation is well recognised and has been shown to be independent of other arteriopathic risk factors including smoking and obesity [Kuller et al., 1996; Tracy, 1998; Ross, 1999; Williams et al., 2004]. While the relationship between inflammation, C-reactive protein and atherosclerotic disease is not entirely clear, arterial wall lipoprotein retention and the subsequent formation of fatty streaks has been implicated [Ross, 1999; Pepys and Hirschfield, 2003]. Once established, fatty streaks can encourage fibrous plaque formation, which in turn can rupture, precipitate

thrombus and result in a cardiac event [Ross, 1999]. Atherosclerotic vessel wall inflammatory infiltrate has been found to have increased levels of a number of inflammatory cells including lymphocytes and dendritic cells [Galkina and Ley, 2009]. Macrophages have also been implicated and, as illustrated in Figure 1.9, have been found to oxidise low density lipoproteins and drive an immune/inflammatory response [Jonasson et al., 1986; Berliner et al., 1990]. Neutrophils and platelets are similarly involved in atherosclerotic disease and have been shown to have a pro-inflammatory role with the subsequent propagation of atheromatous plaque and potential rupture [Ross, 1993; Massberg et al., 2002; Kraaijeveld et al., 2007; von Hundelshausen and Weber, 2007; Zernecke et al., 2008]. These activated cells of the immune system involved in atheroma development also produce inflammatory

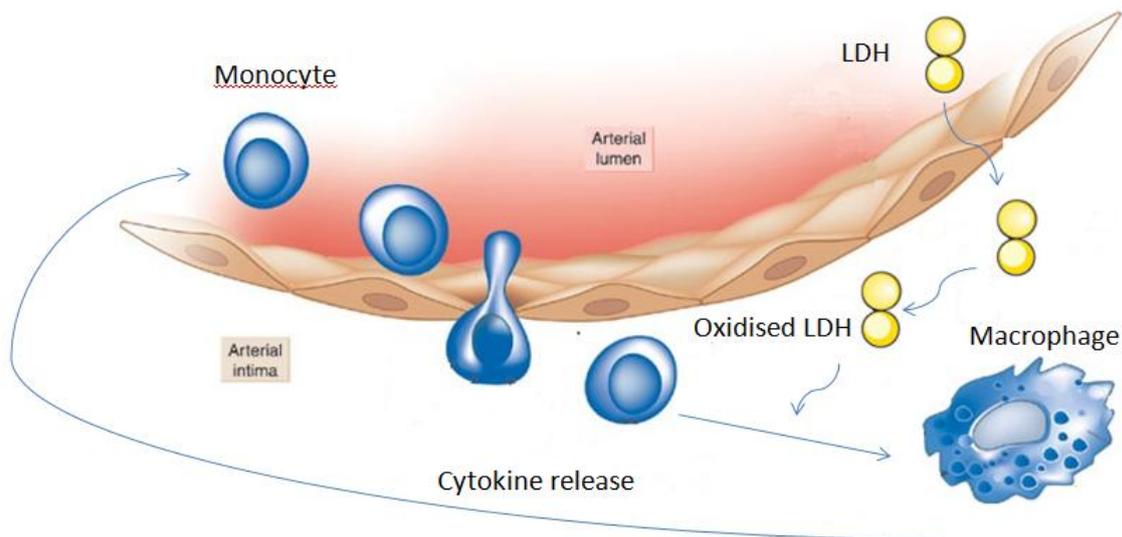


Figure 1.9. Low density lipoprotein, inflammation and atherosclerotic vessel wall infiltrate [Barter et al., 2004]

cytokines including IL-1 and IL-6, and as a consequence, acute phase reactants including C-reactive protein from the liver [Hansson et al., 1989; Clinton and Libby, 1992; Frostegard et

al., 1999]. As a result, both IL-6 and C-reactive protein levels have been shown to be elevated in patients with unstable angina and myocardial infarction [Liuzzo et al., 1994; Lindahl et al., 2000; Danesh et al., 2008]. Traditionally, C-reactive protein levels of 10mg/l or less have been regarded as clinically unimportant, however smaller rises have since been classed as abnormal in those at risk, or with a history of ischaemic heart disease [Haverkate et al., 1997; Gabay and Kushner, 1999]. It is likely that these small rises reflect the local inflammatory process in the artery, however, it is also possible that they represent inflammation at a distal site [Hansson, 2005]. It is also of interest that albumin, a negative acute phase protein, can reduce platelet adhesion and has an anti-oxidant effect that helps clearance of reactive oxygen species that are pathogenic in ischaemic heart disease [Halliwell and Gutteridge, 1990; Porcellati et al., 1995].

1.3.3. Risk Stratification in Cardiovascular and Cerebrovascular Disease

A strategy that assists in the identification of those at increased risk can help individualise treatment and ensure that appropriate therapies are received by those who will benefit [Alderman, 1993]. The optimal prognostic tool would be of low financial cost, openly available and easily interpreted with a high sensitivity and specificity [Parikh and Vasan, 2007].

The extensively validated Framingham Risk Score can be used to calculate the probability of an individual having a major cardiac event and the requirement for therapy [Kannel et al., 1976; Anderson et al., 1991; Brindle et al., 2003; Greenland et al., 2010]. It includes gender, total cholesterol, HDL cholesterol, smoking status as well as systolic blood pressure and is recommended by European and American cardiac societies [Wood et al., 1998; Grundy et al., 1999].

A similar score, the CHADS₂, is used in the risk stratification for future cerebrovascular events in patients with atrial fibrillation. It looks at the presence of congestive heart failure, hypertension, age, diabetes and a history of prior cerebrovascular events and indicates the requirement for anticoagulation therapy [Gage et al., 2004].

As inflammation is pathologically related to atherosclerotic disease, C-reactive protein as a measure of the systemic immune/inflammatory response, is currently recommended by the American Heart Association for cardiovascular screening and statin management [Greenland et al., 2010]. It therefore stands that that other simple, easily obtainable measures of the systemic immune/inflammatory response may also predict outcome in patients with cardiovascular and cerebrovascular disease.

1.4. SERUM MARKERS OF SYSTEMIC INFLAMMATION AND MORTALITY

1.4.1. C-reactive Protein

A number of markers of the systemic immune/inflammatory response have been associated with all-cause mortality. In 274,515 incidentally sampled patients, Marsik and co-workers, demonstrated that a raised baseline serum C-reactive protein concentration was associated with mortality from cancer, cardiovascular and cerebrovascular disease [Marsik et al., 2008]. This study was noted to be particularly robust as patient cause of death was confirmed by a post mortem in approximately 35% of cases. Also, when 22,962 patients with subacute C-reactive protein measurements (<10mg/l) were studied, high sensitivity measurements were found to have additional value to the traditional thresholds (>10mg/l), when predicting all-cause mortality [Currie et al., 2008].

Serum C-reactive protein levels have also been associated with both the risk of developing cancer and a reduced survival in patients with cancer. In a cohort of 22,887 patients, a baseline elevated C-reactive protein level was associated with a greater risk of developing colorectal cancer, with similar studies demonstrating comparative findings for lung cancer [Erlinger et al., 2004; Allin et al., 2009]. Elevated levels of C-reactive protein have also been associated with survival in a number of malignancies including colorectal [Nozoe et al., 1998], oesophageal [Guillem and Triboulet, 2005; Crumley et al., 2006a], pancreatic [Falconer et al., 1995; Jamieson et al., 2005], ovarian [Hefler et al., 2008], endometrial [Schmid et al., 2007], cervical [Polterauer et al., 2007], breast [Albuquerque et al., 1995], renal cell [Casamassima et al., 2005; Komai et al., 2007] and hepatocellular [Hashimoto et al., 2005; Nagaoka et al., 2007] cancer.

Correlations between C-reactive protein concentrations and atherosclerotic disease have also been found. Koenig and co-workers demonstrated, in 936 healthy men, that a raised C-reactive protein was independently associated with an increased risk of developing cardiovascular disease [Koenig et al., 1999]. Also, meta-analysis of 1953 coronary events demonstrated a relative risk of 2.0 for the development of future coronary events if a patient's C-reactive protein was in the top tertile ($>2.4\text{mg/l}$ Vs $<1\text{mg/l}$) [Danesh et al., 2000]. Moreover, many other studies have shown similar associations with a raised C-reactive protein, in patients with unstable angina or myocardial infarction, and future cardiac events and death [Liuzzo et al., 1994; Ridker et al., 2000; Ridker et al., 2001; Danesh et al., 2004].

In patients with cerebrovascular disease, C-reactive protein has been found to have prognostic value in both identifying patients at risk of developing the condition, as well as those who are likely to die as a result. In a cohort of 7901 healthy Japanese men, those with raised C-reactive protein measurements were more likely to have a cerebrovascular event, as well as die from all-cause mortality [Makita et al., 2009]. Similar associations between a raised C-reactive protein and cerebrovascular related mortality have been noted in other studies [Di Napoli et al., 2001; den Hertog et al., 2009].

1.4.2. Albumin

Hypoalbuminaemia has been shown, in 7735 randomly enrolled men, to be associated with an increased risk of all-cause mortality independent of other risk factors [Phillips et al., 1989]. Similarly, hypoalbuminaemia was found to be a risk factor associated with all-cause mortality in the American National Health and Nutrition Examination Survey (NHANES) longitudinal follow-up survey [Gillum and Makuc, 1992] as well as in other large cohorts [Goldwasser and Feldman, 1997; Grimm et al., 2009; Desai et al., 2009].

Hypoalbuminaemia has also long been associated with reduced survival in patients with a variety of cancers [Phillips et al., 1989; Lam et al., 2007; Penel et al., 2008]. A number of studies have demonstrated a link between hypoalbuminaemia and survival in lung cancer. These include work carried out by Maeda and co-workers demonstrating this relationship in 261 patients with advanced Non-small Cell Lung Cancer (NSCLC) [Maeda et al., 2000] as well as a number of other studies with similar results [Hespanhol et al., 1995; Maestu et al., 1997; Tas et al., 1999]. In 1367 patients undergoing surgical treatment for colorectal cancer, hypoalbuminaemia was shown to be independently associated with a reduced survival [Sun et al., 2009] with comparable results being found in other operable and palliative cohorts [Heys et al., 1998; Schindl et al., 2005; Cengiz et al., 2006]. This association, between hypoalbuminaemia and reduced survival, has been shown to persist in a number of solid organ malignancies including gastric [Lien et al., 2004; Alici et al., 2006; Onate-Ocana et al., 2007], oesophageal [Di Fiore et al., 2007; Wang et al., 2009], hepatocellular cancer [Nagaoka et al., 2007; Takahashi et al., 2008; Choi et al., 2008], pancreatic [Siddiqui et al., 2007], ovarian [Gupta et al., 2009] and breast cancer [Wyld et al., 2003; Lis et al., 2003]. Similar studies have also shown this association to occur in haematological malignancies including, Non-Hodgkin's Lymphoma [Alici et al., 2003].

Serum albumin is also known to be associated with cardiovascular mortality in unselected patients and in those with pre-existing renal disease [Goldwasser and Feldman, 1997; Grimm et al., 2009]. Similarly, hypoalbuminaemia has been shown to be associated with an increased risk of death from cardiovascular disease independent of other risk factors [Phillips et al., 1989]. These findings were further confirmed in the American National Health and Nutrition Examination Survey (NHANES) [Gillum and Makuc, 1992] with several other studies also demonstrated this relationship [Kuller et al., 1991; Corti et al.,

1996; Weijenberg et al., 1997]. Hypoalbuminaemia has also been shown, after controlling for stroke risk factors, to predict cerebrovascular incidence as well as mortality in subjects of NHANES epidemiological follow-up study [Gillum et al., 1994].

1.4.3. Liver Function Tests and Calcium

Liver function tests have similarly been associated with outcome and survival. Bilirubin, a potential antioxidant, has been associated with a reduction in the incidence of coronary artery disease [Schwertner et al., 1994]. Higher levels of bilirubin, in approximately 10 000 unselected patients, has been linked with improved all-cause, cancer and cardiovascular survival [Temme et al., 2001]. Conversely, an elevated alkaline phosphatase has been associated with a reduced survival in patients with kidney disease, cardiovascular disease and in the general population [Blayney et al., 2008; Tonelli et al., 2009]. Furthermore, it is of interest that γ -glutamyl transferase (GGT) and serum calcium have also been reported to predict cancer and non-cancer outcomes [Leifsson and Ahren, 1996; Kazemi-Shirazi et al., 2007].

1.4.4. White Cell Count

A white blood cell count has likewise been shown to have prognostic value in previously health individuals. When 2011 men were followed up for an average of thirteen years, their initial white cell count corrected for lifestyle risk factors, was shown to predict all-cause mortality [de Labry et al., 1990]. In addition, a study of 13,555 individuals demonstrated positive correlation between white cell count and all-cause mortality, as well as risk of stroke and cardiovascular disease [Gillum et al., 2005] with a number of other studies demonstrating similar relationships [Tamakoshi et al., 2007].

In cancer cohorts, an elevated white cell count has been shown to have prognostic significance in a range of advanced malignancy including lung [Tibaldi et al., 2008], colorectal [Kohne et al., 2002], pancreatic [Engelken et al., 2003] and other gastrointestinal cancers [O'Gorman et al., 2000]. While these reported associations between a white count and mortality in both cancer and non-cancer groups is of interest, it remains to be seen whether a particular cell type, for example neutrophils, are predominantly responsible for this relationship.

1.4.5. Neutrophil Count

Elevated neutrophil counts have been associated with a reduced survival in renal cell and Non-Small Cell Lung cancer [Donskov, 2007; Teramukai et al., 2009]. In addition, the NHANES epidemiological follow-up study of approximately 5000 patients demonstrated a link between neutrophil count and all-cause and cardiovascular mortality [Gillum et al., 2005].

1.4.6. Lymphocyte Count

Conversely, a raised lymphocytes count seems to be of prognostic benefit and in elderly populations, lymphocytosis has been shown to reduce all-cause mortality [Bender et al., 1986; Izaks et al., 2003; Lukito et al., 2004]. Similarly, an increased lymphocyte count has been associated with an improved survival in patients with metastatic renal cell [Fumagalli et al., 2003], pancreatic cancer [Fogar et al., 2006] and sarcoma [Ray-Coquard et al., 2009]. The prognostic value of a low lymphocyte count has also been studied in patients with cardiac disease and congestive heart failure and has been found to be associated with increased mortality [Acanfora et al., 2001].

1.4.7. Platelet Count

Thrombocytosis, in patients with renal disease, has been shown to be associated with presence of coronary artery disease [Sokunbi et al., 1994]. It has also been linked with both tumour stage [Ito et al., 2006] and prognosis in renal cell carcinoma [O'Keefe et al., 2002; Erdemir et al., 2007] with similar associations in colon [Abbasciano et al., 1995] and lung cancer [Teramukai et al., 2009].

2.0. SUMMARY AND AIMS

2.1. SUMMARY

Local inflammation occurs as a response to injury, infection, autoimmune conditions or malignancy. Through complex pathways involving the innate, humoral and adaptive immune systems, the local immune/inflammatory reaction can result in a systemic immune/inflammatory response. As well as exhibiting clinical manifestations including pyrexia, anorexia and cachexia, the systemic response is also associated with alterations in a number of serum markers of inflammation. These include components of the innate and adaptive immune systems, for example the constituents of a differential white cell count and acute phase reactants such as C-reactive protein, as well as other biochemical parameters including albumin and liver function tests.

Worldwide, approximately 12.4 million new cases of cancer are diagnosed each year with 7.6 million people dying as a result. In the United Kingdom, around one in three people will develop cancer during their lifetime with breast, followed by lung and colorectal malignancies being the most common. Increasing age, male gender and socioeconomic deprivation have been shown to be associated with a reduced survival in the majority of malignancies. Deoxyribonucleic acid (DNA) mutations that result in the interruption of tumour suppression or the activation of pro-oncogenic pathways can occur secondary to a genetic predisposition. However, the majority of genetic mutations are acquired and it is recognised that a local inflammatory microenvironments can lead to DNA damage, neoplastic transformation and cancer development. It has also been shown that cancer progression, and subsequent metastasis, is dependent on complex interactions with the inflammatory tumour microenvironment involving the innate, humoral and adaptive immune systems. The Tumour

Node Metastasis (TNM) staging system utilises evidence of local invasion and distal tumour spread in the risk stratification of patients with cancer. It is however recognised that this system may be open to an element of subjectivity and therefore, there is increasing interest in the study of the host immune/inflammatory response in predicting outcome in patients with malignancy.

Cardiovascular and cerebrovascular atherosclerotic disease is responsible for approximately a third of all deaths in the United Kingdom. Atherosclerotic plaques have a significant inflammatory cell infiltrate and clear relationship between inflammation and vascular disease has been established. Patient risk stratification tools, such as the Framingham Score and CHADS2, are composed of a variety of risk factors including patient age, high density lipoproteins and systolic blood pressure. These scores can be used to individualise therapies in patients with cardiovascular or cerebrovascular disease and help ensure that treatment is received by those who will benefit. Similarly, C-reactive protein as a measure of the systemic immune/inflammatory response, is recommended in the assessment of patients with cardiovascular disease. This therefore raises the possibility that other, widely available serum markers of inflammation may also predict outcome.

A number of commonly sampled serum markers of the systemic immune/inflammatory response have been associated with cancer, cardiovascular, cerebrovascular and all-cause mortality. Serum C-reactive protein levels have been widely studied and shown to be predictive of outcome in a variety of cancers including colorectal, gastroesophageal and pancreatic, with similar associations demonstrated in patients with cardiovascular and cerebrovascular disease. Comparative associations with survival in similar cohorts have been reported when other serum inflammatory markers including

albumin, liver function tests and the constituents of a differential white cell count have been studied.

There has therefore been increasing interest in systemic inflammation-based scores that combine these commonly sampled markers of the systemic immune/inflammatory response and whether they can be used to predict outcome in patients with a range of diseases. A variety of scores have been suggested including the Glasgow Prognostic Score (GPS), the modified Glasgow Prognostic Score, the Neutrophil Lymphocyte Ratio (NLR), the Platelet Lymphocyte Ratio (PLR) and the Prognostic Nutritional Index (PNI). There has however been little work to investigate directly the effect that the presence of cancer has on these markers of systemic inflammation and whether their relationship with survival is a universal phenomenon in all tumour types. It is also unclear as to the optimal constituents of such an inflammation-based prognostic score and whether these would differ for cancer, atherosclerotic disease or all-cause mortality.

2.2. AIMS

To investigate the areas detailed above, in a single cohort of incidentally sampled patients, studies were carried out:

1. To explore the relationship between the presence and site of cancer, an inflammation-based prognostic score and biochemical parameters.
2. To investigate whether an inflammation-based prognostic score can predict cancer survival independent of tumour site and assess any derangement of associated biological parameters.
3. To compare the previously suggested inflammation-based prognostic scores in patients with cancer with regard to their universal utility and predictive value.

4. To investigate whether an inflammation-based score using variables available in large oncological data sets, specifically a white cell and neutrophil count, can be used to predict cancer survival.
5. To investigate the optimal constituents of an inflammation-based prognostic score for use in patients with cancer.
6. To explore whether a similar inflammation-based score can be used to predict cardiovascular, cerebrovascular and all-cause mortality.

3.0. SYSTEMIC INFLAMMATION-BASED PROGNOSTIC SCORES

3.1. ABSTRACT

Introduction: As disease associated inflammation is increasingly recognised as a key determinant of prognosis in patients with cancer, cardiovascular disease and stroke, there has been significant interest in measurements of the systemic immune/inflammatory response in an attempt to help predict patient outcome.

Methods: A systematic review was carried out looking at all published literature between September 1999 and September 2009 in the Pubmed database. The following search term were entered ("glasgow prognostic score" OR "modified glasgow prognostic score" OR "neutrophil lymphocyte ratio" OR "neutrophil to lymphocyte ratio" OR "platelet lymphocyte ratio" OR "platelet to lymphocyte ratio" OR "prognostic index" OR "prognostic nutritional index" AND (survival OR mortality)).

Results: 1095 studies were returned in the pubmed search but only 31 investigated the relationship between inflammation-based scores and mortality. The majority of these (n=19) investigated the prognostic value of the Glasgow Prognostic Score (GPS/mGPS).

Conclusion: The results indicate that good evidence exists demonstrating an association between inflammation-based scores and mortality in certain cancer types as well as cardiovascular disease. It is however unclear as to whether this is a universal phenomenon in all cancers and whether these scores could be utilised in patients with other atherosclerotic conditions such as cerebrovascular disease.

3.2. INTRODUCTION

As disease associated inflammation is increasingly recognised as a key determinant of prognosis in patients with cancer, cardiovascular disease and stroke, there has been significant interest in measurements of the systemic immune/inflammatory response in an attempt to help predict patient outcome. Furthermore, it has been suggested that an optimal biomarker should have a standardised assay, be universally available in clinical practice and financially viable for widespread use [Paoletti et al., 2004]. While single measures of commonly sampled parameters, such as an isolated C-reactive protein, white cell count or albumin, have been shown to have prognostic significance there has been a move towards combining these markers in order to create inflammation-based prognostic scores with improved sensitivity and specificity. Inflammation-based scores that quantify the systemic immune/inflammatory response associated with a disease process have the potential to provide important prognostic information and allow the risk stratification of patients. It is also recognised that the optimal constituents of a score may change depending on the clinical situation. For example, in patients entering large chemotherapeutic trials that routinely record only a differential white cell count, a score consisting of these constituents would be optimal. It is with all this in mind that the main focus of this thesis will be on systemic inflammation-based scores that utilise commonly sampled markers of inflammation including C-reactive protein, liver function tests and the constituents of a differential white cell count.

On review of the current literature, it is evident that there are a number of regularly studied inflammation-based prognostic scores that incorporate multiple, routinely available, markers of the systemic immune/inflammatory response and have been shown to be associated with mortality. The Glasgow Prognostic Score (GPS, Table 2.1), a combination of acute phase proteins namely C-reactive protein and albumin using standard thresholds (C-

reactive protein >10mg/L, albumin <35g/L) has been shown to have prognostic value in certain cancers [Forrest et al., 2003; McMillan et al., 2007]. The GPS was subsequently refined to form the modified Glasgow Prognostic Score (mGPS Table 2.1) [McMillan et al., 2007] when, in patients with primary operable colorectal cancer, hypoalbuminaemia alone was found to have similar prognostic value as a GPS of 0. Several studies have reported the mGPS to have independent prognostic value in patients with a number of different cancers in a variety of clinical scenarios [McMillan, 2009]. Other commonly sampled markers of the systemic immune/inflammatory response have also been combined to form inflammation-based prognostic scores associated with survival in patients with cancer, atherosclerotic disease and all-cause mortality (Table 3.1). These include the Neutrophil Lymphocyte Ratio (NLR), a combination of circulating neutrophil and lymphocyte counts, the Platelet Lymphocyte Ratio (PLR), a combination of circulating platelet and lymphocyte counts, the Prognostic Index (PI), a combination of C-reactive protein and white cell count and finally, Onodera's Prognostic Nutritional Index (PNI), a combination of albumin and lymphocyte count.

The aim of the present Chapter was to examine the relationship between inflammation-based prognostic scores and survival in cancer and non-cancer cohorts.

Table 3.1. Systemic Inflammation-Based Prognostic Scores.

	Score
The Glasgow Prognostic Score	
C-reactive protein ≤ 10 mg/l and albumin ≥ 35 g/l	0
C-reactive protein ≤ 10 mg/l and albumin < 35 g/l	1
C-reactive protein > 10 mg/l and albumin ≥ 35 g/l	1
C-reactive protein > 10 mg/l and albumin < 35 g/l	2
The modified Glasgow Prognostic Score	
C-reactive protein ≤ 10 mg/l and albumin ≥ 35 g/l	0
C-reactive protein ≤ 10 mg/l and albumin < 35 g/l	0
C-reactive protein > 10 mg/l and albumin ≥ 35 g/l	1
C-reactive protein > 10 mg/l and albumin < 35 g/l	2
Neutrophil Lymphocyte Ratio	
Neutrophil count:lymphocyte count $< 5:1$	0
Neutrophil count:lymphocyte count $\geq 5:1$	1
Platelet Lymphocyte Ratio	
Platelet count:lymphocyte count $< 150:1$	0
Platelet count:lymphocyte count 150–300:1	1
Platelet count:lymphocyte count $> 300:1$	2
Prognostic Nutritional Index	
Albumin (g/l) + $5 \times$ total lymphocyte count $\times 10^9/l \geq 45$	0
Albumin (g/l) + $5 \times$ total lymphocyte count $\times 10^9/l < 45$	1

3.3. MATERIALS AND METHODS

A systematic review was carried out looking at all published literature between September 1999 and September 2009 in the Pubmed database. The following search term were entered ("glasgow prognostic score" OR "modified glasgow prognostic score" OR "neutrophil lymphocyte ratio" OR "neutrophil to lymphocyte ratio" OR "platelet lymphocyte ratio" OR "platelet to lymphocyte ratio" OR "prognostic index" OR "prognostic nutritional index" AND (survival OR mortality)) and 1095 results were returned.

Abstracts were searched to identify appropriate citations with subsequent paper acquisition for those manuscripts included. Only papers in English, or those that had been translated into English were included. Only original articles were considered with all reviews being excluded. Multiple papers from the same cohort were included if the comparative findings differed. Any scores including variables other than commonly available serum markers of inflammation were excluded. Only those articles with death as a primary or secondary endpoint were considered and only those reporting disease specific or overall survival were included. Overall survival was preferentially reported, otherwise disease free survival noted. If only "survival" was mentioned then this was taken to be overall survival. Articles that grouped other events with survival (i.e. Major Adverse Cardiac Events) were disregarded. Only studies using multivariate statistical model analysis or relative survival to control for confounding factors (including age and gender) were included. As tumour stage is a major predictor of survival in patients with cancer, only papers correcting for this were included. Hazard ratios, probability values and, where available, 95% confidence intervals were included. Categorical Hazard Ratios (HR) were included where available, otherwise single continuous variables were presented. If tertile or quartile based results were presented then, where available, the HR from the first to the last were presented. HRs were shortened

to two decimal places and P-values to 3 decimal places. In patients with malignancy, where stated in the original paper, the stage of disease was reported. Where stage was not absolutely stated but could be accurately defined from the text/tables then stage is marked * to show implied. If stage could not be accurately identified then this was marked as "stage not defined." Patients with resectional/operable tumours were taken to be stage I-III and denoted with a * unless otherwise stated (excluding ovarian). Each paper's main conclusion or details of original findings were noted. If a score was independent to other factors this was noted. If other factors in a multivariate model did not retain statistical significance then a score was stated to be a superior measure.

3.4. RESULTS - Table 3.2 Systemic review of inflammation-based prognostic scores

Reference	Site	N	Score	Threshold	HR (95% CI)	P- value	Stage	Survival	Comments
Breast									
[Al Murri et al., 2006]	-	96	GPS	0/1/2	2.26 (1.45-3.52)	<0.001	IV	Cancer	GPS independent of palliative chemotherapy and radiotherapy
Gynaecological									
[Sharma et al., 2008]	Ovarian	154	GPS	0/1/2	Not Stated	<0.05	III-IV	Overall	Treated with systemic chemotherapy with or with-out surgery GPS superior to performance status
[Cho et al., 2009a]	Ovarian	192	NLR	>2.60	6.05 (1.77-20.71)	0.004	I-IV	Overall	Treated with surgery NLR Independent of cell type and grade of tumour
Gastroesophageal									
[Nozoe et al., 2002]	Oesophageal	258	PNI	>46	1.80 (1.16-2.81)	0.009	I-IV	Cancer	Treated with surgery PNI independent to lymphatic and vascular invasion
[Crumley et al., 2006b]	Gastro-oesophageal	258	GPS	0/1/2	1.51 (1.22-1.86)	<0.001	I-IV	Cancer	Treated with palliative intent GPS independent of palliative adjuvant therapy
[Han-Geurts et al., 2006]	Oesophageal	400	PNI	Per single unit increase	-	ns	I-III*	Overall	Treated with resection with or without neoadjuvant therapy PNI not independently significant

[Crumley et al., 2008]	Gastro-oesophageal	65	GPS	0/1/2	1.65 (1.10-2.47)	<0.05	III-IV	Cancer	Treated with palliative chemotherapy GPS superior to ECOG-Performance Status
[Kobayashi et al., 2008]	Oesophageal	48	GPS	1-2/0	0.17 (0.06-0.52)	0.002	I-III	Cancer	Treated with surgery and neoadjuvant chemoradiotherapy GPS independent of lymphatic and venous invasion
Renal									
[Ramsey et al., 2007]	-	119	GPS	0/1/2	2.93 (1.88-4.55)	<0.001	IV	Cancer	Treated with immunotherapy GPS independent of MSKCC and MRCCPS scores
Ramsey et al., 2008]	-	23	GPS	0/1-2	2.23 (1.09-4.57)	0.029	IV	Cancer	Treated with immunotherapy GPS superior to interleukin 6
Colorectal									
[Read et al., 2006]	Colorectal	48	mGPS	0/1/2	2.27 (1.09-4.73)	0.028	IV	Overall	Treated with palliative therapy with or without previous therapies mGPS independent of performance status, nutritional assessment tool (PGSGA) and LFTs
[McMillan et al., 2007]	Colorectal	123	mGPS	0/1/2	1.74 (1.20-2.51)	<0.001	II	Cancer	Treated with resection with or without adjuvant therapy mGPS superior to tumour site and delivery of adjuvant therapy

[Leitch et al., 2007]	Colorectal	149	mGPS	0/1/2	2.08 (1.32-3.28)	0.002	I-III	Overall	Treated with colorectal resection, or palliative procedure or chemotherapy or radiotherapy
			NLR	>5	-	ns			mGPS superior to NLR, deprivation and independent to constituents of the differential white cell count
		84	mGPS	0/1/2	1.44 (1.01-2.04)	0.043	IV	Cancer	
			NLR	>5	-	ns			
[Ishizuka et al., 2007]	Colorectal	315	GPS	1-2/0	0.17 (0.04-0.73)	0.018	I-IV	Overall	Treated only with surgery with-out adjuvant therapy GPS superior to tumour markers (CEA, CA19-9, CA72-4)
[Halazun et al., 2008]	Colorectal	440	NLR	>5	2.28 (1.65-3.13)	<0.001	IV	Overall	Treated with colonic surgery then liver surgery and adjuvant chemotherapy NLR superior to positive margin and tumour number
[Leung et al., 2008]	Colorectal	53	mGPS	0/1/2	2.12 (1.34-3.35)	0.001	IV	Cancer	Treated with chemotherapy or supportive care mGPS independent of lipid-soluble antioxidant vitamins
[Crozier et al., 2009]	Colorectal	188	mGPS	0/1/2	2.22 (1.06-4.74)	0.039	II	Cancer	Treated with surgery with or without neoadjuvant chemotherapy. mGPS superior to presentation (emergency/elective), deprivation and adjuvant therapy
[Neal et al., 2009]	Colorectal	174	NLR	>5	-	ns	IV	Overall	Treated with resection of liver metastasis with or without chemotherapy NLR not independently significant

[Kishi et al., 2009]	Colorectal	290	NLR	>5	2.00 (1.30-3.30)	0.005	IV	Overall	Colorectal with liver metastasis with chemotherapy with or without resection NLR independent to number and size of metastasis, history of previous treatment and positive margins
[Roxburgh et al., 2009a]	Colorectal	244	mGPS	0/1/2	2.34 (1.65-3.31)	<0.001	I-III	Cancer	Treated with surgery without or without adjuvant therapy mGPS independent of high risk pathological Petersen's Index
[Roxburgh et al., 2009b]	Colorectal	287	mGPS	0/1/2	2.56 (1.66-4.25)	<0.001	I-III	Cancer	Treated with surgery without or without adjuvant therapy mGPS superior to nodes sampled, adjuvant therapy and Jass criteria and independent to Klintrup criteria
[Ishizuka et al., 2009]	Colorectal	112	mGPS	0/1/2	6.07 (1.63-22.68)	0.007	II-IV	Cancer	Treated with surgery then palliative chemotherapy mGPS superior to liver function tests (AST, ALT, GGT) and tumour markers (CEA, CA19-9, CA 72-4)
[Roxburgh et al., 2009c]	colon	245	mGPS	0/1/2	1.56 (1.03-2.38)	0.038	I-III	Cancer	Treated with surgery with or without adjuvant therapy mGPS superior to adjuvant therapy and independent to Klintrup criteria
	rectum	140			1.76 (1.00-3.10)	0.033			
Hepatocellular									
[Glen et al., 2006]	Pancreatic	187	GPS	0/1/2	1.72 (1.40-2.11)	<0.001	III-IV	Overall	Treated with or without palliative bypass GPS superior to bilirubin

[Clark et al., 2007]	Pancreatic	44	NLR	>5	-	ns	NS	Overall	Treated with surgery NLR not independently significant
[Gomez et al., 2008a]	HCC	96	NLR	>5	-	ns	NS	Overall	Treated with surgery NLR not independently significant
[Smith et al., 2009]	Pancreatic	110	NLR	Per single unit increase	-	ns	NS	Overall	Treated with surgery with or without adjuvant therapy PLR superior to NLR, positive margin, differentiation and independent of lymph node ratio
			PLR	Per single unit increase	1.00 (1.00-1.01)	<0.001			
[Halazun et al., 2009]	HCC	150	NLR	>5	6.10 (2.29-16.29)	<0.001	NS	Overall	Treated with liver transplant with or without adjuvant therapy NLR superior to, tumour grade, vascular invasion and independent to AFP
Lung									
[Forrest et al., 2005]	NSCLC	101	GPS	0/1/2	2.32 (1.52-3.54)	<0.001	III-IV	Overall	Treated with or without palliative chemotherapy or radiotherapy GPS independent of treatment
[Sarraf et al., 2009]	NSCLC	177	NLR	Per single unit increase	1.10 (1.03-1.17)	0.005	I-IV	Overall	Treated with surgery NLR independent to total white cell count
[Teramukai et al., 2009]	NSCLC	388	NLR	>4.74	1.56 (1.09-2.24)	0.011	IIIb-IV	Overall	Treated with palliative chemotherapy NLR independent to smoking history and ECOG performance status

Cardiovascular								
[Duffy et al., 2006]	CVD	1046	NLR	Tertile 1 vs 3	1.60 (1.00-2.55)	0.048	Overall	Patients undergoing PCA
[Gibson et al., 2007]	CVD	1938	NLR	Per single unit increase	1.09 (1.03-1.16)	0.004	Overall	Patients undergoing elective CABG
[Nunez et al., 2008]	CVD	470	NLR	Quintile 1 vs 5	4.2 (1.72-10.21)	0.002	Overall	Patients with STEMI
[Papa et al., 2008]	CVD	422	NLR	>2.42	8.13 (1.42-46.57)	0.0186	Cardiac	Consecutive patients with stable coronary artery disease
[Tamhane et al., 2008]	CVD	2833	NLR	Tertile 1 vs 3	3.88 (3.19-8.06)	<0.001	Overall	Patients admitted with ACS
[Poludasu et al., 2009]	CVD	372	NLR	>3.5	2.11 (1.11-4.02)	0.023	Overall	Patients treated with PCI

3.5. DISCUSSION

The results of the current chapter show that the presence of a systemic immune/inflammatory response, as evidenced by a number of inflammation-based scores, can predict mortality in both cancer and cardiovascular disease. A number of different scores, that all incorporate multiple, routinely available, markers of the systemic immune/inflammatory response have been suggested for use. These include the Glasgow Prognostic Score (GPS and mGPS), Neutrophil Lymphocyte ratio (NLR), Platelet Lymphocyte Ratio (PLR) and Onodera's Prognostic Nutritional Index (PNI).

The majority of studies (n=31) to date (September 2009) have investigated the relationship between inflammation-based scores and cancer. Over half of these (n=18) include patients with malignancies of the alimentary tract including colorectal and gastroesophageal cancer. However, there is also good evidence that inflammation-based scores have a predictive role in pancreatic, hepatobiliary, renal and ovarian cancers. Of these scores the Glasgow Prognostic Score (GPS/mGPS) is the most widely validated with 19 studies all showing a statistically significant association with mortality. The NLR has also been extensively investigated in a variety of tumour types with varying associations with mortality demonstrated. The PLR and PNI have only been included in one study each but, never the less, have also been reported to be associated with mortality. Indeed, it is likely that, while some scores may be more sensitive than others, they are essentially measuring the same process. There have been six studies looking at cardiovascular disease, all of which have demonstrated a positive correlation between the NLR and mortality.

This chapter has detailed evidence demonstrating that inflammation-based scores have predictive value in certain cancer types as well as cardiovascular disease. It is however unclear whether this is a universal phenomenon in all cancers and if these scores could be

utilised in patients with other atherosclerotic conditions such as cerebrovascular disease. The optimal constituents of an inflammation-based score for use in cancer and non-cancer cohorts is yet to be determined.

4.0. THE RELATIONSHIP BETWEEN THE PRESENCE AND SITE OF CANCER, AN INFLAMMATION-BASED PROGNOSTIC SCORE AND BIOCHEMICAL PARAMETERS

4.1. ABSTRACT

Introduction: Biochemical parameters, such as C-reactive protein and liver function tests, have been associated with mortality. The aim of the present chapter was to examine the relationship between the presence and site of cancer, an inflammation-based prognostic score (mGPS) and biochemical parameters.

Methods: Patients (n=223,303) who had a single incidental sample taken for C-reactive protein, albumin, calcium and serum liver function tests where available, between 2000 and 2008 were studied. Those with a pathological diagnosis of cancer (n=22,715) were also identified.

Results: The strongest associations (Spearman's correlation ≥ 0.3) in both the non-cancer and cancer groups were found between albumin, C-reactive protein and Alk phos, AST and ALT, AST and GGT and ALT and GGT (all $p < 0.001$). On multivariate analysis, age, deprivation, C-reactive protein, albumin, adjusted calcium, Alk phos and GGT were associated with the presence of cancer (all $p < 0.01$). When cancer diagnoses were ranked from those with the lowest proportion of mGPS 1 or 2 to those with the highest, the percentage of cases with a mGPS of 1 or 2 ranged from 21% in breast cancer to 68% in pulmonary cancer.

Conclusion: The results indicate that the systemic inflammatory response is increased by the presence of cancer. There is a striking parallel between the proportions of cases with a mGPS of 1 or 2 and reported survival rates in these tumours.

4.2. INTRODUCTION

In the West of Scotland there has been a longstanding interest in the role of the systemic immune/inflammatory response in determining outcome in cancer [McMillan, 2008; McMillan, 2009; Roxburgh and McMillan, 2010]. Although, as detailed in Chapter 1, it is recognised that the development of cancer has a genetic basis, there is increasing evidence that host immune/inflammatory responses play a pivotal role in the development and progression of cancer [Balkwill and Mantovani, 2001; Coussens and Werb, 2002; Mantovani et al., 2008; Colotta et al., 2009; McDonald et al., 2009]. This is consistent with the hypothesis that chronic exposure to inflammatory processes, either through infection, social conditions or lifestyle, enhances the ageing process [Finch and Crimmins, 2004].

As has also been detailed in Chapter 1, the presence of a systemic immune/inflammatory response as evidenced by an elevated C-reactive protein concentration sampled incidentally in a large hospital based cohort, was associated with a shorter duration of cancer and non-cancer survival [Marsik et al., 2008]. Goldwasser and Feldman, amongst others, have reported a similar relationship in a systematic review of the prognostic value of albumin [Goldwasser and Feldman, 1997]. These acute phase proteins, produced exclusively in the liver, have been shown to be a major factor in the progressive nutritional and functional decline of patients with cancer [McMillan, 2008]. Also, C-reactive protein and albumin concentrations have been combined to form the modified Glasgow Prognostic Score (mGPS) and have been reported to be independently associated with reduced survival in patients with a variety of cancers as detailed in Table 3.2 [McMillan, 2009; Roxburgh and McMillan, 2010].

It is also of interest that liver function tests such as bilirubin [Temme et al., 2001], alkaline phosphatase [Tonelli et al., 2009] and γ -glutamyl transferase [Kazemi-Shirazi et al.,

2007] as well as serum calcium [Leifsson and Ahren, 1996], have also been reported to predict cancer and non-cancer outcome (Chapter 1.). There is also some evidence that liver function tests are altered as part of the systemic immune/inflammatory response in patients with lung and gastrointestinal cancer [Brown et al., 2007; Roxburgh et al., 2009a].

The aim of the present Chapter was to examine the effect of cancer on markers of systemic inflammation induced by the liver (mGPS) and on these variables (bilirubin, alkaline phosphatase, aspartate transaminase, alanine transaminase, γ -glutamyl transferase and adjusted calcium).

4.3. MATERIALS AND METHODS

This cohort includes patients who have had a single sample taken for C-reactive protein, albumin and calcium between the 1st of January 2000 and the 1st of November 2008 in the North of Glasgow. The samples were taken incidentally and if more than one set of measurements were available for a given patient, only the initial set was used. Where available, serum liver function tests, including bilirubin and alkaline phosphatase (Alk phos) were recorded. Patients were excluded if they did not have a complete set of identifying details (name, gender, date of birth and hospital number). The limit of detection of C-reactive protein was a concentration of less than 5mg/L. The mGPS was constructed, using C-reactive protein and albumin, as follows; patients with both an elevated C-reactive protein (>10 mg/L) and low albumin (<35 g/L) were allocated a score of 2; patients in whom only C-reactive protein was elevated (>10 mg/L) were allocated a score of 1 and those with a normal C-reactive protein were allocated a score of 0 [McMillan et al., 2007]. The rationale and basis of the mGPS has been previously described [McMillan, 2008]. Serum C-reactive protein, albumin and liver function tests, including bilirubin, Alk phos, AST, ALT and GGT, as well as calcium adjusted for albumin [Ashby et al., 1986], were classified in accordance

with the NHS Greater Glasgow and Clyde biochemistry laboratory reference ranges. These analytes were measured using identical standard operating procedures on identical automated platforms using reagents from the same manufacturer (Abbot Diagnostics). Analysis was carried out in three linked laboratories in the North of Glasgow which take part in external quality assurance schemes for routine biochemical analysis. All samples were refrigerated, as stable for a number of days in these conditions, and analysed within 6 hours.

All patient records fulfilling these criteria were then cross referenced with the local pathology database to identify those in whom a tissue diagnosis of cancer had been made either before or after their initial blood sample (Appendix 1). Systematized Nomenclature of Medicine – Clinical Terms (SNOMED CT) codes, which share certain codes with the International Classification of Diseases, contained in the pathology database were used to identify the site and type of tumour in those with a tissue diagnosis of cancer. Patients were then grouped into those with a pathological diagnosis of cancer and those without. The cancer group were then grouped into those patients with a clear primary cancer diagnosis, those with multiple tumours or those with no clear indication of the primary tumour site. Patients with incomplete identifying details (name, date of birth and hospital number), under 16 years of age, or those with multiple malignancies, metastatic disease or cancer of an unknown origin were excluded. For the purpose of analysis, cancers were grouped according to tumour site; bladder, breast, colorectal, dermatological, endocrinological, gastroesophageal, gynaecological, haematological, head and neck, hepatocellular, musculoskeletal and soft tissue, pancreaticobiliary, pulmonary, prostatic, renal and testicular. The study was approved by the Research Ethics Committee, North Glasgow NHS Trust.

The Scottish Index of Multiple Deprivation (SIMD) 2006, as detailed in Chapter 1. and recommended by the ISD on behalf of NHS Scotland and the Scottish Government

Department of Health, was used to measure deprivation with the least deprived being scored as 1 to the most deprived scoring 5 [Bishop J et al., 2004].

SNOMED CT codes were used to identify the site and type of tumour in those with a tissue diagnosis of cancer. Each pathology sample had two SNOMED CT codes, one for type of tumour (morphology) and one for site of tumour (topography). Examples of SNOMED CT cancer morphology codes, of which there are over 400, include metastatic carcinoma (M80106), large cell carcinoma (M80123), small cell carcinoma (M80413-M80416), squamous cell carcinoma (M80703-M80763), adenocarcinoma (M81402-M81406), cholangiocarcinoma (M81603), renal cell carcinoma (M83123-M83126), intraductal carcinoma (M85002-M85003), malignant melanoma (M87202-M87433), mesothelioma (M90501-M90503), lymphoma (M95903-M97003) and leukaemia (M98003-M99403). Examples of SNOMED CT topography codes include bladder (T74000-T74400), breast (T04000-T04400), colorectal (colon T67000- T67995, rectum T68000-T68200, anus T69000-T69200) dermatological (T02100-02870), endocrine (adrenal T93000-T93100, thyroid T96000-T97800), gastroesophageal (oesophagus T62000-T62910, stomach T63000-T63700), gynaecological (uterus with or without cervix T82000- T82900, uterus and ovaries T82920-T82922, cervix T83000-T83300, endometrium T84000-T85000, ovary T86920-T87800), haematological (bone marrow: T06000-T06600, lymph node T08000-T09600), head and neck (pharynx T60000-T61300, upper respiratory tract T21000-T24920, mouth and salivary glands T51000-T55550), liver (T56000-T56020), musculoskeletal and soft tissue (skeletal bone T10000-T12720, skeletal muscle T13000) pancreaticobiliary (gallbladder T57000, bile duct T57600-T58500, pancreas and ampulla T58700-T59300) pulmonary(T26000-T29900), prostatic (T77000-T77350), renal (T71000-T72000) testicular (T78000-T78020). The combination of SNOMED CT morphology and topography codes were used to identify the tumour type and site.

With regards to statistical analysis, the relationships between patient demographics, the presence of cancer, biochemical parameters, the mGPS, tumour site and other biochemical parameters were analysed using the Pearson's chi-square test (linear by linear association). Spearman's rank correlation was used to measure the strength of the relationship between age and biochemical parameters and the interrelationships between biochemical parameters. The relationship between the presence of cancer and patient demographics and biochemical parameters was examined using Cox proportional hazards model multivariate regression analysis; only patients who had a diagnosis of cancer made within two years following their blood test were included in this analysis. This was based on the premise that this group of patients were likely to have had an ongoing malignant process. Analysis was performed using SPSS software (SPSS Inc., Chicago, IL, USA).

4.4. RESULTS

In total 223,303 patients were studied. The majority, 144,900 (65%), were under 65 years of age. There were 120,115 (54%) females and 103,188 (46%) males. Of those patients with an identifiable postcode (86%), the majority were from the Greater Glasgow area (88%). In this cohort, when measured by the SIMD 2006, 14% of cases were from affluent areas (least deprived quintile of the Scottish population) and 45% from deprived areas (most deprived quintile of the Scottish population).

The demographics of non-cancer and cancer patients are shown in Table 4.1. Patients with cancer, when compared to those without a diagnosis of cancer were older and more likely to be female ($p < 0.001$). The cancer group had a higher proportion of cases from less deprived areas than those in the non-cancer group ($p < 0.001$).

Of 22,715 with a diagnoses of cancer 19,476 (86%) had a single primary tumour while 1299 (6%) had multiple malignancies. In 1940 (8%) cases SNOMED CT codes gave

no definitive indication of the primary site of malignancy. Examples of this include: 1497 cases were SNOMED CT codes stated lung carcinoma, lung adenocarcinoma or lung metastatic carcinoma; 269 cases were SNOMED CT codes stated liver carcinoma, liver adenocarcinoma or liver metastatic carcinoma; 209 cases had SNOMED CT codes stating lymph node carcinoma, lymph node adenocarcinoma or lymph node metastatic carcinoma and 160 cases had SNOMED CT codes stating metastatic carcinoma at other sites.

When the cancer group was categorised (n=22,715) into different tumour sites the following main groups were observed; bladder (n=939, 4%), breast (n=3849, 17%), colorectal (n=1899, 8%), dermatological (n=4907, 22%), endocrinological (n=303, 1%), gastroesophageal (n=1018, 5%), gynaecological (n=1136, 5%), haematological (n=1165, 5%), head and neck (n=610, 3%), hepatocellular (n=55, 0%), musculoskeletal and soft tissue (n=164, 1%), pancreaticobiliary (n=456, 2%), prostate (n=957, 4%), pulmonary (n=1393, 6%), renal (n=527, 2%) and testicular (n=97, 1%).

The relationship between markers of the systemic immune/inflammatory response, biochemical parameters and the presence of cancer is shown in Table 4.2. There were higher circulating concentrations of C-reactive protein and lower albumin levels (and thus a higher proportion of mGPS 1 or 2), in those patients with cancer (all $p < 0.001$). Patients in the cancer group were also found to have lower AST and ALT levels but higher adjusted calcium, Alk phos and GGT levels compared with the non-cancer cohort (all $p < 0.001$).

The interrelationships between markers of the systemic immune/inflammatory response and biochemical parameters in patients without and with cancer are shown in Tables 4.3 and 4.4. In the non-cancer cohort age was significantly associated with all biochemical parameters (all $p < 0.001$) (Table 4.3). Similarly, all biochemical parameters were significantly associated with each other (all $p < 0.001$). In the non-cancer cohort the strongest

associations (Spearman's correlation > 0.3) were between C-reactive protein and albumin, C-reactive protein and Alk phos, AST and ALT, AST and GGT and ALT and GGT.

There were similar interrelationships in the cancer cohort (Table 4.4) with the strongest associations (Spearman's correlation > 0.3) existing between C-reactive protein and albumin, C-reactive protein and Alk phos, bilirubin and AST, Alk phos and GGT, AST and ALT, AST and GGT and ALT and GGT.

The temporal relationships between the date of blood samples and cancer diagnoses are shown in Table 4.5. In order to give a clearer indication of the alterations that occur in biochemical parameters around the diagnosis of cancer only patients with a diagnosis within two years of their blood sample were included in the analysis, pre-diagnosis (n=8083) and post-diagnosis (n=5971). Compared with those patients who had pre-diagnosis levels, post-diagnosis patients were more likely to be female and affluent (all $p < 0.001$). Patients following a diagnosis of cancer had lower albumin levels and thus higher mGPS ($p < 0.001$). Also, post diagnosis patients were more likely to have lower adjusted calcium, bilirubin, Alk Phos, AST, ALT and GGT levels (all $p < 0.05$).

The relationship between patient demographics, biochemical parameters and the absence (n= 200,588) or presence (n= 8083) of cancer within two years pre-diagnosis group is shown in Table 4.6. On univariate analysis, those over 65 years of age and living in least deprived areas were more likely to have cancer (all $p < 0.001$). Patients with an elevated C-reactive protein, adjusted calcium, Alk phos and GGT levels or low albumin levels were more likely to have cancer (all $p < 0.001$). On multivariate analysis, these associations with the presence of cancer persisted in age, deprivation, C-reactive protein, albumin, adjusted calcium, Alk phos and GGT (all $p < 0.01$).

The relationships between the mGPS and tumour site in the two year pre-diagnosis group is shown in cancers with more than 50 cases in Table 4.7. The cancers are ranked from those with the highest proportion of mGPS 0 to those with the lowest. The percentage of cases with a mGPS of 1 or 2 ranges from 21% in breast cancer, to 46% in prostate cancer and to 68% in pulmonary cancer. Compared with breast cancer the mGPS was significantly higher in dermatological, bladder, endocrinological, gynaecological, prostate, musculoskeletal, gastroesophageal, haematological, renal, colorectal, head and neck, pancreaticobiliary and pulmonary cancers (all $p < 0.001$).

4.5. DISCUSSION

The results of the current Chapter show that the presence of a systemic immune/inflammatory response, as evidenced by an elevated C-reactive protein concentration ($>10\text{mg/l}$), was present in 40% and hypoalbuminaemia ($<35\text{g/l}$) was present in 14% of 223 303 patients who were incidentally sampled. Adjusted calcium was elevated in 2.2% (1.8% unadjusted), Alk phos was elevated in 12%, ALT was elevated in 11% and GGT was elevated in 26% of patients studied.

These results are consistent with previous studies including an Austrian cohort of approximately 280,000 patients, in whom 35% had an elevated C-reactive protein concentration [Marsik et al., 2008] and 13% had hypoalbuminaemia [Grimm et al., 2009]. They also reported that 33% had an elevated GGT concentration [Kazemi-Shirazi et al., 2007]. In a Swedish population based cohort of 33,346 patients, 1% were found to have an elevated unadjusted serum calcium [Leifsson and Ahren, 1996]. In a North American cohort of 18,835 patients approximately 9% had an elevated Alk phos [Tonelli et al., 2009]. Another population based cohort from the United States of 14,950 patients reported an

elevated ALT level in 14% of patients [Ruhl and Everhart, 2009]. Therefore, the results of the present Chapter are similar to those previously reported in other large cohort studies.

In the present cohort there were more females in the cohort as a whole, more females in the cancer group and more females had measurements taken post cancer diagnosis. However, gender was not significantly associated with the presence of cancer. The present Chapter shows that when compared with non-cancer, the presence of cancer was associated with a significantly higher proportion of cases with an elevated C-reactive protein (>10mg/L, 47% vs 39%) and lower albumin (<35g/L, 19% vs 14%) concentrations, higher adjusted calcium (>2.60mmol/L, 4% vs 2%), Alk phos (>280U/L, 16% vs 12%) and GGT (>40U/L in females, >70U/L in males, 29% vs 25%), and lower ALT levels (>50U/L, 10% vs 12%). Furthermore, when samples taken following a diagnosis of cancer were compared with those taken prior to a diagnosis of cancer the proportion of cases with albumin (>35g/L, 74% vs 81%) adjusted calcium (>2.60mmol/L, 3% vs 5%), Alk phos (>280U/L, 15% vs 18%), ALT (>50U/L, 10% vs 11%). and GGT (>40U/L, in females, >70U/L in males 29% vs 31%) levels were all lower. When these biochemical parameters together with bilirubin, and ALT were regressed against the presence of cancer, C-reactive protein, albumin, adjusted calcium, Alk phos and GGT were all shown to be independently associated with the presence of cancer. These results would suggest that the presence of cancer influences a number of biochemical parameters that have been previously associated with all-cause mortality [Leifsson and Ahren, 1996; Kazemi-Shirazi et al., 2007; Marsik et al., 2008; Grimm et al., 2009; Tonelli et al., 2009].

The present Chapter also showed that all these biochemical parameters were significantly associated. In particular, C-reactive protein was associated with albumin and Alk phos and GGT associated with AST and ALT in both cancer and non-cancer cohorts. Recently the combination of C-reactive protein and albumin, termed the mGPS, was shown

to be associated in a similar manner in patients with lung cancer [Brown et al., 2007] and predict cancer specific survival in a number of operable [Roxburgh and McMillan, 2010] and inoperable cancers [McMillan, 2009]. These results raise the question of whether the above reported association of these biochemical parameters and all-cause mortality are indeed independent of each of other. Further studies using this cohort will aim to address this question.

In the present Chapter the proportions of mGPS 1 or 2 were greater following a diagnosis of cancer. This appears to be secondary to a reduction in albumin which may be secondary to a number of factors including operative management, adjuvant oncological therapies or tumour progression, a pattern recognised in previous studies within specific tumour types [McMillan et al., 2003; Jamieson et al., 2005]. Interestingly in patients who had blood tests taken following a diagnosis of cancer, adjusted calcium, bilirubin and Alk phos, AST, ALT and GGT levels were all more likely to be within normal limits. The reasons for this are unclear but it is possible that removal or treatment of the underlying malignant process in a proportion of these patients may be responsible.

In the present Chapter it was also of interest to note that, prior to diagnosis, there were significant variations in the proportions of cases with a mGPS of 1 or 2. Indeed, the proportion of patients with an elevated mGPS varied from 21% in breast cancer, to 46% in prostate cancer and to 68% in pulmonary cancer. There is a striking parallel between the proportions of cases with a mGPS of 1 or 2 and reported survival rates in these tumours as detailed in Chapter 1 [Cancer Research UK, 2007]. For example, the five year survival rate for breast cancer is approximately 84% and the proportion of patients in the present Chapter with a mGPS of 0 was 79%. In contrast the 5 year survival for lung cancer is 8% and the proportion of patients in the present Chapter with a mGPS of 0 was 32%. These results may suggest that, in addition to the mGPS having prognostic value within tumour types

[McMillan, 2009] it may also have prognostic value across tumour types. Further work is required to confirm this hypothesis.

The present cohort has a number of limitations. The patients were selected on the basis that measurements of C-reactive protein, albumin and calcium had been performed and were therefore not necessarily representative of all non-cancer and cancer patients treated in the North Glasgow area and therefore gives no meaningful information on cancer prevalence. Similarly, as this cohort was not a true representation of the North of Glasgow population, a direct comparison between SIMD measured deprivation and previously reported cancer incidence (figure 1.7) was not possible. Furthermore, sampling was incidental and not performed at a standard time during the course of the disease in the cancer group. It is also recognised that patients in both groups may have concurrent morbidity causing a rise in their C-reactive protein and derangement of their albumin and other biochemical parameters. As inclusion in the cancer cohort was dependent on a pathological diagnosis of cancer it is acknowledged that a small number in the non-cancer group may have had a clinical diagnosis of malignancy. In order to provide more detailed follow up with regards to survival it is important to link the present data with patient outcomes.

In summary, the results of the present Chapter indicate that the systemic immune/inflammatory response is common in a large patient cohort, increased by the presence of cancer and associated with the perturbation of a number of biochemical parameters previously reported to be associated with mortality. Taken together with the myriad effects of the systemic immune/inflammatory response on host metabolism [Gabay and Kushner, 1999] these results have a number of clinical implications particularly for clinical epidemiological studies. Future studies should incorporate a measure of the systemic immune/inflammatory response, such as C-reactive protein or the mGPS, in the study design,

analysis and interpretation. This is required to account for the likely confounding effect of this response on the other measured variables and their relationship with outcomes.

Table 4.1. The relationship between patient demographics and the presence of cancer

		Non-Cancer	Cancer	p value
n=223,303		n (%)	n (%)	
Age	<65 years	135,531 (67)	9369 (41)	<0.001
	65-74 years	31,593 (16)	6412 (28)	
	>75 years	33,464 (17)	6934 (31)	
Sex	Male	93,313 (46)	9875 (43)	<0.001
	Female	107,275 (54)	12,840 (57)	
SIMD 2006	1 (least deprived)	22,902 (14)	3233 (15)	<0.001
	2	18,693 (11)	2578 (12)	
	3	21,148 (12)	2771 (13)	
	4	30,850 (18)	3872 (18)	
	5 (most deprived)	77,448 (45)	8844 (42)	

Table 4.2. The relationship between markers of the systemic immune/inflammatory response, biochemical parameters and cancer

		Non-Cancer	Cancer	p value
n=223 303		n (%)	n (%)	
		200,588	22,715	
C-reactive protein	≤10mg/l	122,317 (61)	12,050 (53)	<0.001
	>10mg/l	78,271 (39)	10,665 (47)	
Albumin	<35g/l	27,515 (14)	4275 (19)	<0.001
	≥35g/l	173,073 (86)	18,440 (81)	
mGPS	0	122,317 (61)	12,050 (53)	<0.001
	1	56,357 (28)	7033 (31)	
	2	21,914 (11)	3632 (16)	
Adjusted calcium	<2.10mmol/l	4963 (3)	577 (2)	<0.001
	2.10-2.60mmol/l	191,429 (95)	21,323 (94)	
	>2.60mmol/l	4196 (2)	815 (4)	
Bilirubin	<20μmol/l	170,447 (88)	19,322 (88)	0.008
	≥20μmol/l	24,258 (12)	2593 (12)	
Alkaline phosphatase	<80U/l	28,168 (14)	2463 (11)	<0.001
	80-280U/l	148,245 (74)	16,626 (73)	
	>280U/l	23,954 (12)	3603 (16)	
Aspartate transaminase	<40U/l	162,269 (84)	18,676 (86)	<0.001
	≥40U/l	30,531 (16)	3129 (14)	
Alanine transaminase	<50U/l	144,468 (88)	15,872 (90)	<0.001
	≥50U/l	19,043 (12)	1740 (10)	
γ-glutamyl transferase	M <70 U/l, F <40U/l	145,658 (75)	15,689 (71)	<0.001
	M ≥70U/l, F ≥40U/l	49,393 (25)	6325 (29)	

Table 4.3. The interrelationships between markers of the immune/inflammatory response and biochemical parameters in patients without cancer

n=200,588	C-reactive protein	Albumin	Adjusted Calcium	Bilirubin	Alkaline phosphatase	Aspartate transaminase	Alanine transaminase	γ-glutamyl transferase
Age	0.199***	-0.286***	0.052***	0.099***	0.177***	0.078***	-0.047***	0.148***
C-reactive protein		-0.362***	-0.108***	0.167***	0.320***	0.103***	0.040***	0.193***
Albumin			-0.060***	0.040***	0.022***	0.009***	0.072***	-0.116***
Adjusted calcium				-0.077***	-0.147***	-0.081***	-0.028***	0.077***
Bilirubin					0.143***	0.297***	0.216***	0.169***
Alkaline phosphatase						0.225***	0.152***	0.243***
Aspartate transaminase							0.736***	0.437***
Alanine transaminase								0.550***

***p<0.001

Table 4.4. The interrelationships between markers of the immune/inflammatory response and biochemical parameters in patients with cancer

n=22,715	C-reactive protein	Albumin	Adjusted Calcium	Bilirubin	Alkaline phosphatase	Aspartate transaminase	Alanine transaminase	γ-glutamyl transferase
Age	0.103***	-0.203***	0.024***	0.127***	0.079***	0.020**	-0.162***	-0.062***
C-reactive protein		-0.439***	-0.078***	0.166***	0.308***	0.104***	0.034***	0.239***
Albumin			-0.086***	0.004	0.028***	0.007	0.073***	-0.123***
Adjusted calcium				-0.095***	-0.087***	-0.053***	-0.013	0.106***
Bilirubin					0.166***	0.307***	0.225***	0.177***
Alkaline phosphatase						0.253***	0.213***	0.346***
Aspartate transaminase							0.720***	0.403***
Alanine transaminase								0.505***

***p<0.001

Table 4.5. The relationship between time of diagnosis and biochemical parameters in patients with cancer

n=14,054		Pre-diagnosis n (%)	Post-diagnosis n (%)	p value
Age	<65 years	3516 (43)	2690 (45)	0.732
	65-74 years	2413 (30)	1626 (27)	
	>75 years	2154 (27)	1655 (28)	
Sex	Male	3806 (47)	2482 (42)	<0.001
	Female	4277 (53)	3489 (58)	
SIMD 2006	1 (least deprived)	1140 (15)	951 (17)	<0.001
	2	958 (12)	724 (13)	
	3	1080 (14)	802 (15)	
	4	1428 (19)	1003 (18)	
	5 (most deprived)	3042 (40)	2011 (37)	
C-reactive protein	<10mg/l	4082 (51)	2963 (50)	0.304
	>10mg/l	4001 (49)	3008 (50)	
Albumin	<35g/l	1566 (19)	1545 (26)	<0.001
	>35g/l	6517 (81)	4426 (74)	
mGPS	0	4082 (50)	2963 (50)	<0.001
	1	2666 (33)	1643 (27)	
	2	1335 (17)	1365 (23)	
Adjusted Calcium	<2.10mmol/l	183 (2)	151 (3)	<0.001
	2.10-2.60	7501 (93)	5614 (94)	
	>2.60mmol/l	399 (5)	206 (3)	
Bilirubin	<20µmol/l	6844 (87)	5163 (90)	<0.001
	≥20µmol/l	994 (13)	587 (10)	
Alkaline phosphatase	<80U/l	788 (10)	743 (12)	0.003
	80-280U/l	5805 (72)	4324 (73)	
	>280U/l	1482 (18)	898 (15)	
Aspartate transaminase	<40U/l	6614 (85)	4941 (86)	0.030
	≥40U/l	1175 (15)	788 (14)	
Alanine transaminase	<50U/l	5729 (89)	4215 (90)	0.004
	≥50U/l	734 (11)	450 (10)	
γ-glutamyl transferase	M<70 U/l, F<40U/l	5440 (69)	4100 (71)	0.012
	M≥70U/l, F≥40U/l	2432 (31)	1667 (29)	

Table 4.6. The relationship between patient demographics, biochemical parameters and the presence of cancer

n=208,671		Univariate analysis			Multivariate analysis		
		Odds Ratio	95% CI	p-value	Odds Ratio	95% CI	p-value
Age	<65/	1		<0.001	1		<0.001
	65-74	2.94	2.79-3.12	<0.001	2.60	2.46-2.75	<0.001
	>75	2.48	2.35-2.62	<0.001	2.17	2.05-2.30	<0.001
Sex	Male	1		0.317			
	Female	0.98	0.94-1.02	0.317			
SIMD	5 (most deprived)	1		<0.001	1		<0.001
	4	1.18	1.11-1.26	<0.001	1.16	1.09-1.24	<0.001
	3	1.30	1.21-1.40	<0.001	1.32	1.22-1.42	<0.001
	2	1.31	1.21-1.41	<0.001	1.36	1.26-1.46	<0.001
	1 (least deprived)	1.27	1.18-1.36	<0.001	1.27	1.18-1.36	<0.001
C-reactive protein	<10mg/l	1		<0.001	1		<0.001
	≥10mg/l	1.53	1.47-1.60	<0.001	1.22	1.16-1.28	<0.001
Albumin	<35g/l	1.51	1.43-1.60	<0.001	1.11	1.04-1.19	0.002
	≥35g/l	1		<0.001	1		0.002
Adjusted calcium	<2.10mmol/l	0.94	0.81-1.09	0.425			
	2.10-2.60mmol/l	1		<0.001			<0.001
	>2.60mmol/l	2.43	2.18-2.70	<0.001	1.90	1.70-2.21	<0.001
Bilirubin	<20µmol/l	1		0.558			
	≥20µmol/l	1.02	0.95-1.09	0.558			
Alkaline phosphatase	<80U/l	0.71	0.66-0.77	<0.001	0.81	0.75-0.87	<0.001
	80-280U/l	1		<0.001	1		<0.001
	>280U/l	1.58	1.49-1.68	<0.001	1.30	1.22-1.39	<0.001
Aspartate transaminase	<40U/l	1		0.075			
	≥40U/l	0.94	0.89-1.01	0.075			
Alanine transaminase	<50U/l	1		0.477			
	≥50U/l	0.97	0.90-1.05	0.477			
γ-glutamyl transferase	M<70 U/l,F<40U/l	1		<0.001	1		
	M≥70U/l, F≥40U/l	1.32	1.26-1.38	<0.001	1.09	1.03-1.15	0.003

Table 4.7. The relationship between tumour site and an inflammation-based prognostic score sampled prior to diagnosis

Tumour Site	Total n	mGPS n (%)			p value*
		0	1	2	
Breast	1199	950 (79)	207 (17)	42 (4)	
Dermatological	1126	693 (62)	310 (27)	123 (11)	<0.001
Bladder	296	174 (59)	76 (26)	46 (15)	<0.001
Endocrinological	145	84 (58)	41 (28)	20 (14)	<0.001
Gynaecological	197	108 (55)	60 (30)	29 (15)	<0.001
Prostate	267	145 (54)	91 (34)	31 (12)	<0.001
Musculoskeletal	78	41 (53)	15 (19)	22 (28)	<0.001
Gastroesophageal	503	249 (50)	162 (32)	92 (18)	<0.001
Haematological	499	245 (49)	157 (32)	97 (19)	<0.001
Renal	294	132 (45)	105 (36)	57 (19)	<0.001
Colorectal	784	342 (44)	250 (32)	192 (24)	<0.001
Head and neck	156	67 (43)	58 (37)	31 (20)	<0.001
Pancreaticobiliary	321	122 (38)	101 (31)	98 (31)	<0.001
Pulmonary	820	260 (32)	387 (47)	173 (21)	<0.001

*compared with breast cancer

5.0. AN INFLAMMATION-BASED PROGNOSTIC SCORE (mGPS) PREDICTS CANCER SURVIVAL INDEPENDENT OF TUMOUR SITE

5.1. ABSTRACT

Introduction: The modified Glasgow Prognostic score (a combination of C-reactive protein and albumin) as well as liver function tests including bilirubin, alkaline phosphatase and γ -glutamyl transferase have been reported to predict cancer survival. The aim of the present study was to examine the relationship between the mGPS, biochemical parameters and survival.

Methods: Patients (n=21,669) sampled between 2000 and 2006 for C-reactive protein, albumin, liver function tests where available, as well as a diagnosis of cancer were identified. Of this group 9608 patients who had an ongoing malignant process were studied (sampled within two years following diagnosis).

Results: On follow up, there were 6005 (63%) deaths of which 5122 (53%) were cancer deaths. The median time from blood sampling to diagnosis was 1.4 months. Increasing age, male gender, and increasing deprivation was associated with reduced 5 year overall and cancer specific survival (all $p < 0.001$). An elevated mGPS, adjusted calcium, bilirubin, Alk phos, AST, ALT and GGT were associated with a reduced 5 year overall and cancer specific survival, independent of age, sex and deprivation (all $p < 0.001$). An increasing mGPS was predictive of a reduced cancer specific survival in all cancers (all $p < 0.001$).

Conclusion: The results of the present study indicate that the mGPS is a powerful prognostic factor when compared with other biochemical parameters and independent of tumour site in patients with cancer.

5.2. INTRODUCTION

As detailed in Chapter 3, measurement of the systemic immune/inflammatory response has been refined using a selective combination of C-reactive protein and albumin (termed the modified Glasgow Prognostic score, mGPS) and has been shown to have prognostic value, independent of tumour stage, in lung, gastrointestinal and renal cancers [McMillan, 2008; McMillan, 2009]. As has been detailed in Chapter 1, it is also of interest that liver function tests such as bilirubin [Temme et al., 2001], alkaline phosphatase [Tonelli et al., 2009] and γ -glutamyl transferase [Kazemi-Shirazi et al., 2007] as well as serum calcium [Leifsson and Ahren, 1996], have also been reported to predict cancer and non-cancer mortality in large cohort studies.

In Chapter 4 it has been demonstrated, in a large cohort of more than 200,000 patients, that the mGPS is elevated in patients with cancer when compared to those without cancer [Proctor et al., 2010]. Moreover, there were significant interrelationships between the above biochemical parameters, including the constituents of the mGPS [Proctor et al., 2010]. The question is therefore raised whether the mGPS and other routine biochemical parameters have independent prognostic value in patients with cancer and whether this applies across different tumour sites.

The aim of the present Chapter was to examine the relationship between an inflammation based prognostic score (mGPS), biochemical parameters, tumour site and survival in patients with cancer.

5.3. MATERIALS AND METHODS

From a cohort previously described in Chapter 4 [Proctor et al., 2010], patients in North Glasgow who were sampled incidentally for the pre-mentioned markers of the systemic immune/inflammatory response as well as liver function tests between the 1st January 2000 and the 31st December 2006 to correspond with Cancer Registry follow-up data, were identified.

Only patients with blood samples taken within two years following their cancer diagnosis were included. This was done with the premise that a prognostic score would be of clinical use during this time period. Biochemical variable thresholds were categorised as detailed in Chapter 4 [Proctor et al., 2010].

Cancer diagnosis was established through linkage with the Scottish Cancer Registry using exact matches of patients' forename, surname and date of birth followed by a Soundex phonetic matching algorithm if initial exact matching was unsuccessful (Appendix 2). At the time of data collection the Scottish Cancer Registry held complete pathological and clinical cancer diagnosis records from the 1st of January 1980 until 31st December 2006 and mortality follow-up until 30th June 2009. In those who had died, cancer specific deaths were classified as patients whose primary cause of death matched their primary cancer diagnosis. All other deaths were classed as non-cancer specific deaths. Only patients who had complete Cancer Registry follow-up were included. Patient numbers in this Chapter differ from those in Chapter 4 as the matching process was through the cancer registry and not the North of Glasgow pathology database.

Cancers were coded in accordance with the International Classification of Disease 10 and broadly grouped according to tumour site. These include breast (C50), bladder (C67), gynaecological (C51-57), prostate (C61), gastroesophageal (C15-16), haematological (C81-96), renal (C64-65), colorectal (C18-20), head and neck (C00-14, C30-32), hepatopancreaticobiliary (C22-25) and pulmonary (C34, C45) cancer. These groups were listed in order of the magnitude of their inflammatory status as shown previously [Proctor et al., 2010]. Deprivation was measured with the Scottish Index of Multiple Deprivation (SIMD) as detailed in Chapter 1 with the least deprived scoring 1 and the most deprived scoring 5.

Statistical analysis was carried out as follows. Survival, overall and cancer specific, was calculated from the time of cancer diagnosis to death. Cancer groups with less than 150 cancer

specific deaths were excluded to ensure statistical power. Analysis was carried on all cancer patients as well as on a sub-group of patients who had a blood sample taken within 2 months of their cancer diagnosis. This was carried out in order to examine the relationships between the mGPS, biochemical parameters and survival in all patients with a history of malignancy (patients sampled within 2 years following a diagnosis of cancer) and those at the time of diagnosis (patients sampled within 2 months following a diagnosis of cancer). When investigating the prognostic value of the systemic immune/inflammatory response and associated biochemical variables only blood samples taken after a diagnosis of cancer were included. This was carried out under the premise that it would be within this time period that the prognostic value of these variable would be of clinical utility. The mGPS and biochemical proportionality assumptions were explored using log-log plots and were found to be satisfactory. Kaplan-Meier estimator was used to analyse the relationship between patient characteristics, mGPS, biological parameters, tumour site and overall and cancer specific survival (Tables 5.3 and 5.4) as well as the relationship between mGPS and survival (Figure 5.2 and Table 5.9). Cox proportional hazards model multivariate regression analysis (stratified by tumour site) was used to determine the relationship between patient characteristics, the mGPS, and each biochemical parameter and survival (Tables 5.5, 5.6, 5.7, 5.8). Chi square (linear by linear) association was used to analyse the relationship between the mGPS, patient characteristics and biochemical parameters (Table 5.9). Due to the number of statistical comparisons a P value of less than 0.01 was considered significant. Analysis was performed using SPSS software (SPSS Inc., Chicago, IL, USA). The study was approved by the Research Ethics Committee, North Glasgow NHS Trust.

5.4. RESULTS

From Glasgow Inflammation Outcome Study of 223,303 patients originally described (Proctor et al, 2010), 21,668 patients were identified as having a diagnosis of cancer in the Scottish Cancer Registry and a blood sample taken between January 2000 and December 2006. There were 9608 patients in this group who had been sampled within two years following a diagnosis of cancer and included in the present Chapter. The majority, 7516 (78%), were under 75 years of age. There were 5116 (53%) females and 4492 (47%) males. All patients had an identifiable post code corresponding to a SIMD 2006 score with 15% of cases being from affluent areas (least deprived quintile of the Scottish population) and 38% being from deprived areas (most deprived quintile of the Scottish population). The minimum follow-up was 29 months and the maximum 112 months (median 61 months for survivors).

The relationship between patient characteristics, tumour site, mGPS, biochemical parameters and mortality in patients with blood samples taken within two years, as well as within two months, following a diagnosis is shown in Table 5.1 and Table 5.2. In those sampled within two years (n=9608), there were 5122 (53%) cancer deaths and the median time from blood sample to diagnosis was 1.4 months. In those sampled within two months (n=5397), there were 2993 (56%) cancer deaths and the median time from blood sample to diagnosis was 0.4 months. Breast and lung cancer were among the most common tumour types studied.

The relationship between the mGPS, biochemical parameters, tumour site and five year survival in patients with blood samples taken within two years, as well as two months, following a diagnosis of cancer is shown in Table 5.3, Table 5.4 and Figure 5.1. Increasing age, male gender and increasing deprivation was associated with reduced 5 year overall and cancer specific survival (all $p < 0.001$). An elevated mGPS, adjusted calcium, bilirubin, Alk phos, AST, ALT and GGT were associated with a reduced 5 year overall and cancer specific survival (all $p < 0.001$). These tables demonstrate the dramatic survival difference between tumours at

different sites as well as in those with raised markers of the systemic immune/inflammatory response.

In the present cohort sampled within two years, the majority of patients with a low albumin concentration (n=2701) also had an elevated C-reactive protein concentration (n=2419, 90%). Few patients had a low albumin but a C-reactive protein concentration in the normal range (n=282). A low albumin concentration alone was not significantly associated with cancer specific survival in bladder (p=0.913), gynaecological (p=0.737), prostate (p=0.500), gastroesophageal (p=0.893), renal (p=0.945), colorectal (p=0.133), head and neck (p=0.740), and hepatopancreaticobiliary (p=0.209) cancers. In Cox multivariate regression analysis the relationship between a mGPS of 2 and cancer specific survival (HR 3.01, p<0.001) was stronger than that between C-reactive protein alone and cancer specific survival (HR 2.29, p<0.001).

The relationship between the mGPS and cancer specific survival in breast (n=1956), bladder (n=466), gynaecological (n=533), prostate (n=491), gastroesophageal (n=869), haematological (n=974), renal (n=424), colorectal (n=1065), head and neck (n=501), hepatopancreaticobiliary (n=605) and pulmonary (n=1724) cancers are shown in Figures 5.2.

The relationship between the mGPS, biochemical parameters and survival in patients sampled within two years, as well as two months, following cancer diagnosis, adjusted for age, sex, deprivation and stratified by tumour site, is shown in Table 5.5 and Table 5.6. On survival analysis, a raised mGPS, adjusted calcium, bilirubin, Alk phos, AST, ALT and GGT were all associated with increased overall and cancer specific mortality independent of age, sex and deprivation (all p<0.001). These tables illustrate the associations between markers of the systemic immune/inflammatory response, biochemical variables and survival and highlight the relative strength of the mGPS as a predictive factor.

The relationship between the mGPS, patient demographics and survival in patients sampled within two years, as well as within two months, following cancer diagnosis is shown in Table 5.7 and 5.8. On multivariate survival analysis, stratified by tumour site, increasing age and mGPS were associated with increased overall and cancer specific mortality (all $p < 0.001$). Patients in the most deprived quintile (5) had a reduced overall and cancer specific survival (both $p < 0.01$) but a significant linear relationship across deprivation categories was not observed. These tables illustrate the associations between the mGPS, patient demographics and survival and highlight the relative strength of the mGPS, when compared to patient characteristics, as a predictive factor.

In the present cohort of patients sampled within two months following cancer diagnosis, only a limited number of patients had staging information available from the Scottish Cancer Registry. Tumour staging was available in 533 (39%) patients with breast cancer, 430 (76%) patients with colorectal cancer and 158 (13%) patients with pulmonary cancer. All other cancer groups had no staging available. Therefore, only in colorectal cancer was staging available in over 50% of patients. In these colorectal cancer patients there were 30 Dukes A, 113 Dukes B, 131 Dukes C and 156 Dukes D and 236 died of their cancer. When Dukes stage was included in the multivariate analysis, both Dukes stage (HR 3.59, 95% CI 2.95-4.39, $p < 0.001$) and mGPS (HR 1.49, 95% CI 1.26-1.76, $p < 0.001$) remained independently associated with survival.

The associations between an activated systemic immune/inflammatory response, as measured by an elevated mGPS, and other biochemical parameters in patients with blood samples taken within two years following a diagnosis of cancer is shown in Table 5.9. An increasing mGPS was also associated with increasing age, male gender, increasing adjusted calcium, bilirubin, Alk phos, AST, ALT and GGT (all $p < 0.001$). An increasing mGPS was shown to be associated with a reduction in overall and cancer specific survival (both $p < 0.001$).

This table illustrates the correlation between an activated systemic immune/inflammatory response and derangement of a number of other biochemical variables.

Figure 5.2 illustrates the relationship between the mGPS and cancer specific survival ($p < 0.001$) in each tumour site. This clearly demonstrates that the mGPS is predictive of survival independent of tumour site.

5.5. DISCUSSION

Previously, in large cohort studies, a number of biochemical parameters (other than C-reactive protein and albumin that compose the mGPS) including bilirubin [Temme et al., 2001], Alk phos [Tonelli et al., 2009], GGT [Kazemi-Shirazi et al., 2007] and calcium [Leifsson and Ahren, 1996], have been reported to predict overall and cancer specific survival.

In the present Chapter, the mGPS and the above biochemical parameters were shown to have prognostic value in all patients with a history of malignancy as well as those around the time of diagnosis. Moreover, a mGPS of 2 was associated with an approximate 160% reduction in both overall and cancer specific survival independent of tumour site. In contrast, an increase in adjusted calcium was associated with an approximate 130% reduction in both overall and cancer specific survival; an increase in bilirubin was associated with an approximate 50% reduction in both overall and cancer specific survival; an increase in Alk phos was associated with an approximate 110% reduction in both overall and cancer specific survival: an increase in AST was associated with an approximate 70% reduction in both overall and cancer specific survival; an increase in ALT was associated with an approximate 40% increase in both overall and cancer specific survival and an increase in GGT was associated with an approximate 80% reduction in both overall and cancer specific survival. These results indicate that the mGPS and the biochemical parameters measured have prognostic significance in the tumour sites studied.

Moreover, the results show that a raised mGPS is most closely associated with a reduction in both overall and cancer specific survival, independent of tumour site.

The GPS was originally developed, from the combination of C-reactive protein and albumin, in a cohort of patients with advanced non-small lung cancer [Forrest et al., 2004]. In this Chapter, they were combined to give a score of 0 for both a normal C-reactive protein and albumin, 1 for either an abnormal C-reactive protein alone or albumin alone and 2 for both an abnormal C-reactive protein and albumin. It was clear from this analysis that a low albumin alone was uncommon, accounting for less than 10% of all observations, and raised the possibility that this was not associated with a reduced survival. When examined in a cohort of patients undergoing potentially curable resection for colorectal cancer [McMillan et al., 2007] the results showed that the relationship between an abnormal albumin alone and cancer specific survival was similar to that of a normal albumin. Therefore, the GPS was modified (mGPS) to give a score of 1 only for an elevated C-reactive protein concentration. In the present Chapter, in a much larger cohort, a low albumin alone was associated with poor survival in some tumours (breast, haematological and pulmonary) but not others (bladder, gynaecological, prostate, gastroesophageal, renal, colorectal, head and neck, hepatopancreaticobiliary). Therefore, the present results would suggest the greater consistency of the mGPS and suggest its general use rather than that of the GPS.

In the present Chapter we examined the relationship between the mGPS, biochemical parameters and survival in all patients with cancer as well as a sub group of patients who were sampled within two months following a diagnosis of cancer. This was carried out with the premise that patients sampled within two years following a diagnosis would have a mGPS associated with disease prognosis and those sampled within two months would have a mGPS associated with their cancer diagnosis. With reference to the mGPS, the hazard ratios associated

with survival remained consistent in both analyses and confirms the temporal utility of this inflammation-based prognostic score.

To date the mGPS has been shown to have prognostic value, independent of TNM stage, in lung, gastrointestinal and renal cancers [McMillan, 2008; McMillan, 2009]. In the present Chapter there was insufficient staging information available on all cancer groups, apart from colorectal, to demonstrate that the mGPS is universally prognostic independent of stage. However, the results of the present and previous studies as detailed in Chapter 3 would suggest that the mGPS might also have independent prognostic in other cancer types [McMillan, 2009; Roxburgh and McMillan, 2010]. Further detailed studies, including tumour stage, in breast, bladder, gynaecological, prostate, haematological, head and neck, and hepatopancreaticobiliary cancers are required to confirm this hypothesis. Nevertheless, if this were to prove to be the case, then similar to the TNM staging system, the mGPS may be implemented universally in the assessment of cancer patients. Moreover, that they (TNM stage and mGPS) may be combined in a single staging system which would not only account for tumour stage but also the host systemic immune/inflammatory response.

The present cohort has a number of limitations. The patients were selected on the basis that measurements of C-reactive protein, albumin and calcium had been performed and were therefore not necessarily representative of all cancer patients diagnosed and treated in the North Glasgow area. It is also recognised that patients with cancer may have concurrent morbidity causing a rise in their C-reactive protein and derangement of their albumin and other biochemical parameters. Moreover, a rise in inflammatory markers may have occurred as a result of the initiation of oncological treatments including resectional surgery, radiotherapy and chemotherapy.

In summary, the results of the present Chapter indicate that, in a large patient cohort, the systemic immune/inflammatory response, as evidenced by the mGPS, is common and that the

mGPS is a powerful prognostic factor compared with other biochemical parameters, independent of tumour site in patients with cancer.

Table 5.1. The relationship between patient characteristics, mGPS, biochemical parameters, tumour site and mortality in patients sampled within two years following cancer diagnosis

			Patients n= 9608 (%)	All deaths n= 6005	Cancer deaths n=5122
Age	<65 years		4577 (48)	2258	2056
	65–74 years		2936 (30)	2041	1729
	≥ 75 years		2095 (22)	1706	1337
Sex	Male		4492 (47)	3269	2794
	Female		5116 (53)	2736	2328
SIMD 2006	1 (least deprived)		1419 (15)	722	616
	2		1198 (13)	631	543
	3		1445 (15)	850	739
	4		1858 (19)	1222	1038
	5 (most deprived)		3688 (38)	2580	2186
Tumour site	Breast		1956 (20)	452	328
	Bladder		466 (5)	250	155
	Gynaecological		533 (6)	298	256
	Prostate		491 (5)	244	165
	Gastroesophageal		869 (9)	772	719
	Haematological		974 (10)	539	418
	Renal		424 (5)	265	229
	Colorectal		1065 (11)	673	571
	Head and Neck		501 (5)	316	220
	Hepatopancreaticobiliary		605 (6)	563	536
	Pulmonary		1724 (18)	1633	1525
Inflammation based score	mGPS	0	3985 (42)	1647	1315
		1	3204 (33)	2325	2039
		2	2419 (25)	2033	1768
Biochemical parameters	Adjusted calcium	<2.10mmol/l	329 (4)	229	177
		2.10-2.60mmol/l	8856 (92)	5399	4601
		>2.60mmol/l	423 (4)	377	344
	Bilirubin	<20µmol/l	7936 (83)	4693	3978
		≥20µmol/l	1342 (14)	1071	942
	Alkaline phosphatase	<80U/l	177 (2)	56	48
		80-280U/l	6846 (71)	3704	3068
		>280U/l	2569 (27)	2236	1999
	Aspartate transaminase	<40U/l	7609 (18)	4439	3734
		≥40U/l	1678 (79)	1340	1206
Alanine transaminase	<50U/l	6350 (66)	3588	3036	
	≥50U/l	992 (10)	741	677	
γ-glutamyl transferase	M<70 U/l, F<40U/l	6082 (63)	3336	2765	
	M≥70U/l, F≥40U/l	3264 (34)	2488	2212	

Table 5.2. The relationship between patient characteristics, mGPS, biochemical parameters, tumour site and mortality in patients sampled within two months following cancer diagnosis

			Patients n= 5397 (%)	All deaths n= 3405	Cancer deaths n= 2993
Age	<65 years		2452 (46)	1192	1093
	65–74 years		1686 (31)	1172	1035
	≥ 75 years		1259 (23)	1041	865
Sex	Male		2315 (43)	1829	1620
	Female		3082 (57)	1576	1373
SIMD 2006	1 (least deprived)		802 (15)	392	335
	2		685 (13)	344	304
	3		766 (14)	429	388
	4		974 (18)	648	572
	5 (most deprived)		2170 (40)	1592	1394
Tumour site	Breast		1383 (26)	235	166
	Bladder		181 (3)	97	60
	Gynaecological		188 (3)	129	108
	Prostate		159 (3)	106	75
	Gastroesophageal		456 (8)	400	371
	Haematological		427 (8)	255	211
	Renal		179 (3)	117	107
	Colorectal		569 (11)	363	309
	Head and Neck		191 (4)	130	94
	Hepatopancreaticobiliary		486 (9)	455	438
	Pulmonary		1178 (22)	1118	1054
Inflammation based score	mGPS	0	2243 (41)	890	743
		1	1815 (34)	1385	1252
		2	1339 (25)	1130	998
Biochemical parameters	Adjusted calcium	<2.10mmol/l	165 (3)	116	91
		2.10-2.60mmol/l	4971 (92)	3058	2685
		>2.60mmol/l	261 (5)	231	217
	Bilirubin	<20µmol/l	4366 (81)	2558	2230
		≥20µmol/l	855 (16)	720	655
	Alkaline phosphatase	<80U/l	98 (2)	28	25
		80-280U/l	3788 (70)	2039	1746
		>280U/l	1502 (30)	1333	1217
	Aspartate transaminase	<40U/l	4191 (78)	2409	2084
		≥40U/l	1034 (19)	876	812
	Alanine transaminase	<50U/l	3740 (69)	2127	1847
≥50U/l		636 (12)	514	483	
γ-glutamyl transferase	M<70 U/l, F<40U/l	3393 (63)	1841	1581	
	M≥70U/l, F≥40U/l	1867 (35)	1472	1338	

Table 5.3. The relationship between patient characteristics, mGPS, biochemical parameters, tumour site and five year survival in patients sampled within two years following cancer diagnosis

n=9608			5 year overall survival %	p-value	5 year cancer survival %	p-value
Age	<65 years		50		54	
	65–74 years		30		38	
	≥ 75 years		19	<0.001	30	<0.001
Sex	Male		27		34	
	Female		47	<0.001	53	<0.001
SIMD	1 (least deprived)		50		55	
	2		47		52	
	3		41		46	
	4		34		41	
	5 (most deprived)		30	<0.001	37	<0.001
Tumour site	Breast		77		83	
	Bladder		57		64	
	Gynaecological		44		52	
	Prostate		50		64	
	Gastroesophageal		11		13	
	Haematological		46		54	
	Renal		39		38	
	Colorectal		37		43	
	Head and Neck		34		50	
	Hepatopancreaticobiliary		6		7	
	Pulmonary		5	<0.001	7	<0.001
Inflammation based score	mGPS	0	59		65	
		1	28		34	
		2	15	<0.001	21	<0.001
Biochemical parameters	Adjusted calcium	<2.10mmol/l	30		41	
		2.10-2.60mmol/l	39		45	
		>2.60mmol/l	10	<0.001	14	<0.001
	Bilirubin	<20µmol/l	41		47	
		≥20µmol/l	20	<0.001	25	<0.001
	Alkaline phosphatase	<80U/l	62		66	
		80-280U/l	46		53	
		>280U/l	13	<0.001	17	<0.001
	Aspartate transaminase	<40U/l	42		48	
		≥40U/l	20	<0.001	24	<0.001
Alanine transaminase	<50U/l	43		50		
	≥50U/l	25	<0.001	28	<0.001	
γ-glutamyl transferase	M<70 U/l, F<40U/l	45		52		
	M≥70U/l, F≥40U/l	23	<0.001	28	<0.001	

Table 5.4. The relationship between patient characteristics, mGPS, biochemical parameters, tumour site and five year survival in patients sampled within two months following cancer diagnosis

n=5397			5 year overall survival %	p-value	5 year cancer survival %	p-value
Age	<65 years		51		54	
	65–74 years		30		35	
	≥ 75 years		18	<0.001	26	<0.001
Sex	Male		21		26	
	Female		49	<0.001	54	<0.001
SIMD	1 (least deprived)		52		57	
	2		50		54	
	3		43		47	
	4		32		38	
	5 (most deprived)		27	<0.001	32	<0.001
Tumour site	Breast		83		87	
	Bladder		48		63	
	Gynaecological		33		39	
	Prostate		32		46	
	Gastroesophageal		12		15	
	Haematological		40		47	
	Renal		35		39	
	Colorectal		37		43	
	Head and Neck		29		43	
	Hepatopancreaticobiliary		5		6	
	Pulmonary		5	<0.001	7	<0.001
Inflammation based score	mGPS	0	60		65	
		1	24		29	
		2	15	<0.001	20	<0.001
Biochemical parameters	Adjusted calcium	<2.10mmol/l	29		39	
		2.10-2.60mmol/l	38		43	
		>2.60mmol/l	10	<0.001	13	<0.001
	Bilirubin	<20µmol/l	41		47	
		≥20µmol/l	15	<0.001	19	<0.001
	Alkaline phosphatase	<80U/l	68		69	
		80-280U/l	46		52	
		>280U/l	11	<0.001	14	<0.001
	Aspartate transaminase	<40U/l	42		48	
		≥40U/l	15	<0.001	18	<0.001
Alanine transaminase	<50U/l	43		48		
	≥50U/l	19	<0.001	21	<0.001	
γ-glutamyl transferase	M<70 U/l, F<40U/l	46		51		
	M≥70U/l, F≥40U/l	20	<0.001	25	<0.001	

Table 5.5. The relationship between the mGPS, biochemical parameters and survival in patients sampled within two years following cancer diagnosis. Adjusted for age, sex, deprivation and stratified by tumour site

n=9608		Overall survival		Cancer specific survival	
		HR	p value	HR	p value
mGPS	0	1	<0.001	1	<0.001
	1	1.79	<0.001	1.92	<0.001
	2	2.87	<0.001	3.01	<0.001
Adjusted calcium	<2.10mmol/l	1.25	0.001	1.15	0.070
	2.10-2.60mmol/l	1	<0.001	1	<0.001
	>2.60mmol/l	2.30	<0.001	2.38	<0.001
Bilirubin	<20µmol/l	1	<0.001	1	<0.001
	≥20µmol/l	1.53	<0.001	1.55	<0.001
Alkaline phosphatase	<80U/l	0.71	0.012	0.74	0.037
	80-280U/l	1	<0.001	1	<0.001
	>280U/l	2.12	<0.001	2.19	<0.001
Aspartate transaminase	<40U/l	1	<0.001	1	<0.001
	≥40U/l	1.66	<0.001	1.73	<0.001
Alanine transaminase	<50U/l	1	<0.001	1	<0.001
	≥50U/l	1.36	<0.001	1.39	<0.001
γ-glutamyl transferase	M<70 U/l, F<40U/l	1	<0.001	1	<0.001
	M≥70U/l, F≥40U/l	1.85	<0.001	1.85	<0.001

Table 5.6. The relationship between the mGPS, biochemical parameters and survival in patients sampled within two months following cancer diagnosis. Adjusted for age, sex, deprivation and stratified by tumour site

n=5397		Overall survival		Cancer specific survival	
		HR	p value	HR	p value
mGPS	0	1	<0.001	1	<0.001
	1	1.67	<0.001	1.74	<0.001
	2	2.36	<0.001	2.40	<0.001
Adjusted calcium	<2.10mmol/l	1.17	0.107	1.07	0.530
	2.10-2.60mmol/l	1	<0.001	1	<0.001
	>2.60mmol/l	1.96	<0.001	2.04	<0.001
Bilirubin	<20µmol/l	1	<0.001	1	<0.001
	≥20µmol/l	1.49	<0.001	1.49	<0.001
Alkaline phosphatase	<80U/l	0.78	0.016	0.68	0.053
	80-280U/l	1	<0.001	1	<0.001
	>280U/l	1.93	<0.001	1.96	<0.001
Aspartate transaminase	<40U/l	1	<0.001	1	<0.001
	≥40U/l	1.66	<0.001	1.71	<0.001
Alanine transaminase	<50U/l	1	<0.001	1	<0.001
	≥50U/l	1.34	<0.001	1.37	<0.001
γ-glutamyl transferase	M<70 U/l, F<40U/l	1	<0.001	1	<0.001
	M≥70U/l, F≥40U/l	1.79	<0.001	1.79	<0.001

Table 5.7. The relationship between patient characteristics, mGPS and survival in patients sampled within two years following cancer diagnosis. Multivariate analysis stratified by tumour site

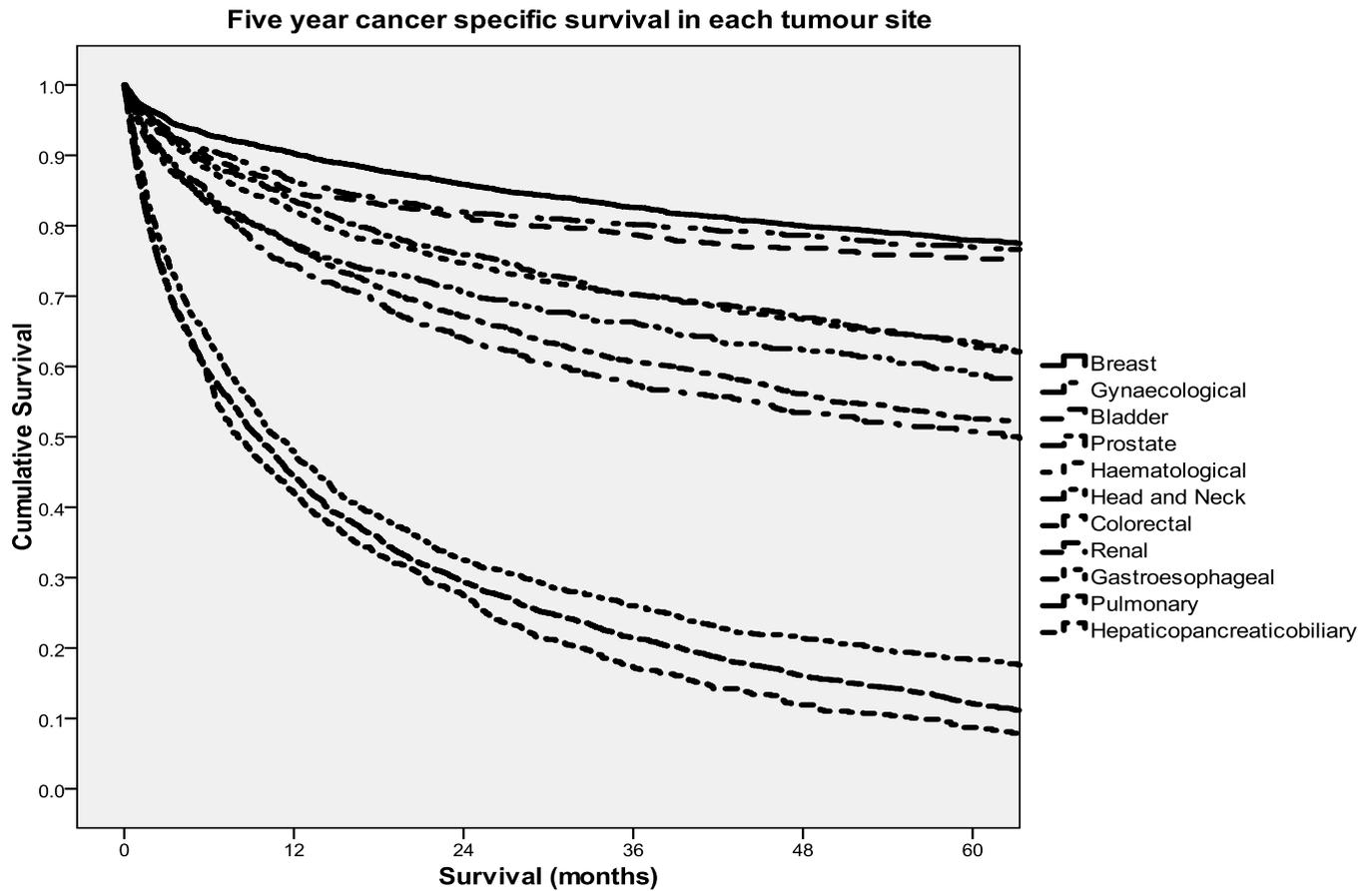
n=9608		Overall survival			Cancer specific survival		
		HR	95 % CI	p value	HR	95 % CI	p value
Age	<65 years	1		<0.001	1		<0.001
	65–74 years	1.35	1.26-1.43	<0.001	1.20	1.13-1.29	<0.001
	≥ 75 years	1.93	1.80-2.06	<0.001	1.59	1.48-1.70	<0.001
Sex	Male	1		0.037	1		0.132
	Female	0.94		0.037	0.95		0.132
SIMD	1 (least deprived)	1		<0.001	1		0.003
	2	1.02		0.684	1.02		0.773
	3	1.09		0.090	1.09		0.136
	4	1.16	1.06-1.27	0.002	1.12		0.025
	5 (most deprived)	1.22	1.12-1.33	<0.001	1.17	1.06-1.28	0.001
mGPS	0	1		<0.001	1		<0.001
	1	1.79	1.68-1.91	<0.001	1.92	1.79-2.06	<0.001
	2	2.87	2.68-3.08	<0.001	3.01	2.79-3.24	<0.001

Table 5.8. The relationship between patient characteristics, mGPS and survival in patients sampled within two months following cancer. Multivariate analysis stratified by tumour site

		Overall survival			Cancer specific survival		
		HR	95 % CI	p value	HR	95 % CI	p value
n=5397							
Age	<65 years	1		<0.001	1		<0.001
	65–74 years	1.31	1.20-1.42	<0.001	1.22	1.12-1.33	<0.001
	≥ 75 years	1.92	1.76-2.10	<0.001	1.70	1.55-1.87	<0.001
Sex	Male	1		0.340	1		0.507
	Female	0.96		0.340	0.97		0.507
SIMD	1 (least deprived)	1		<0.001	1		<0.001
	2	1.05		0.503	1.08		0.364
	3	1.07		0.342	1.11		0.179
	4	1.20	1.06-1.36	0.005	1.20	1.05-1.38	0.008
	5 (most deprived)	1.30	1.16-1.45	<0.001	1.28	1.14-1.45	<0.001
mGPS	0	1		<0.001	1		<0.001
	1	1.67	1.53-1.83	<0.001	1.74	1.59-1.91	<0.001
	2	2.36	2.15-2.59	<0.001	2.40	2.17-2.65	<0.001

Table 5.9. The relationship between the mGPS, patient characteristics, biochemical parameters and survival in patients sampled within two years following cancer diagnosis

		mGPS			p value
		0 n (%)	1 n (%)	2 n (%)	
n=9608					
Age	<65 years	2193 (55)	1464 (46)	920 (38)	
	65–74 years	1091 (27)	1007 (31)	838 (35)	
	≥ 75 years	701 (18)	733 (23)	661 (27)	<0.001
Sex	Male	1516 (38)	1672 (52)	1304 (54)	
	Female	2469 (62)	1532 (48)	1115 (46)	<0.001
SIMD 2006	1 (least deprived)	739 (19)	383 (12)	297 (12)	
	2	592 (15)	342 (11)	264 (11)	
	3	633 (16)	460 (14)	352 (15)	
	4	727 (18)	613 (19)	518 (21)	
	5 (most deprived)	1294 (32)	1406 (44)	988 (41)	<0.001
Adjusted calcium	<2.10mmol/l	76 (2)	87 (3)	166 (7)	
	2.10-2.60mmol/l	3835 (96)	2972 (93)	2049 (85)	
	>2.60mmol/l	74 (2)	145 (4)	204 (8)	<0.001
Bilirubin	<20µmol/l	3564 (92)	2621 (85)	1751 (76)	
	≥20µmol/l	324 (8)	466 (15)	552 (24)	<0.001
Alkaline phosphatase	<80U/l	93 (2)	16 (0)	68 (3)	
	80-280U/l	3428 (86)	2163 (68)	1255 (52)	
	>280U/l	457 (12)	1020 (32)	1092 (45)	<0.001
Aspartate transaminase	<40U/l	3491 (90)	2501 (81)	1617 (70)	
	≥40U/l	399 (10)	593 (19)	686 (30)	<0.001
Alanine transaminase	<50U/l	3016 (91)	1982 (85)	1352 (79)	
	≥50U/l	301 (9)	337 (15)	354 (21)	<0.001
γ-glutamyl transferase	M<70 U/l, F<40U/l	3071 (79)	1875 (60)	1136 (49)	
	M≥70U/l, F≥40U/l	839 (21)	1242 (40)	1183 (51)	<0.001
Overall survival in months (mean, CI)		70 (68-72)	37 (36-39)	22 (21-24)	<0.001
Cancer specific survival in months (mean, CI)		77 (75-79)	43 (41-45)	27 (26-29)	<0.001



Number at risk

Alive	9608	5492	4566	3710	2732	1845
Overall deaths	0	4116	5042	5898	6876	7763
Cancer deaths	0	3658	4452	4795	4961	5052

Figure 5.1. The relationship between tumour site and cancer specific five year survival. Tumours from top to bottom; breast, gynaecological, bladder, prostate, haematological, head and neck, colorectal, renal, gastroesophageal, pulmonary and hepatopancreaticobiliary

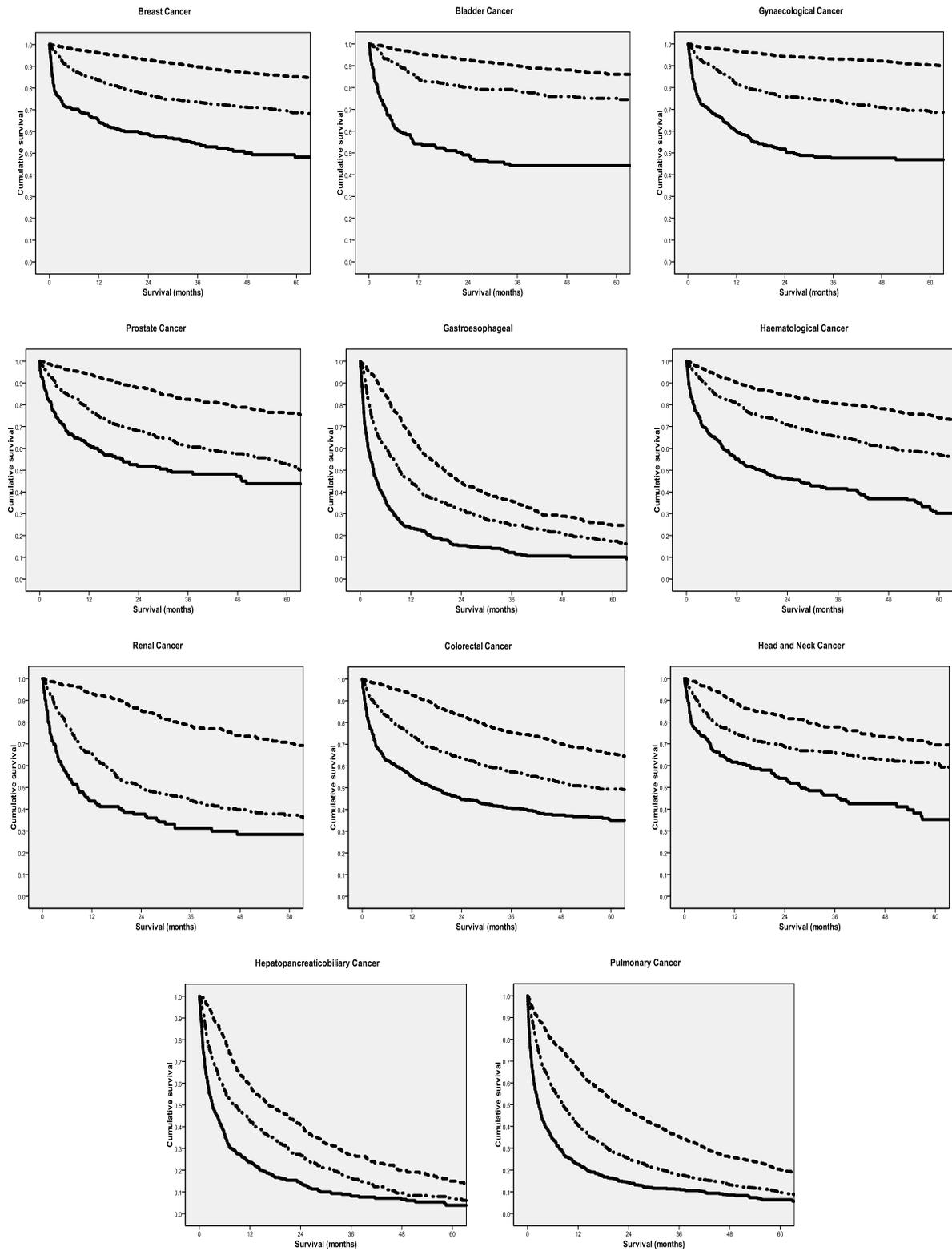


Figure 5.2. The relationship between the mGPS - 0 (top, small dash line), mGPS 1 (middle, large dash line) and mGPS 2 (bottom, solid line) - and cancer specific survival ($p < 0.001$) in each tumour site

6.0. A COMPARISON OF INFLAMMATION-BASED PROGNOSTIC SCORES IN PATIENTS WITH CANCER

6.1. ABSTRACT

Introduction: Components of the systemic inflammatory response, combined to form inflammation-based prognostic scores including the modified Glasgow Prognostic Score (mGPS), Neutrophil Lymphocyte Ratio (NLR), Platelet Lymphocyte Ratio (PLR), Prognostic Index (PI) and Prognostic Nutritional Index (PNI), have been associated with cancer specific survival. The aim of the present chapter was to compare the prognostic value of these scores.

Methods: Patients (n=27,031) who had an incidental blood sample taken between 2000 and 2007 for C-reactive protein, albumin, white cell, neutrophil, lymphocyte and platelet counts, as well as a diagnosis of cancer (Scottish Cancer Registry) were identified. Of this group 8759 patients who had been sampled within two years following their cancer diagnosis were studied.

Results: On follow up there were 5163 deaths, 4417 (86%) of which were cancer related. The median time from blood sampling to diagnosis was 1.7 months. An elevated mGPS, NLR, PLR, PI and PNI were predictive of a reduced cancer specific survival independent of age, sex and deprivation and tumour site (all $p < 0.001$). The area under the receiver operator curves was greatest for mGPS and PI. Specifically, in colorectal cancer, an elevated mGPS and PI were predictive of a reduced cancer specific survival independent of age, sex, deprivation and tumour stage (both $p < 0.001$).

Conclusion: The results of the present study show that systemic inflammation-based scores, in particular the mGPS and PI, have prognostic value in cancer independent of tumour site.

6.2. INTRODUCTION

As has been detailed in Chapter 1, while the development of cancer has a genetic basis, there is increasing evidence that the host immune/inflammatory response plays an important role in the development and progression of cancer [Coussens and Werb, 2002; Mantovani et al., 2008; Colotta et al., 2009; McDonald et al., 2009; Tenesa et al., 2010]. The modified Glasgow Prognostic Score (mGPS) has been shown to have prognostic value, independent of tumour stage, in a number of tumour types including lung, gastrointestinal and renal cancers [McMillan, 2008; McMillan, 2009]. As shown in Table 3.1, it is also of interest that other haematological components of the systemic immune/inflammatory response have been combined to form inflammation-based prognostic scores that have been associated with survival in patients with cancer. In addition to those detailed in Chapter 3, the Neutrophil Lymphocyte Ratio (NLR), a combination of circulating neutrophil and lymphocyte counts, has been associated with survival in colorectal [Chua et al., 2011a], lung [Sarraf et al., 2009; Kao et al., 2010a] and ovarian [Cho et al., 2009b] cancers. The Platelet Lymphocyte Ratio (PLR), a combination of circulating platelet and lymphocyte counts, has been associated with survival in patients with pancreatic cancer [Smith et al., 2009]. The combination of C-reactive protein and white cell count in a Prognostic Index (PI) has been associated with survival in patients with lung cancer [Kasymjanova et al., 2010]. Finally, Onodera's Prognostic Nutritional Index (PNI) has also been associated with survival in patients with pancreatic [Kanda et al., 2011] and gastric [Nozoe et al., 2010] cancer.

In Chapter 4 and 5, it has been shown that the mGPS is elevated in patients with cancer [Proctor et al., 2010] and predictive of survival across all tumour sites studied independent of age, sex and deprivation [Proctor et al., 2011b]. Therefore, it is of considerable interest to compare the prognostic value of the mGPS, NLR, PLR, PI and PNI across different tumour sites.

The aim of the current Chapter was to compare the prognostic value of the mGPS, NLR, PLR, PI and PNI adjusted for age, sex, deprivation and tumour site in the Glasgow Inflammation Outcome Study.

6.3. MATERIALS AND METHODS

From a cohort previously described in Chapter 4. [Proctor et al., 2010], patients in the North Glasgow who were sampled incidentally for C-reactive protein, albumin, calcium, white cell, neutrophil, lymphocyte and platelet counts between the 1st January 2000 and the 31st December 2007 were included. This time period differed from previous Chapters to correspond with the available follow-up data from the Scottish Cancer Registry at the time of analysis. The limit of detection of C-reactive protein was a concentration of less than 5mg/L. The mGPS, NLR, PLR, PI and PNI were constructed as described in Table 6.1.

Cohort linkage, with the Scottish Cancer Registry, classification of deaths and cancer coding were carried out as described in Chapter 5 and only those patients who had complete Cancer Registry follow-up were included [Proctor et al., 2011b]. Survival was calculated as described in Chapter 5 with cancer registry follow-up available until June the 30th 2009 [Proctor et al., 2011b]. Cancer groups with less than 150 cancer specific deaths were excluded to ensure statistical power. Analysis was carried on all cancer patients who had a blood sample taken within two years of diagnosis as well as on a sub-group of patients who had a sample taken within two months of a diagnosis of cancer. This was carried out in order to examine the relationships between the mGPS, NLR, PLR, PI, PNI and survival in all patients with a recently diagnosed malignancy (patients sampled within two years following a diagnosis of cancer) and those at the time of diagnosis (patients sampled within two months following a diagnosis of cancer). Cancer stage was extracted from the registry where available.

Deprivation was measured with the Scottish Index of Multiple Deprivation (SIMD) as detailed in Chapter 1. Patients with incomplete identifying details (name, date of birth and hospital number), under 16 years of age, or those with multiple malignancies, metastatic disease or cancer of an unknown origin or those without all haematological variables available were excluded. Overall patient numbers were different in this Chapter, when compared to previous, as the inclusion time period was longer as extended cancer registry follow-up was available. The study was approved by the Research Ethics Committee, North Glasgow NHS Trust.

Statistical analysis was carried out as follows. The mGPS, NLR, PLR, PI, PNI proportionality assumptions were explored using log-log plots and were found to be satisfactory. Kaplan-Meier estimator was used to analyse the relationship between patient characteristics, mGPS, NLR, PLR, PI, PNI, tumour site and overall and cancer specific survival (Table 6.2, Figures 6.1, 6.2). Cox proportional hazards model multivariate regression analysis (stratified by tumour site) was used to correct for age, sex and deprivation and determine the relationship between patient characteristics, mGPS, NLR, PLR, PI, PNR and survival (Tables 6.3, 6.4). To determine whether one of the independently significant variables was more predictive than the other, the area under the receiver operating characteristic (ROC) curve was calculated. Analysis was performed using SPSS software (SPSS Inc., Chicago, IL, USA).

6.4. RESULTS

From the Glasgow Inflammation Outcome Study of 223,303 patients originally described in Chapter 4. [Proctor et al., 2010], 27,031 patients were identified as having a diagnosis of cancer in the Scottish Cancer Registry and a blood sample for taken for C-reactive protein, albumin, white cell, neutrophil, lymphocyte and platelet counts between

January 2000 and December 2007. Within this group there were 8759 patients who had been sampled for all these variables within two years following a diagnosis of cancer and were included in the present study. The majority, 6857 (78%), were under 75 years of age. There were 4644 (53%) females and 4115 (47%) males. Fifteen per cent of patients lived in affluent areas (least deprived quintile of the Scottish population) and 36% in deprived areas (most deprived quintile of the Scottish population). The minimum follow-up from cancer diagnosis was 18 months and the maximum 115 months (median 51 months for survivors).

The relationship between patient characteristics, tumour site, inflammatory-based prognostic scores and survival is shown in Table 6.2 and Figure 6.1. In total 8759 patients were studied. On follow-up, there were 5163 deaths of which 4417 (86%) were cancer deaths. The median time from diagnosis to blood sampling was 1.7 months, suggesting that most scores reflect status at diagnosis rather than at a later stage. Increasing age, male gender, and increasing deprivation were associated with reduced 5 year overall and cancer specific survival (all $p < 0.001$). An elevated mGPS, NLR, PLR, PI and PNI were also associated with a reduced 5 year overall and cancer specific survival (all $p < 0.001$). When this analysis was repeated on a subgroup of patients ($n=4674$) who were sampled within two months following their cancer diagnosis (median time to diagnosis 0.5 months) the above proportions remained similar and the associations significant (all $p < 0.001$). The relationship between the mGPS, NLR, PLR, PI and PNI and cancer specific survival in individual tumour groups is shown in Figure 6.2. All were predictive of cancer specific survival in bladder, breast, colorectal, gastroesophageal, gynaecological, prostate, pulmonary and renal cancer (all $p < 0.001$). Additionally, the mGPS, NLR, PI and PNI were predictive of cancer specific haematological, head and neck, and hepatopancreaticobiliary cancer survival (all $p < 0.001$).

The relationship between inflammation-based prognostic scores and survival, adjusted for age, sex and deprivation and stratified by tumour site is shown in Table 6.3. On survival

analysis of all patients (n=8759), a raised mGPS, NLR, PLR, PI and PNI were associated with reduced overall and cancer specific survival independent of age, sex and deprivation (all $p<0.001$). When this analysis was repeated on a subgroup of patients (n=4674) who were sampled two months following their cancer diagnosis the above proportions remained similar and the associations significant (all $p<0.001$).

In all patients, using overall mortality at as an endpoint, the area under the receiver operator curve was for the mGPS 0.718 (95% CI, 0.707-0.729, $p<0.001$); NLR 0.650 (95% CI, 0.638-0.661, $p<0.001$); PLR 0.632 (95% CI, 0.620-0.644, $p<0.001$); PI 0.715 (95% CI, 0.704-0.726, $p<0.001$) and PNI 0.678 (95% CI, 0.666-0.689, $p<0.001$). Using cancer-specific mortality at as an endpoint, the area under the receiver operator curve was for the mGPS 0.698 (95% CI, 0.687-0.709, $p<0.001$); NLR 0.632 (95% CI, 0.620-0.643, $p<0.001$); PLR 0.632 (95% CI, 0.620-0.644, $p<0.001$); PI 0.694 (95% CI, 0.682-0.705, $p<0.001$) and PNI 0.660 (95% CI, 0.649-0.672, $p<0.001$).

When this analysis was repeated on a subgroup of patients (n=4674) who were sampled within two months following their cancer diagnosis, using overall mortality as an endpoint, the area under the receiver operator curve was for the mGPS 0.730 (95% CI, 0.715-0.745, $p<0.001$); NLR 0.657 (95% CI, 0.641-0.672, $p<0.001$); PLR 0.644 (95% CI, 0.628-0.660, $p<0.001$); PI 0.724 (95% CI, 0.724-0.753, $p<0.001$) and PNI 0.692 (95% CI, 0.677-0.708, $p<0.001$). Using cancer-specific mortality as an endpoint, the area under the receiver operator curve was for the mGPS 0.712 (95% CI, 0.697-0.727, $p<0.001$); NLR 0.640 (95% CI, 0.624-0.656, $p<0.001$); PLR 0.638 (95% CI, 0.622-0.654, $p<0.001$); PI 0.719 (95% CI, 0.704-0.733., $p<0.001$) and PNI 0.673 (95% CI, 0.658-0.689, $p<0.001$).

In the present cohort only a limited number of patients had staging information available from the Scottish Cancer Registry. Of those patients sampled within two months

following cancer diagnosis, tumour staging was available in 470 (35%) patients with breast cancer, 374 (75%) patients with colorectal cancer and 108 (12%) patients with pulmonary cancer. All other cancer groups had no staging available. Therefore, only in colorectal cancer was staging available in over 50% of patients. In this group there were 31 Dukes A, 113 Dukes B, 113 Dukes C and 117 Dukes D with 227 dying of their cancer on follow-up. The relationship between inflammation-based prognostic scores and survival in colorectal cancer patients sampled within two months following cancer diagnosis adjusted for age, sex, deprivation and Dukes stage is shown in Table 6.4. Given the small numbers of Dukes A, Dukes A and B were grouped together. On survival analysis (n=374), only a raised mGPS and PI were associated with reduced overall and cancer specific survival independent of age, sex, deprivation and Dukes stage (all $p < 0.001$).

As mGPS and PI both contain C-reactive protein, the prognostic value of a low albumin ($< 35\text{g/l}$) or high white cell count ($> 11 \times 10^9/\text{l}$), independent of each other was of interest. On Cox-regression multivariate survival analysis of the same group of patients (n=374), both a high white cell count (HR 1.57, $p = 0.011$) and low albumin (HR 1.59, $p = 0.004$) were predictive of cancer specific survival independent of age, sex, deprivation and Dukes stage.

6.5. DISCUSSION

The results of the present Chapter show clearly that systemic inflammation-based prognostic scores, whether it be the mGPS, NLR, PLR, PI or PNI, predict cancer specific outcome in most cancers. Moreover, scores including C-reactive protein (mGPS and PI) were superior, in terms of differentiating good from poor prognostic groups, to those based on components of the circulating white cell count (NLR, PLR) or in combination with albumin (PNI). Therefore, any further development of prognostic scores, based on the

systemic immune/inflammatory response, should include the prototypical acute phase protein [Gabay and Kushner, 1999], C-reactive protein. Also, based on the present results and the existing validation literature [McMillan, 2009], the mGPS should be included in the routine assessment of all patients with cancer.

In the present Chapter, it was of interest that the combination of C-reactive protein and albumin (mGPS) had similar prognostic value to that of C-reactive protein and white cell count (PI). Although few patients had a low albumin (5%) or a high white cell count (3%) in the presence of a normal C-reactive protein concentration ($<10\text{mg/l}$), both a low albumin concentration and a high white cell count both had prognostic value independent of C-reactive protein. These results would suggest that a white cell count, rather than the NLR or PLR, may be a useful addition to the comprehensively validated mGPS.

It was also of interest that a nutritional index previously shown to have prognostic value (PNI) and used extensively in Japan [Nozoe et al., 2002; Nozoe et al., 2010; Kanda et al., 2011], behaved, in terms of prognostic value, very similar to the systemic inflammation-based prognostic scores (mGPS, NLR, PLR and PI). The present results, while only demonstrating correlation and not causation, add further weight to the proposal that the systemic immune/inflammatory response is a major factor in the relationship between nutritional decline and poor outcome in patients with cancer [McMillan, 2008; McMillan, 2009].

Taken together, the results of the present Chapter highlight the associations between systemic inflammation and outcome in patients with cancer. Therefore, we believe that the present results provide good evidence of systemic inflammation acting as a “common soil” [Fidler and Poste, 2008] promoting the fatal progression in most, if not all, cancers. If this

proves to be the case then moderation of the systemic immune/inflammatory response will become, in the future, as important a therapeutic target as the tumour itself.

In this context it is of interest that Chechlinska, Kowalewska and co-workers [Chechlinska et al., 2010] have proposed that “systemic inflammation is a confounding factor in the interpretation of the biomarker profile of cancer patients” and “to assess the independent predictive value of a biomarker, it should be validated against its expression in inflammatory conditions, and examined in the context of unspecific parameters of systemic inflammation”. Also, they have concluded that if this is not done “we will end up using advanced technologies to assess inflammatory reactions in cancer patients”, an opinion shared by the authors of the present body of work.

In summary, the results of the present Chapter show that systemic inflammation-based scores mGPS, NLR, PLR, PI and PNI have prognostic value in a variety of cancers. However, in terms of differentiating good from poor prognostic groups in a variety of tumour sites, the extensively validated mGPS is superior. A measurement of systemic inflammation, in particular the mGPS, should be included in the routine assessment of all patients with cancer.

Table 6.1. Systemic inflammation-based prognostic scores

The modified Glasgow Prognostic Score	Score
C-reactive protein ≤ 10 mg/l and albumin ≥ 35 g/l	0
C-reactive protein ≤ 10 mg/l and albumin < 35 g/l	0
C-reactive protein > 10 mg/l	1
C-reactive protein > 10 mg/l and albumin < 35 g/l	2
Neutrophil Lymphocyte Ratio	
Neutrophil count:lymphocyte count $< 5:1$	0
Neutrophil count:lymphocyte count $\geq 5:1$	1
Platelet Lymphocyte Ratio	
Platelet count:lymphocyte count $< 150:1$	0
Platelet count:lymphocyte count 150–300:1	1
Platelet count:lymphocyte count $> 300:1$	2
Prognostic Index	
C-reactive protein ≤ 10 mg/l and white cell count $\leq 11 \times 10^9/l$	0
C-reactive protein ≤ 10 mg/l and white cell count $> 11 \times 10^9/l$	1
C-reactive protein > 10 mg/l and white cell count $\leq 11 \times 10^9/l$	1
C-reactive protein > 10 mg/l and white cell count $> 11 \times 10^9/l$	2
Prognostic Nutritional Index	
Albumin (g/L) + 5 \times total lymphocyte count $\times 10^9/l \geq 45$	0
Albumin (g/L) + 5 \times total lymphocyte count $\times 10^9/l < 45$	1

Table 6.2. The relationship between patient characteristics, tumour site, inflammatory-based prognostic scores and survival

			Patients n=8759 (%)	Overall survival 5 year % (n of deaths) n=5163	p- value	Cancer survival 5 year % (n of deaths) n=4417	p- value
Age	<65 years		4237 (48)	52 (1977)		55 (1808)	
	65–74 years		2620 (30)	33 (1703)		41 (1439)	
	≥ 75 years		1902 (22)	21 (1483)	<0.001	31 (1170)	<0.001
Sex	Male		4115 (47)	29 (2844)		36 (2432)	
	Female		4644 (53)	49 (2319)	<0.001	55 (1985)	<0.001
SIMD 2006	1 (least deprived)		1278 (15)	51 (609)		57 (523)	
	2		1138 (13)	48 (579)		54 (495)	
	3		1391 (16)	43 (779)		48 (683)	
	4		1786 (20)	37 (1110)		44 (940)	
	5 (most deprived)		3166 (36)	33 (2086)	<0.001	40 (1776)	<0.001
Tumour site	Breast		1853 (21)	79 (361)		85 (263)	
	Bladder		437 (5)	48 (226)		63 (149)	
	Gynaecological		460 (5)	45 (248)		51 (217)	
	Prostate		456 (5)	53 (206)		64 (153)	
	Gastroesophageal		874 (10)	12 (754)		15 (697)	
	Haematological		817 (10)	48 (418)		57 (320)	
	Renal		400 (5)	38 (242)		44 (214)	
	Colorectal		996 (11)	39 (583)		45 (493)	
	Head and Neck		555 (7)	34 (344)		51 (239)	
	Hepatopancreaticobiliary		474 (5)	7 (430)		8 (410)	
	Pulmonary		1437 (16)	5 (1351)	<0.001	7 (1262)	<0.001
Inflammation based scores	mGPS	0	3673 (42)	61 (1349)		68 (1083)	
		1	2436 (28)	32 (1640)		39 (1425)	
		2	2650 (30)	16 (2174)	<0.001	22 (1909)	<0.001
	NLR	0	5151 (59)	51 (2401)		58 (2021)	
		1	3608 (41)	23 (2762)	<0.001	29 (2396)	<0.001
	PLR	0	2734 (31)	52 (1253)		60 (996)	
		1	3522 (40)	42 (1993)		48 (1716)	
		2	2503 (29)	23 (1917)	<0.001	28 (1705)	<0.001
	PI	0	3084 (35)	64 (1042)		70 (832)	
		1	3460 (40)	31 (2303)		38 (1994)	
		2	2215 (25)	17 (1818)	<0.001	23 (1591)	<0.001
	PNI	0	4342 (50)	57 (1806)		63 (1487)	
		1	4417 (50)	21 (3357)	<0.001	27 (2930)	<0.001

Table 6.3. The relationship between inflammation-based prognostic scores and survival. Adjusted for age, sex, deprivation and stratified by tumour site

All patients (n=8759)

		Overall survival		Cancer specific survival	
		HR	p value	HR	p value
mGPS	0	1	<0.001	1	<0.001
	1	1.74	<0.001	1.85	<0.001
	2	2.91	<0.001	3.06	<0.001
NLR	0	1	<0.001	1	<0.001
	1	1.93	<0.001	1.97	<0.001
PLR	0	1	<0.001	1	<0.001
	1	1.22	<0.001	1.31	<0.001
	2	1.89	<0.001	2.08	<0.001
PI	0	1	<0.001	1	<0.001
	1	2.03	<0.001	2.15	<0.001
	2	2.87	<0.001	3.03	<0.001
PNI	0	1	<0.001	1	<0.001
	1	2.24	<0.001	2.34	<0.001

Patients sampled within two months following cancer diagnosis (n=4674)

		Overall survival		Cancer specific survival	
		HR	p value	HR	p value
mGPS	0	1	<0.001	1	<0.001
	1	1.65	<0.001	1.74	<0.001
	2	2.35	<0.001	2.44	<0.001
NLR	0	1	<0.001	1	<0.001
	1	1.76	<0.001	1.77	<0.001
PLR	0	1	<0.001	1	<0.001
	1	1.19	<0.001	1.24	<0.001
	2	1.71	<0.001	1.82	<0.001
PI	0	1	<0.001	1	<0.001
	1	1.78	<0.001	1.87	<0.001
	2	2.44	<0.001	2.51	<0.001
PNI	0	1	<0.001	1	<0.001
	1	1.98	<0.001	2.01	<0.001

Table 6.4. The relationship between inflammation-based prognostic scores and survival in colorectal cancer patients sampled within two months following cancer diagnosis. Adjusted for age, sex, deprivation and Dukes stage

n=374		Overall survival		Cancer specific survival	
		HR	p value	HR	p value
mGPS	0	1	<0.001	1	<0.001
	1	1.81	0.004	1.91	<0.001
	2	2.30	<0.001	2.51	<0.001
NLR	0	1	0.102	1	0.146
	1	1.27	0.102	1.25	0.146
PLR	0	1	0.786	1	0.560
	1	1.16	0.487	1.30	0.281
	2	1.13	0.596	1.23	0.403
PI	0	1	<0.001	1	<0.001
	1	1.69	0.012	1.92	<0.001
	2	2.83	<0.001	3.07	<0.001
PNI	0	1	0.059	1	0.095
	1	1.33	0.059	1.31	0.095

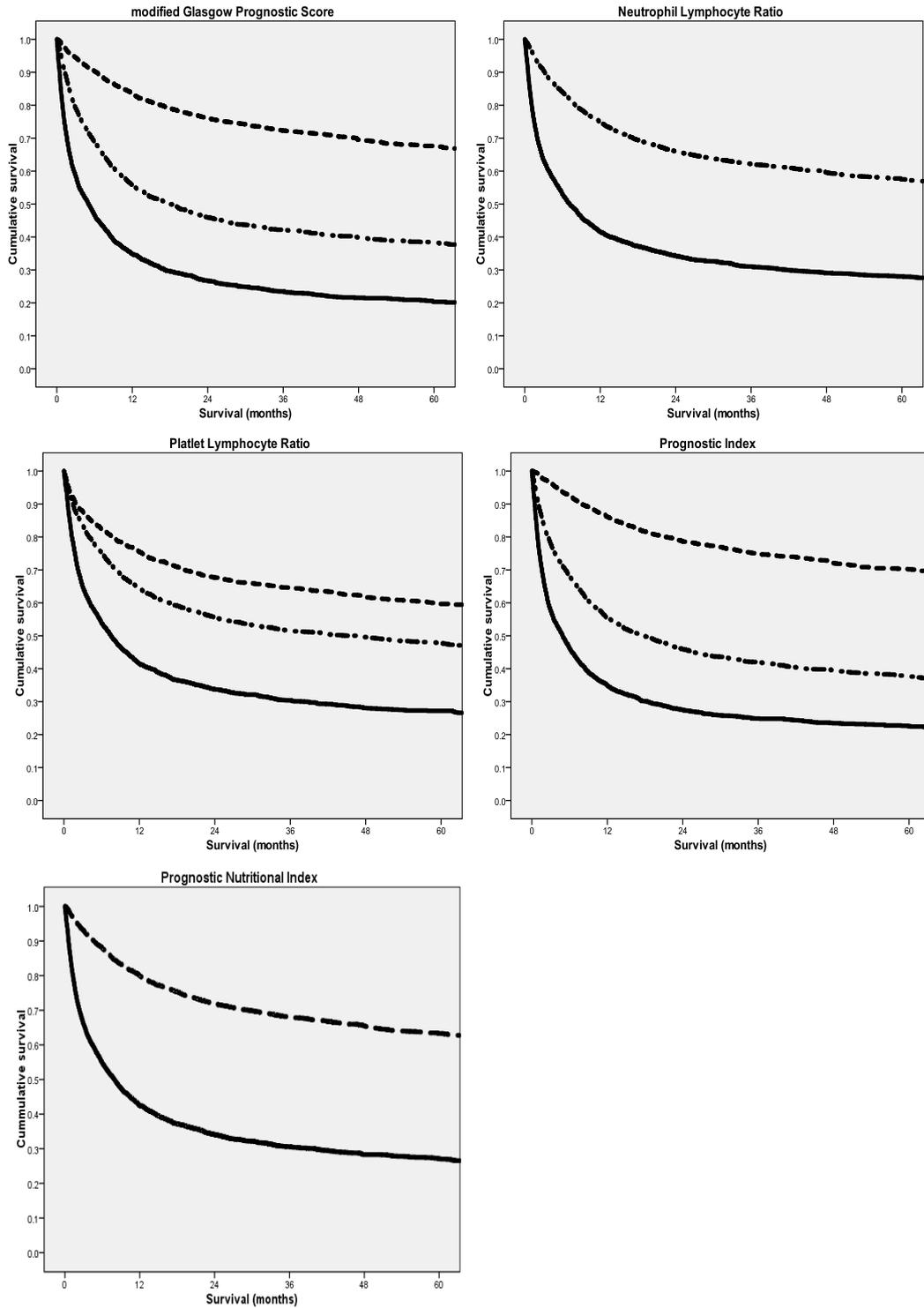
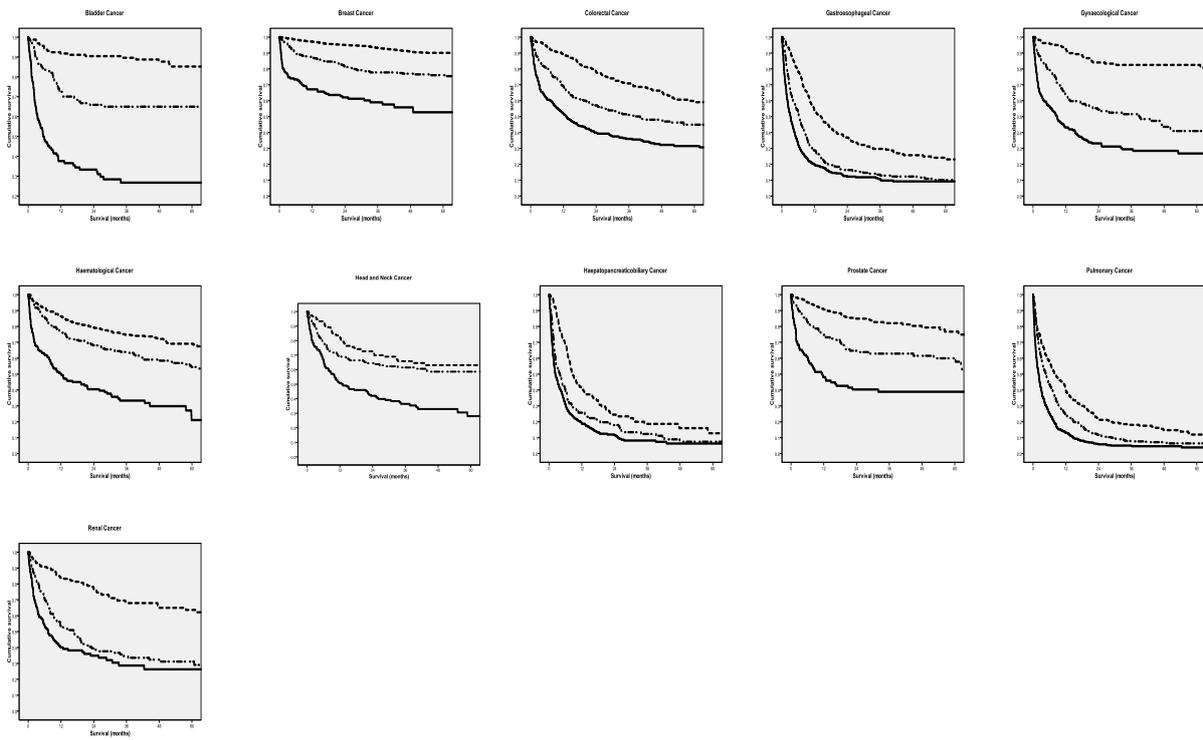
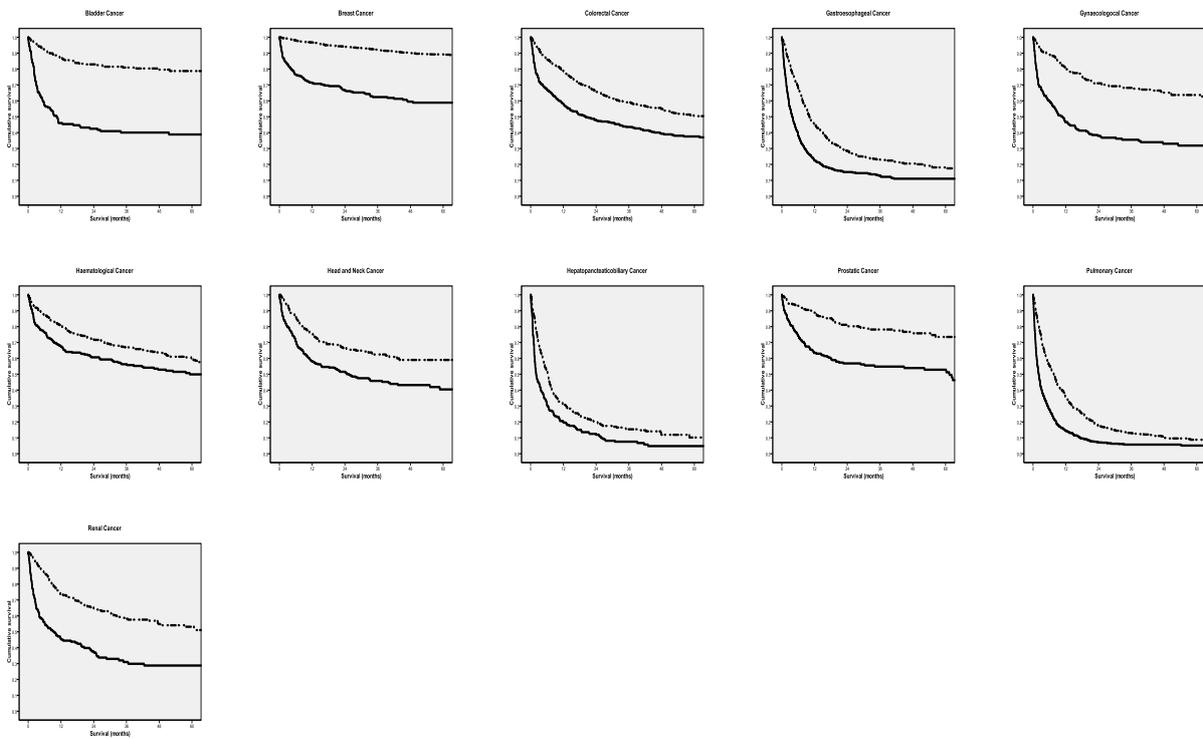


Figure 6.1. The relationship between the mGPS (0-top, small dash line; 1-middle, large dash line; 2-bottom, solid line), NLR (0-top, large dash line; 1-bottom, solid line), PLR (0-top, small dash line; 1-middle, large dash line; 2-bottom, solid line), PI (0-top, small dash line; 1-middle, large dash line; 2-bottom, solid line), PNI (0-top, large dash line; 1-bottom, solid line) and cancer specific survival in all patients (all <math><0.001</math>).

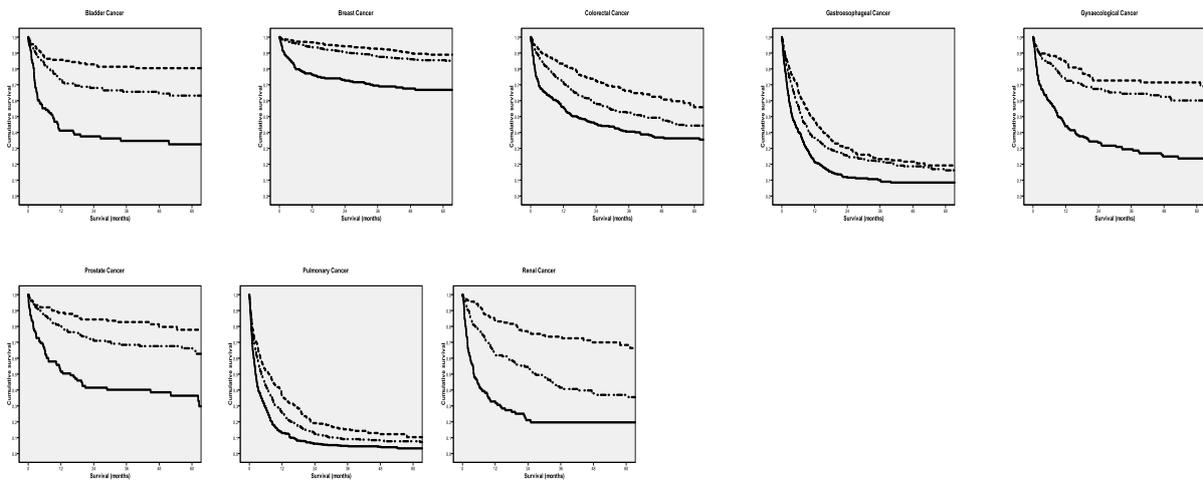
modified Glasgow Prognostic Score



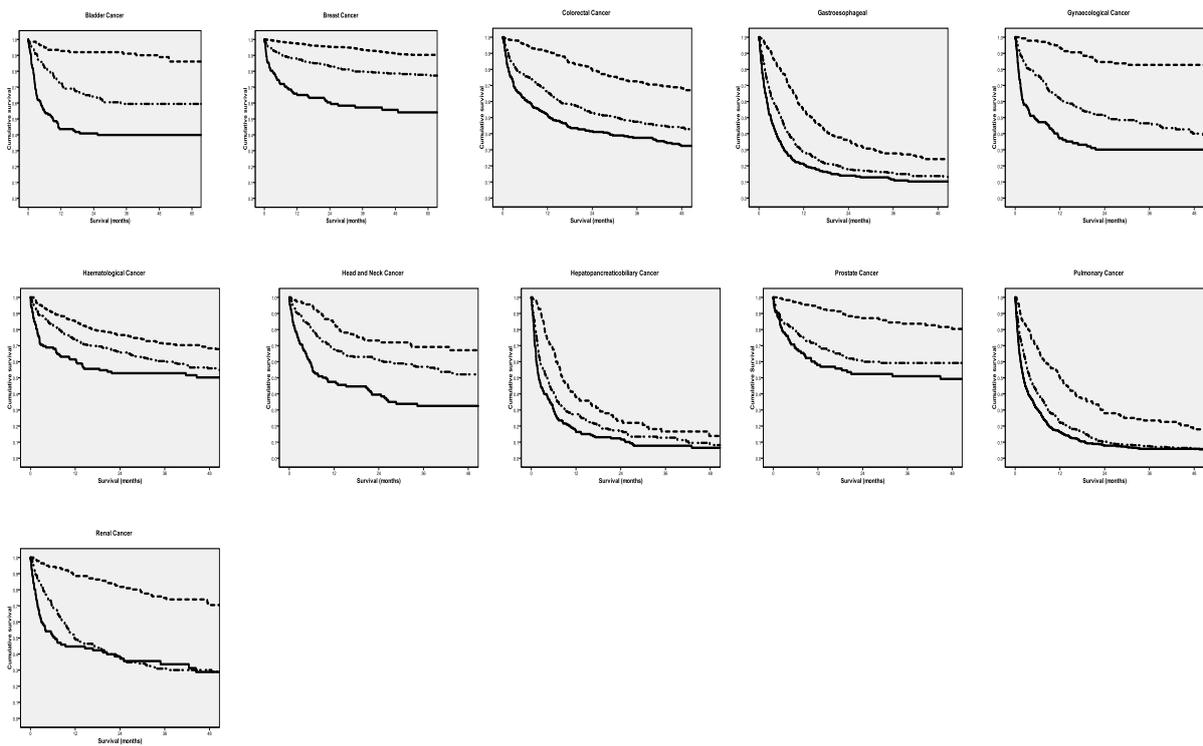
Neutrophil Lymphocyte Ratio



Platelet Lymphocyte Ratio



Prognostic Index



Prognostic Nutritional Index

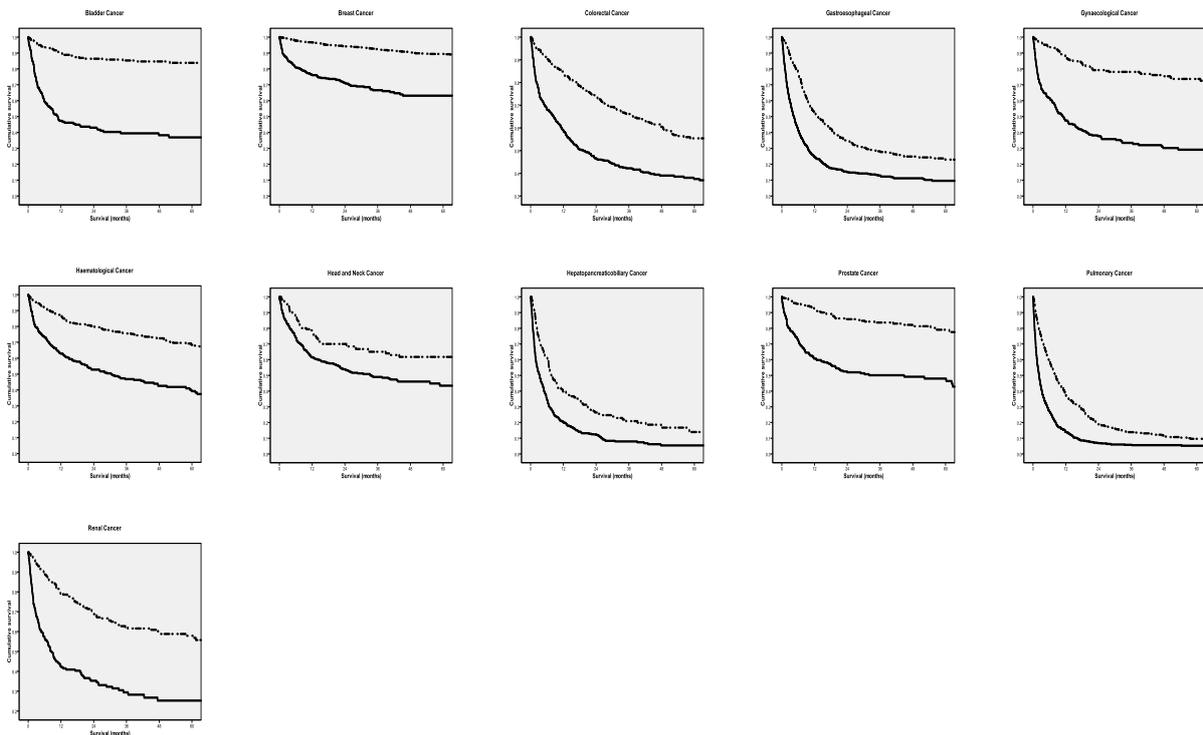


Figure 6.2.

The relationship between the mGPS, NLR, PLR, PI and PNI and survival illustrated by Kaplan–Meier survival plots in each individual tumour type with only those demonstrating statistical significance ($p < 0.001$) being shown.

For the modified Glasgow Prognostic Score in each plot, the top (small dash) line represents mGPS of 0, the middle (large dash) line represents a mGPS of 1 and the bottom (solid) line represents a mGPS of 2.

For the Neutrophil Lymphocyte Ratio in each plot, the top (large dash) line represents an NLR of 0 and the bottom (solid) line represents an NLR of 1.

For the Platelet Lymphocyte Ratio in each plot, the top (small dash) line represents PLR of 0, the middle (large dash) line represents a NLR of 1 and the bottom (solid) line represents a NLR of 2.

For the Prognostic Index in each plot, the top (small dash) line represents PNI of 0, the middle (large dash) line represents a PNI of 1 and the bottom (solid) line represents a PNI of 2.

For the Prognostic Nutritional Index in each plot, the top (large dash) line represents an PNI of 0 and the bottom (solid) line represents an NLR of 1.

7.0. A DERIVED NEUTROPHIL TO LYMPHOCYTE RATIO PREDICTS SURVIVAL IN PATIENTS WITH CANCER

7.1. ABSTRACT

Introduction: The Neutrophil Lymphocyte Ratio (NLR) has prognostic value in patients with a variety of cancers. Many chemotherapeutic trial databases hold information on white cell and neutrophil counts only. The aim of the present study was to compare the prognostic value of the NLR, with a derived score (dNLR), composed of white cell and neutrophil counts.

Methods: Patients (n=27,031) who were sampled incidentally between 2000 and 2007 for neutrophil, lymphocyte and white cell counts, and also had a diagnosis of cancer (Scottish Cancer Registry), were identified. Of this group 12,118 patients who had been sampled within two years of their cancer diagnosis were studied.

Results: On follow up, there were 7366 deaths, of which 6198 (84%) were cancer deaths. The area under the Receiver Operating Characteristic (ROC) curve for cancer specific survival was 0.650 for the NLR and 0.640 for the dNLR. The NLR and dNLR were independently associated with survival in all cancers studied (all $p < 0.001$).

Conclusion: The results of the present study show that the dNLR, based on white cell and neutrophil counts, has similar prognostic value to the NLR. Therefore, the universally available dNLR is to be commended for use in the risk stratification of patients undergoing chemotherapy.

7.2. INTRODUCTION

While it has been detailed that the host immune/inflammatory response plays an important role in carcinogenesis and disease progression there is increasing interest in simple, objective methods of measuring this response in patients with cancer. The Neutrophil Lymphocyte Ratio (NLR), as detailed in Chapter 3 and 6, has been well established as a prognostic score in patients with a wide variety of cancers including those undergoing chemotherapy for cancer [Kao et al., 2010b; Chua et al., 2011b]. Although, apparently inferior to other measures of the systemic immune/inflammatory response, such as the mGPS [Proctor et al., 2011a], the NLR does have the advantage of its components being inexpensive and routinely measured in day to day oncological practice and in current chemotherapeutic cancer trials. Clearly, if such extensive data were to confirm the prognostic value and clinical utility of the NLR this would be an important, relevant, clinical translational advance in the identification of cancer patients at high risk [Clarke et al., 2011].

However, on patient entry to chemotherapeutic trials, despite having a differential white cell count carried out, only white cell and neutrophil counts are routinely entered into the clinical trial databases. The differential white cell count is broadly composed of neutrophils (50-60%), lymphocyte (20-40%), monocytes (2-6%), eosinophils (1-4%) and basophils (0.5-1%) In an attempt to allow the widespread utilisation of a similar inflammation-based score in such settings, we aimed to investigate the prognostic value of a derived NLR (dNLR), from a white cell and neutrophil count. Therefore, the aim of the present study was to compare the prognostic value of the NLR and dNLR adjusted for age, sex, deprivation and tumour site in the Glasgow Inflammation Outcome Study.

7.3. MATERIALS AND METHODS

From a cohort previously described in Chapter 6 [Proctor et al., 2011a], patients in the North Glasgow who were sampled incidentally for C-reactive protein, albumin, calcium and a differential white cell count between the 1st January 2000 and the 31st December 2007 were included. This time period was as such to correspond with the available follow-up data from the Scottish Cancer Registry at the time of analysis as previously detailed. The NLR was constructed as follows: $NLR = \text{Neutrophil count} / \text{Lymphocyte count}$. The dNLR was constructed as follows: $dNLR = \text{Neutrophil Count} / (\text{White Cell Count} - \text{Neutrophil count})$.

Patients with blood samples taken within two years of their cancer diagnosis were included in the analysis, and split into those sampled prior to and following cancer diagnosis. The dNLR was derived from the assumption that the white cell count is made up primarily of lymphocytes and neutrophils, and therefore, the white cell count minus the neutrophil count would be broadly similar to the lymphocyte count. As different thresholds have been suggested in the past [Ding et al., 2010; Ohno et al., 2010; Kim et al., 2010; Sharaiha et al., 2011], several were examined in order to ascertain the optimal NLR and dNLR.

Cohort linkage, with the Scottish Cancer Registry, classification of deaths and cancer coding were carried out as described in Chapter 5 and only those patients who had complete Cancer Registry follow-up were included [Proctor et al., 2011b]. Survival was calculated as described in Chapter 5 [Proctor et al., 2011b]. Patient inclusion criteria has been previously detailed and only cancer groups previously studied were included [Proctor et al., 2011a]. Cancer groups with less than 150 cancer specific deaths were excluded to ensure statistical power. Cancer stage was extracted from the registry where available.

Deprivation was measured with the Scottish Index of Multiple Deprivation (SIMD) as detailed in Chapter 1. Overall patient numbers were different in this Chapter, when compared to some previous, as the inclusion time period differed to some and longer cancer registry follow-up was available (until June 2009). The study was approved by the Research Ethics Committee, North Glasgow NHS Trust.

Statistical analysis was carried out as follows. Kaplan-Meier estimator was used to analyse the relationship between patient characteristics, tumour site and overall and cancer specific survival (Table 7.1). Receiver Operating Characteristic (ROC) curve was used to determine the sensitivity and specificity similarities between the NLR and dNLR (Figure 7.1). Cox proportional multivariate regression analysis, corrected for age, sex, deprivation and tumour site, as well as area under the ROC curve, were used to determine the relationship between different NLR and dNLR thresholds (in whole numbers) and survival in patients sampled before and after diagnosis (Tables 7.2). Box plot was used to demonstrate the relationship between the NLR, dNLR and Dukes stage in patients with colorectal cancer (Figure 7.2). Cox proportional multivariate regression analysis, corrected for age, sex, deprivation and tumour site, as well as area under ROC curve, were used to determine the relationship between optimal NLR and dNLR thresholds and survival in patients with advanced (Dukes C and D) colorectal cancer (Table 7.3). Analysis was performed using SPSS software (SPSS Inc., Chicago, IL, USA).

7.4. RESULTS

From the Glasgow Inflammation Outcome Study cohort of 223,303 patients previously described in Chapter 4 [Proctor et al., 2010], 27,031 patients were identified as also having a diagnosis of cancer. Within this group, 12,119 patients had been sampled

within two years of their cancer diagnosis. Of this group, 3859 (32%) were sampled prior to a diagnosis and 8260 (68%) were sampled following diagnosis and the possible initiation of surgical or chemotherapeutic interventions.

The majority of patients were under the age of 75 years ($n=9097$, 75%), were female ($n=6217$, 51%) and were from the most deprived SIMD quintile ($n=4692$, 39%). The minimum follow-up from cancer diagnosis was 18 months and the maximum 132 months (median 52 months for survivors).

The relationship between patient characteristics, tumour site, and overall and cancer specific survival in all patients is shown in Table 7.1. In total 12,119 patients were studied. On follow-up, there were 7368 deaths of which 6200 (84%) were cancer related. The median time from blood sampling to diagnosis was 1.6 months in those sampled prior to diagnosis and 2.2 months in those sampled following diagnosis, suggesting that most scores reflect status at diagnosis. Increasing age, male gender, and increasing deprivation were associated with reduced 5 year overall and cancer specific survival (all $p<0.001$).

The ROC curves, using cancer specific death as an end-point ($n=12,119$), for NLR and dNLR is shown in Figure 7.1. The ROC curves for NLR and dNLR were 0.650 ($p<0.001$) and 0.640 ($p<0.001$) respectively. The Spearman rank correlation between the NLR and dNLR was 0.962 ($p<0.001$).

The relationship between NLR and dNLR thresholds and survival in patients sampled prior to and following diagnosis, adjusted for age, sex, deprivation and stratified by tumour site, is shown in Table 7.2. The optimal thresholds for the NLR ($>4:1$) and the dNLR ($>2:1$) in both patients sampled prior to and following diagnosis, as measured by the hazard ratios and area under the ROC curve, are highlighted.

In the present cohort, a limited number of patients had stage related data available with only colorectal cancer having stage data for over 50% of patients. Stage was only available in 29% (n=621) of patients with breast cancer and 10% (n=234) of patients with pulmonary cancer. Other cancer groups had no stage available. Of the colorectal cancer patients with stage available, 90 (9%) were Dukes A, 264 (27%) were Dukes B, 327 (33%) were Dukes C and 312 (31%) were Dukes D.

The relationship between the NLR, dNLR and Dukes stage in all patients is shown in Figure 7.2. The relationship between the NLR, dNLR and survival in patients with advanced colorectal cancer (Dukes C and D) adjusted for age, sex and deprivation is shown in Table 7.3. These patients were studied as they are likely to be similar to those entered into chemotherapeutic trials [Chua et al., 2011c]. On survival analysis (n=639), both the NLR and dNLR were associated with reduced overall and cancer specific survival independent of age, sex and deprivation (all $p < 0.001$) with similar Hazard Ratios and area under the ROC curve (both $p < 0.001$).

7.5. DISCUSSION

The results of the present Chapter show clearly that the NLR and dNLR have similar prognostic value and that they can be used similarly to predict survival in a large cohort of unselected cancer patients. Furthermore, the NLR and dNLR had similar predictive value, in all cancers as well as advanced colorectal cancer. Recently, Chua and co-workers reported that the NLR, as a marker of the systemic immune/inflammatory response, predicted clinically meaningful outcomes in patients with advanced colorectal cancer receiving chemotherapy [Chua et al., 2011a]. Taken together, these results would indicate that the derived NLR is suitable for the risk stratification of patients in chemotherapeutic trials, in particular, colorectal cancer studies.

In the present Chapter, although the ROC analysis of the NLR and dNLR were similar, the prognostic value of different thresholds was examined. It was of interest that, in contrast to the most commonly used NLR threshold of 5:1 [Walsh et al., 2005; Halazun et al., 2008; Gomez et al., 2008b], in the present Chapter 4:1 was found to have superior prognostic value in terms of Hazard Ratio and area under the ROC curve. These results are consistent with the varying NLR threshold reported across and within different tumour types. It is also of interest that the optimal threshold for the dNLR was 2:1, an expected shift from that of the NLR explained by the method of derivation. The results of the present Chapter also suggest that the dNLR has considerable potential to be adopted universally as a stratification factor in all current cancer clinical trials. Moreover, if it were shown to have such clinical utility, the dNLR would identify patients who may respond to anti-inflammatory interventions.

In the present Chapter it was of interest that there was a small but persistent superiority of the prognostic value of the NLR over the dNLR. The basis of this observation is not clear. However, in the dNLR the use of (WBC - neutrophil) as the denominator is broadly mixing two cell types, lymphocytes and monocytes, with possible opposing effects in terms of predictive value. In the normal range the relative proportion of lymphocytes to monocytes is approximately 6:1. In cancer patients there may be a fall in the absolute proportion of lymphocytes and an increase in the absolute proportion of monocytes. Even so their relative proportion is unlikely to fall below 3:1 even in advanced disease [Leitch et al., 2007]. Therefore (WBC – neutrophil) is dominated by lymphocytes and is likely to be a reasonable approximation to the lymphocyte fraction and the potential error introduced by the presence of monocytes in the fraction is therefore likely to be small. Given that different aspects of the differential white cell count have been reported to predict survival [Leitch et al., 2007; Proctor et al., 2011b; Lee et al., 2012] it is possible to derive other ratios such as

the neutrophil: white cell count ratio. However, of the differential white cell count parameters the neutrophil: lymphocyte ratio has been the most extensively validated and it was this that we were attempting to recapitulate when investigating the dNLR in the present Chapter. Clearly, where the NLR is available it should be used. However, there is a wealth of clinical trial data, where only white cell and neutrophil counts have been recorded in computer databases that could be used to examine the clinical value of the systemic immune/inflammatory response. As the present Chapter validates the use of the dNLR, this may help unlock the residual value of such clinical trial databases, and encourage the widespread utilisation of a similarly based systemic inflammation-based scores in such settings.

The results of the present Chapter also confirm the hypothesis that a total white cell count may be a useful addition to currently established prognostic markers of the systemic immune/inflammatory response such as, C-reactive protein, albumin, neutrophil and lymphocytes counts and their combinations [Proctor et al., 2011a]. It was also of interest that the hazard ratios and areas under the curves were greater from patients sampled after diagnosis, compared to patients sampled prior to diagnosis. These results would suggest that the systemic immune/inflammatory response is a more potent stimulator of cancer progression in established disease. This is consistent with the long standing observations on the “seed and soil” nature of cancer progression and metastasis [Fidler and Poste, 2008].

The present cohort has a number of limitations. The patients were selected on the presence of haematological and biochemical variables and were therefore not necessarily representative of all cancer patients in general. Patients may also have concurrent morbidity, including infection, causing alterations in their haematological variables. It is also recognised that demographic variables, such as race, that appear to influence neutrophil counts were not

available. Nevertheless, the optimal prognostic threshold for the NLR in the range of 4-5:1 has been consistently validated in different cancer cohorts (Clarke et al., 2011). It remains to be determined whether a dNLR of 2:1 will be similarly validated in different cancer cohorts.

In summary, the results of the present Chapter show that a derived NLR, based on a white cell and neutrophil counts, has similar prognostic value to the NLR. Therefore, the universally available dNLR is to be commended for use in the risk stratification of patients undergoing chemotherapy.

Table 7.1. The relationship between patient characteristics, tumour site, and overall and cancer specific survival in all patients

		n= 12,119 (%)	Overall survival 5 year % (n of deaths) n= 7368	p- value	Cancer survival 5 year % (n of deaths) n= 6200	p- value
Age	<65 years	5425 (45)	50 (2573)		54 (2316)	
	65–74 years	3672 (30)	32 (2412)		39 (2041)	
	≥ 75 years	3022 (25)	20 (2383)	<0.001	30 (1843)	<0.001
Sex	Male	5902 (49)	28 (4096)		35 (3437)	
	Female	6217 (51)	46 (3272)	<0.001	52 (2763)	<0.001
SIMD 2006	1 (least deprived)	1698 (14)	48 (860)		54 (732)	
	2	1486 (12)	45 (781)		51 (655)	
	3	1804 (15)	42 (1011)		48 (866)	
	4	2439 (20)	34 (1541)		42 (1284)	
	5 (most deprived)	4692 (39)	30 (3175)	<0.001	38 (2663)	<0.001
Tumour site	Breast	2147 (18)	78 (442)		84 (304)	
	Bladder	562 (5)	48 (286)		62 (195)	
	Gynaecological	639 (5)	47 (325)		53 (281)	
	Prostate	709 (6)	51 (323)		64 (222)	
	Gastroesophageal	1085 (9)	11 (932)		14 (851)	
	Haematological	1210 (10)	47 (627)		58 (456)	
	Renal	552 (4)	39 (324)		44 (285)	
	Colorectal	1413 (12)	39 (820)		46 (680)	
	Head and Neck	738 (6)	35 (452)		52 (296)	
	Hepatopancreaticobiliary	721 (6)	6 (660)		7 (626)	
	Pulmonary	2343 (19)	6 (2177)	<0.001	8 (2004)	<0.001

Table 7.2. The relationship between NLR and dNLR thresholds and survival. Adjusted for age, sex, deprivation and stratified by tumour site

			Overall survival			Cancer survival		
			HR	p-value	AUC	HR	p-value	AUC
Patient sampled prior to cancer diagnosis n=3859								
NLR	<1:1	78 (2)	1			1		
	>1:1	3781 (98)	1.03	0.840	0.507	0.967	0.859	0.507
	<2:1	589 (15)	1			1		
	>2:1	3270 (85)	1.49	<0.001	0.563	1.48	<0.001	0.547
	<3:1	1354 (35)	1			1		
	>3:1	2505 (65)	1.55	<0.001	0.593	1.52	<0.001	0.560
	<4:1	1987 (51)	1			1		
	>4:1	1872 (49)	1.57	<0.001	0.598	1.52	<0.001	0.558
	<5:1	2435 (63)	1			1		
	>5:1	1424 (37)	1.50	<0.001	0.579	1.44	<0.001	0.542
dNLR	<1:1	214 (6)	1			1		
	>1:1	3645 (94)	1.17	0.098	0.518	1.16	0.163	0.516
	<2:1	1399 (36)	1			1		
	>2:1	2460 (64)	1.54	<0.001	0.593	1.53	<0.001	0.563
	<3:1	2461 (64)	1			1		
	>3:1	1398 (36)	1.47	<0.001	0.575	1.43	<0.001	0.543
	<4:1	2960 (77)	1			1		
	>4:1	899 (23)	1.46	<0.001	0.552	1.41	<0.001	0.527
	<5:1	3267 (85)	1			1		
	>5:1	592 (15)	1.40	<0.001	0.538	1.33	<0.001	0.517
Patient sampled following cancer diagnosis n=8260								
NLR	<1:1	297 (4)	1			1		
	>1:1	7963 (96)	1.16	0.094	0.505	1.89	0.069	0.506
	<2:1	1480 (18)	1			1		
	>2:1	6779 (82)	1.63	<0.001	0.575	1.67	<0.001	0.566
	<3:1	2984 (36)	1			1		
	>3:1	5275 (64)	1.85	<0.001	0.641	1.93	<0.001	0.629
	<4:1	4104 (50)	1			1		
	>4:1	4155 (50)	1.86	<0.001	0.661	1.92	<0.001	0.646
	<5:1	4872 (59)	1			1		
	>5:1	3387 (41)	1.82	<0.001	0.657	1.86	<0.001	0.642
dNLR	<1:1	656 (8)	1			1		
	>1:1	7604 (92)	1.35	<0.001	0.521	1.39	<0.001	0.519
	<2:1	3083 (37)	1			1		
	>2:1	5176 (63)	1.76	<0.001	0.630	1.83	<0.001	0.620
	<3:1	4978 (60)	1			1		
	>3:1	3281 (40)	1.74	<0.001	0.644	1.78	<0.001	0.631
	<4:1	6034 (73)	1			1		
	>4:1	2225 (27)	1.75	<0.001	0.619	1.78	<0.001	0.609
	<5:1	6632 (80)	1			1		
	>5:1	1627 (20)	1.76	<0.001	0.595	1.80	<0.001	0.589

Table 7.3. The relationship between the NLR, dNLR and survival in patients with advanced colorectal cancer (Dukes C and D). Adjusted for age, sex and deprivation.

		Overall survival			Cancer specific survival			
		n = 639 (%)	HR	p-value	AUC	HR	p-value	AUC
NLR	<4:1	278 (44)	1			1		
	>4:1	361 (56)	1.60	<0.001	0.584	1.60	<0.001	0.565
dNLR	<2:1	194 (30)	1			1		
	>2:1	445 (70)	1.61	<0.001	0.575	1.67	<0.001	0.566

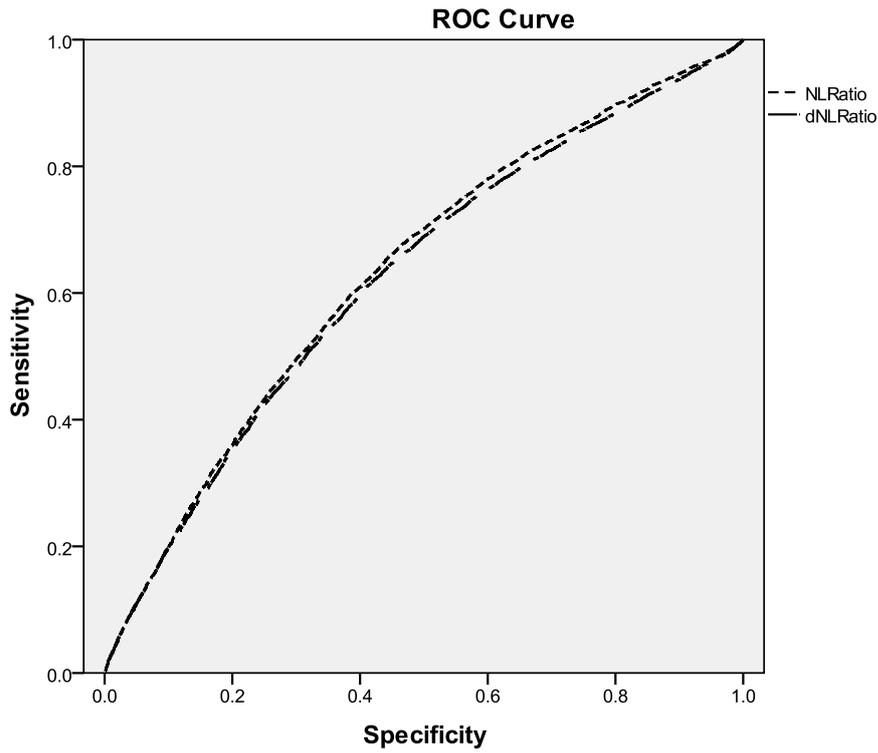


Figure 7.1. Receiver Operating Characteristic curve for cancer specific survival.

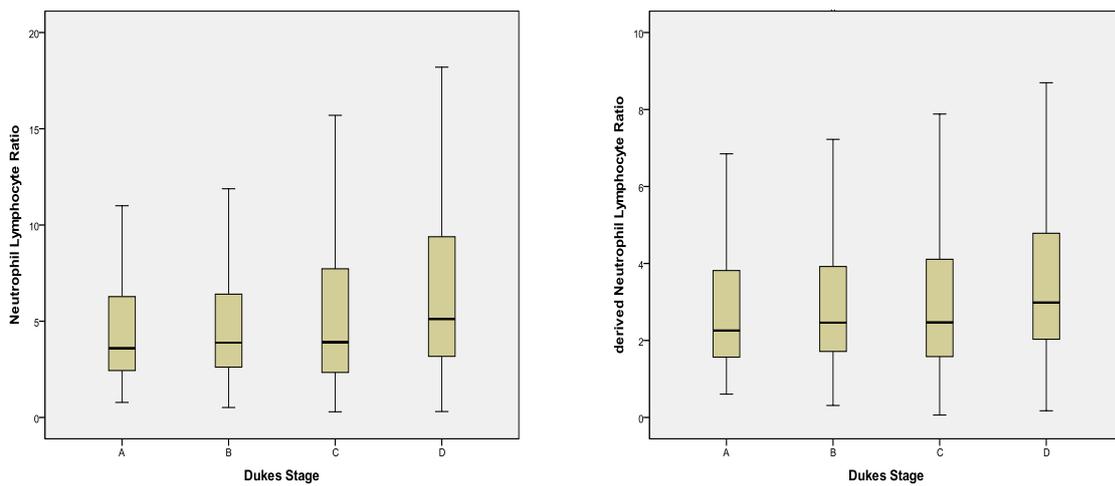


Figure 7.2. The relationship between NLR, dNLR and Dukes stage in patients with colorectal cancer.

CHAPTER 8. OPTIMISATION OF THE SYSTEMIC INFLAMMATION-BASED GLASGOW PROGNOSTIC SCORE IN PATIENTS WITH CANCER (oGPS)

8.1. ABSTRACT

Introduction: The modified Glasgow Prognostic Score (mGPS), an inflammation-based prognostic score that uses thresholds of C-reactive protein ($>10\text{mg/l}$) and albumin ($<35\text{g/l}$), is independently prognostic of survival in cancer. The aim of the present chapter was to establish whether the addition of a differential white cell count and a high sensitivity C-reactive protein measurement enhances the prognostic value of the mGPS.

Methods: Patients ($n=12,119$) who had an incidental blood sample taken between 2000 and 2007 for C-reactive protein, albumin and a differential white cell count, as well as a diagnosis of cancer made within two years were identified. This group were studied for the prognostic value of neutrophil, lymphocyte, and platelet counts. Also, patients ($n=2743$) sampled following the introduction of high sensitivity C-reactive protein measurements were studied for the prognostic value of different thresholds.

Results: The prognostic value of the mGPS (HR 2.61, $p<0.001$, AUC 0.715), using cancer specific survival as an end-point, was improved by the addition of neutrophil and platelet counts (HR 4.86, $p<0.001$, AUC 0.734) and a high sensitivity C-reactive protein measurement ($>3\text{mg/l}$, HR 5.77, $p<0.001$, AUC 0.734).

Conclusion: The results of the present chapter showed that the addition of neutrophil and platelet counts, as well as a high sensitivity C-reactive protein measurement, enhanced the prognostic value of the Glasgow Prognostic Score.

8.2. INTRODUCTION

As has been detailed in previous Chapters, the host immune/inflammatory response plays an important role in carcinogenesis and disease progression. It has also been demonstrated that a range of inflammation-based scores have prognostic value in patients with cancer [Proctor et al., 2011a]. Moreover, it appeared that the inclusion of a white cell count, may add prognostic value to the validated mGPS [Proctor et al., 2011a]. Also, the recent introduction of high sensitivity C-reactive protein measurements in routine clinical laboratory analysis has resulted in the threshold sensitivity being lowered from approximately 5mg/l to 0.05mg/l. This has raised the possibility that the combination of C-reactive protein and albumin may be further modified to take account of the potential additional prognostic value that could be derived from the components of a differential white cell count (neutrophil, lymphocyte, platelet counts) and high sensitivity C-reactive protein measurement.

The aim of the present Chapter was to establish whether the addition of a differential white cell count or a high sensitivity C-reactive protein measurement could enhance the prognostic value of the Glasgow Prognostic Score.

8.3. MATERIALS AND METHODS

From a cohort previously described in Chapter 6 [Proctor et al., 2011a], patients in North Glasgow who were sampled incidentally for C-reactive protein, albumin, calcium and a differential white cell count between the 1st January 2000 and the 31st December 2007 were included. This time period was as such to correspond with the available follow-up data from the Scottish Cancer Registry at the time of analysis. Prior to the introduction of high-sensitivity C-reactive protein measurements, the limit of detection was a concentration of less

than 5mg/L and 0.05mg/l thereafter. Only those patients undergoing blood sampling after the introduction of the routine measurement of high sensitivity C-reactive protein (1st of August 2006), were included in the analysis investigating the prognostic value of high sensitivity thresholds (<5mg/l). Standardised thresholds for white cell count ($>11 \times 10^9/l$), neutrophil count ($>7.5 \times 10^9/l$), lymphocyte count ($>3 \times 10^9/l$) and platelet count ($>400 \times 10^9/l$) were used. High sensitivity C-reactive protein thresholds, <1mg/l and <3mg/l, previously reported to be associated with survival were investigated [Ridker, 2008]. The validated GPS and mGPS were constructed as described in Table 3.1. Patients with blood samples taken within two years of their cancer diagnosis were included in the analysis, and split into those sampled prior to and following cancer diagnosis. Cohort linkage, with the Scottish Cancer Registry, classification of deaths and cancer coding were carried out as described in Chapter 5 and only those patients who had complete Cancer Registry follow-up were included [Proctor et al., 2011b]. Survival was calculated as described in Chapter 5 [Proctor et al., 2011b]. Cancer groups with less than 150 cancer specific deaths were excluded to ensure statistical power. Cancer stage was extracted from the registry where available.

Deprivation was measured with the Scottish Index of Multiple Deprivation (SIMD) as detailed in Chapter 1 [Proctor et al., 2010]. Patient inclusion criteria has been previously detailed and only cancer groups previously studied were included [Proctor et al., 2011a]. Overall patient numbers were different in this Chapter, when compared to previous, as the inclusion time period differed to some and longer cancer registry follow-up was available (until July 2009). The study was approved by the Research Ethics Committee, North Glasgow NHS Trust.

Statistical analysis was carried out as follows. The relationship between patient demographics, tumour site, markers of the systemic immune/inflammatory response and

survival was assessed using the Kaplan-Meier and Cox proportional hazards methods and Area Under the receiver operating characteristic Curve (AUC). Cox proportional hazards model stratified by tumour site, adjusted for age, sex, deprivation, time of sample and stratified by tumour site, and AUC was used to analyse the relationship between the components of a differential white cell count and high sensitivity C-reactive protein and survival. In those patients with a high sensitivity C-reactive protein measurement, the prognostic value of the addition of the components of a differential white cell count and high sensitivity C-reactive protein to the GPS was examined with Cox proportional hazards model stratified for age, sex, deprivation, time of sample and stratified by tumour and the AUC. Given the number of statistical comparisons, a p value of <0.01 was considered to have statistical significance in the analysis. Analysis was performed using SPSS software (SPSS, Chicago, IL, USA).

8.4. RESULTS

From GIOS cohort of 223,303 patients originally described [Proctor et al., 2010], 27,031 patients were identified as having a diagnosis of cancer in SCR and a blood sample for C-reactive protein, albumin and a differential white cell count taken between January 2000 and December 2007. Within this group, 12,119 patients had been sampled within two years of their cancer diagnosis, 3859 (32%) sampled prior to and 8260 (68%) following. Most patients were under 75 years of age (n=9097, 75%) and female (n=6217, 51%). Thirty nine percent of patients (n=4692) were SIMD 5.

The relationship between patient demographics, tumour site, markers of systemic inflammation and five year survival in all patients (n=12,119) is shown in Table 8.1. There were 7368 deaths in total with 6200 being cancer specific. The median time from blood

sample to diagnosis was 2 months, suggesting that most scores reflect status at diagnosis. The minimum follow-up from cancer diagnosis for all patients was 18 months and the maximum 138 months (median 52 months for survivors). Increasing age, male gender, and increasing deprivation were associated with reduced 5 year overall and cancer specific survival (all $p < 0.001$). High C-reactive protein ($>10\text{mg/l}$), white cell, neutrophil and platelet count and low albumin and lymphocyte count were associated with a reduced 5 year overall and cancer specific survival (all $p < 0.001$).

The relationship between patient demographics, standard threshold C-reactive protein measurement ($>10\text{mg/l}$), albumin, white cell, lymphocyte, neutrophil and platelet count and survival, stratified by tumour site and time of sample, is shown in Table 8.2. Increasing age, sample prior to diagnosis as well as high C-reactive protein ($>10\text{mg/l}$), neutrophil and platelet count, as well as low albumin were all independently associated with a reduction in overall and cancer specific survival (all $p < 0.01$).

When the relationship between high sensitivity C-reactive protein thresholds and survival was investigated in patients sampled following the routine introduction of high sensitivity measurements ($n=2742$), no statistical difference between a high sensitivity C-reactive protein concentration of $<1\text{mg/L}$ and $1\text{-}3\text{mg/L}$ was found. A threshold of $>3\text{mg/L}$ was therefore used.

The relationship between patient demographics, high sensitivity C-reactive protein measurements ($>3\text{mg/l}$), albumin, white cell, lymphocyte, neutrophil and platelet counts and survival ($n=2743$): stratified by tumour site in only those patients sampled following the routine introduction of high sensitivity measurements is shown in Table 8.3. Increasing age, sample prior to diagnosis as well as high C-reactive protein ($>3\text{mg/l}$), neutrophil and platelet

count, as well as low albumin were all independently associated with a reduction in overall and cancer specific survival (all $p < 0.01$).

The hazard ratios and the areas under the receiver operator characteristic (ROC) curve in patients sampled following the introduction of high sensitivity C-reactive protein measurements ($n=2743$), adjusted for age, sex, deprivation, tumour site and time of sampling, for a high sensitivity C-reactive protein threshold of $>3\text{mg/l}$ for overall and cancer specific mortality were calculated. These were HR 2.47 (95% CI 2.08 - 2.94), AUC 0.636 ($p < 0.001$) and HR 2.64 (95% CI 2.19-3.18), AUC 0.625 ($p < 0.001$) respectively compared with the HR 2.13 (95% CI 1.89-2.40), AUC 0.676 for overall ($p < 0.001$) and cancer specific HR 2.22 (95% CI 1.96-2.52), AUC 0.665 ($p < 0.001$) mortality using the standard threshold of 10mg/l . Therefore, since they were similar, when optimising the GPS, both the highly sensitive and standard threshold for C-reactive protein (10mg/l) were taken forward with other measures of the systemic immune/inflammatory response.

When the interrelationships between inflammatory markers in patients sampled following the introduction of high sensitivity C-reactive protein, were explored ($n=2743$), C-reactive protein was inversely associated with albumin ($r_s=0.691$, $p < 0.001$), lymphocyte count ($r_s=0.334$, $p < 0.001$) and directly associated with white cell count ($r_s=0.334$, $p < 0.001$), neutrophil count ($r_s=0.399$, $p < 0.001$) and platelet count ($r_s=0.172$, $p < 0.001$). Albumin was inversely associated with white cell count ($r_s=0.268$, $p < 0.001$), neutrophil count ($r_s=0.352$, $p < 0.001$), platelet count ($r_s=0.145$, $p < 0.001$) and directly associated with lymphocyte count ($r_s=0.332$, $p < 0.001$). White cell count was directly associated with lymphocyte count ($r_s=0.0187$, $p < 0.001$), neutrophil count ($r_s=0.951$, $p < 0.001$) and platelet count ($r_s=0.346$, $p < 0.001$). Lymphocyte count was inversely associated with neutrophil count ($r_s=0.049$,

p<0.05) and directly associated with platelet count (rs=0.179, p<0.001). Neutrophil count was directly associated with platelet count (rs=0.335, p<0.001).

The optimisation of the Glasgow Prognostic Score in patients sampled following the introduction of high sensitivity C-reactive protein is shown in Table 8.4 (n=2743). All combinations of inflammatory markers were predictive of overall and cancer specific survival (all p<0.001) independent of age, sex, time of sample and tumour site. A score combining C-reactive protein, albumin, and neutrophil count had an improved prognostic value, in terms of increased hazard ratio and area under the ROC curve, when compared to the GPS or mGPS. Further addition of a high sensitivity C-reactive improved the hazard ratio range, and thus offers the potential to better differentiate between those who will do well and those that will do badly. However, it did not increase the area under the ROC curve and therefore did not improve its overall prognostic value.

The relationship between the Glasgow Prognostic Scores and survival in patients studied following the introduction of high sensitivity C-reactive protein, is shown in Figure 8.1 (n=2743).

8.5. DISCUSSION

The results of the present Chapter show that, in a large cohort (n=12,119) of patients with cancer, routine objective markers of the systemic immune/inflammatory response, elevated neutrophil and platelet counts, enhance the prognostic value of the Glasgow Prognostic Score independent of age, sex, deprivation, tumour site and time of sampling. In a selected cohort of patients (n=2743) with high sensitivity C-reactive protein measurements, a lower threshold for C-reactive protein was associated with a larger range of hazard ratios, thus potential better differentiation of patients who will do particularly well, but no

improvement in the overall AUC of the Glasgow Prognostic Score, when combined with neutrophil and platelet counts.

Taken together, these additions to the GPS, have led to an increase in the hazard ratios associated with overall and cancer specific survival by approximately 100%. Therefore, the refinement of the GPS with other components of the systemic immune/inflammatory response appears to have improved the prognostic value and clinical utility of the Glasgow Prognostic Score.

It is of interest that in the present Chapter of patients with cancer, it was identified that, high sensitivity C-reactive protein measurements $>3\text{mg/l}$ were associated with reduced survival. This is consistent with recent publications in cancer cohorts using the same threshold [Koenig et al., 2008; Ito et al., 2011; Nakamura et al., 2012]. The results of the present Chapter are also in agreement with previous workers [Gagnon et al., 2010] who reported, in patients with non-small cell lung cancer, that while an increase risk of mortality occurs with a C-reactive protein measurement above approximately 4mg/l , the risk increase is steeper between 8 and 16 mg/l . Indeed, some workers have adopted other thresholds for C-reactive protein such as $>5\text{mg/l}$ into the GPS [Toiyama et al., 2011].

The same high sensitivity C-reactive protein threshold ($>3\text{mg/l}$) has also been used to predict survival in non-cancer cohorts [Tuomisto et al., 2006; Zacho et al., 2010]. Taken together, the results would suggest that a C-reactive protein of 3mg/l is a potentially important threshold in both cancer and non-cancer cohorts. It raises the possibility that inflammation-based prognostic scores such as those examined in the present Chapter will have prognostic value in non-cancer cohorts. Further work in large non-cancer cohorts is required to examine the hypothesis that the Glasgow Prognostic Score, in combination with

neutrophil and platelet counts, as well as high sensitivity C-reactive protein, also independently predicts non-cancer specific survival.

Over the last decade, with the validation and establishment of simple, objective systemic inflammation-based prognostic scores it has become recognised that activation of the systemic immune/inflammatory response has a detrimental impact on the outcome of cancer patients [Roxburgh and McMillan, 2010]. More recently, there has been renewed interest on the beneficial impact of an activated local immune/inflammatory response on the outcome of cancer patients [Pinato et al., 2012]. Therefore, the relationship between the systemic and local immune/inflammatory responses is likely to be of considerable and increasing interest in future years.

The present cohort has a number of limitations. The patients were selected on the basis that measurements of C-reactive protein, albumin and a differential white cell count had been performed and were therefore not necessarily representative of all cancer patients diagnosed and treated in the North Glasgow area. It is also recognised that patients with cancer may have concurrent morbidity causing a rise in their C-reactive protein and derangement of their albumin and other haematological parameters.

In summary, the addition of the constituents of a differential white cell count and high sensitivity C-reactive protein measurements can be used to improved the prognostic value and clinical utility of the Glasgow Prognostic Score. This optimised Glasgow Prognostic Score can be adopted to predict cancer survival where differential white cell counts and high sensitivity C-reactive protein measurements are available.

Table 8.1. The relationship between patient demographics, tumour site, inflammatory markers and survival in all patients

n=12,119			Patients n=12 119 (%)	Overall survival 5 year % (n of deaths) n= 7368	p- value	Cancer survival 5 year % (n of deaths) n= 6200	p- value
Age	<65 years		5425 (45)	50 (2573)		54 (2316)	
	65–74 years		3672 (30)	32 (2412)		39 (2041)	
	≥ 75 years		3022 (25)	20 (2383)	<0.001	30 (1843)	<0.001
Sex	Male		5902 (49)	28 (4096)		35 (3437)	
	Female		6217 (51)	46 (3272)	<0.001	52 (2763)	<0.001
SIMD 2006	1 (least deprived)		1698 (14)	48 (860)		54 (732)	
	2		1486 (12)	45 (781)		51 (655)	
	3		1804 (15)	42 (1011)		48 (866)	
	4		2439 (20)	34 (1541)		42 (1284)	
	5 (most deprived)		4692 (39)	30 (3175)	<0.001	38 (2663)	<0.001
Tumour site	Breast		2147 (18)	78 (442)		84 (304)	
	Bladder		562 (5)	48 (286)		62 (195)	
	Gynaecological		639 (5)	47 (325)		53 (281)	
	Prostate		709 (6)	51 (323)		64 (222)	
	Gastroesophageal		1085 (9)	11 (932)		14 (851)	
	Haematological		1210 (10)	47 (627)		58 (456)	
	Renal		552 (4)	39 (324)		44 (285)	
	Colorectal		1413 (12)	39 (820)		46 (680)	
	Head and Neck		738 (6)	35 (452)		52 (296)	
	Hepatopancreaticobiliary		721 (6)	6 (660)		7 (626)	
Inflammatory markers	C-reactive protein	<10mg/l	5146 (42)	56 (2143)		63 (1708)	
		>10mg/l	6973 (58)	23 (5225)	<0.001	30 (4492)	<0.001
	Albumin	>35g/l	8354 (69)	46 (4388)		53 (3627)	
		<35g/l	3765 (31)	17 (2980)	<0.001	23 (2573)	<0.001
	White cell count	<11×10 ⁹ /l	9104 (75)	42 (5043)		49 (4213)	
		>11×10 ⁹ /l	3015 (25)	21 (2325)	<0.001	28 (1987)	<0.001
	Neutrophil count	<7.5×10 ⁹ /l	8334 (69)	44 (4444)		51 (3699)	
		>7.5×10 ⁹ /l	3785 (31)	21 (2924)	<0.001	28 (2501)	<0.001
	Lymphocyte count	<3×10 ⁹ /l	11 554 (95)	37 (7071)		44 (5957)	
		>3×10 ⁹ /l	565 (5)	44 (297)	<0.001	51 (243)	<0.001
Platelet count	<400×10 ⁹ /l	10 254 (85)	40 (5889)		48 (4884)		
	>400×10 ⁹ /l	1865 (15)	19 (1479)	<0.001	24 (1316)	<0.001	

Table 8.2. The relationship between patient demographics, C-reactive protein (>10mg/l), albumin, differential white cell count and survival. Stratified by tumour site

n=12,119		Overall survival					Cancer specific survival				
		Univariate		Multivariate		AUC	Univariate		Multivariate		AUC
		HR	p-value	HR	p-value		HR	p-value	HR	p-value	
Age	<65/ 65-74 >75	1 1.37 2.09	 <0.001 <0.001	1 1.28 1.93	 <0.001 <0.001		1 1.24 1.73	 <0.001 <0.001	1 1.17 1.60	 <0.001	
Sex	Male Female	1 0.96	 0.101				1 0.97	 0.268			
SIMD	1 2 3 4 5	1 1.04 1.05 1.16 1.23	 0.343 0.226 <0.001 <0.001				1 1.02 1.04 1.11 1.17	 0.735 0.471 0.029 <0.001			
Time of sample	Pre diag Post diag	1 0.73	 <0.001	1 0.67	 <0.001		1 0.78	 <0.001	1 0.70	 <0.001	
C-reactive protein	<10mg/l >10mg/l	1 1.94	 <0.001	1 1.53	 <0.001	0.671	1 2.01	 <0.001	1 1.61	 <0.001	0.653
Albumin	>35mg/l <35mg/l	1 1.82	 <0.001	1 1.55	 <0.001	0.620	1 1.82	 <0.001	1 1.53	 <0.001	0.607
White cell count	<11×10 ⁹ /l >11×10 ⁹ /l	1 1.49	 <0.001	1 1.06	 0.166		1 1.48	 <0.001	1 1.05	 0.237	
Lymphocyte count	<3×10 ⁹ /l >3×10 ⁹ /l	1 0.77	 <0.001	1 0.86	 0.012		1 0.75	 <0.001	1 0.84	 0.010	
Neutrophil count	<7.5×10 ⁹ /l >7.5×10 ⁹ /l	1 1.59	 <0.001	1 1.28	 <0.001	0.608	1 1.58	 <0.001	1 1.26	 <0.001	0.593
Platelet count	<400×10 ⁹ /l >400×10 ⁹ /l	1 1.37	 <0.001	1 1.15	 <0.001	0.560	1 1.41	 <0.001	1 1.17	 <0.001	0.560

Table 8.3. The relationship between patient demographics, high sensitivity C-reactive protein (>3mg/l), albumin, differential white cell count and survival. Stratified by tumour site

n=2743		Overall survival				AUC	Cancer specific survival				
		Univariate HR	p-value	Multivariate HR	p-value		Univariate HR	p-value	Multivariate HR	p-value	AUC
Age	<65/	1		1			1		1		
	65-74	1.20	0.003	1.11	0.104		1.12	0.085	1.03	0.657	
	>75	1.73	<0.001	1.55	<0.001		1.50	<0.001	1.34	<0.001	
Sex	Male	1					1				
	Female	1.04	0.554				1.04	0.547			
SIMD	1	1					1				
	2	0.94	0.558				0.94	0.576			
	3	1.07	0.513				1.08	0.490			
	4	1.15	0.148				1.14	0.205			
	5	1.21	0.034				1.17	0.093			
Time of sample	Pre Diag	1		1			1		1		
	Post Diag	0.86	<0.001	0.62	<0.001		0.87	<0.001	0.63	<0.001	
C-reactive protein	<3mg/l	1		1			1		1		
	>3mg/l	2.54	<0.001	1.63	<0.001	0.636	2.68	<0.001	1.72	<0.001	0.625
Albumin	>35mg/l	1		1			1		1		
	<35mg/l	2.37	<0.001	1.97	<0.001	0.700	2.42	<0.001	2.01	<0.001	0.685
White cell count	<11×10 ⁹ /l	1		1			1		1		
	>11×10 ⁹ /l	1.65	<0.001	1.04	0.672		1.65	<0.001	1.03	0.748	
Neutrophil count	<7.5×10 ⁹ /l	1		1			1		1		
	>7.5×10 ⁹ /l	1.78	<0.001	1.39	<0.001	0.621	1.76	<0.001	1.35	<0.001	0.609
Lymphocyte count	<3×10 ⁹ /l	1					1				
	>3×10 ⁹ /l	0.76	0.043				0.69	0.012			
Platelet count	<400×10 ⁹ /l	1		1			1		1		
	>400×10 ⁹ /l	1.58	<0.001	1.26	0.001	0.575	1.68	<0.001	1.31	<0.001	0.578

Table 8.4. Optimisation of the Glasgow Prognostic Score in patients sampled following the introduction of high sensitivity C-reactive protein measurement. Survival adjusted for age, sex, time of sample and stratified by tumour site

n=2743				Cancer specific survival			Overall survival		
				HR	p-value	AUC	HR	p-value	AUC
GPS	0	955 (35)		1			1		
	1	621 (23)		1.86			1.88		
	2	1166 (42)		3.18	<0.001	0.731	3.33	<0.001	0.715
mGPS	0	1204 (44)		1			1		
	1	372 (14)		1.30			1.32		
	2	1166 (42)		2.49	<0.001	0.707	2.61	<0.001	0.695
CRP > 10mg/l +Albumin >35g/L +Neutrophils >7.5×10 ⁹ /l	0	847 (30.9)		1			1		
	1	537 (19.6)		1.64			1.70		
	2	790 (28.8)		2.65			2.80		
	3	568 (20.7)		3.96	<0.001	0.746	4.12	<0.001	0.726
CRP > 10mg/l + Albumin >35g/L + Neutrophils >7.5×10 ⁹ /l + Platelets >400×10 ⁹ /l	0	813 (29.6)		1			1		
	1	513 (18.7)		1.59			1.65		
	2	706 (25.7)		2.49			2.63		
	3	505 (18.4)		3.59			3.79		
	4	205 (7.5)		4.58	<0.001	0.751	4.86	<0.001	0.734
hs-CRP >3mg/l + Albumin >35g/L + Neutrophils >7.5×10 ⁹ /l + Platelets >400×10 ⁹ /l	0	508 (18.5)		1			1		
	1	635 (23.2)		1.61			1.62		
	2	815 (29.7)		2.72			2.88		
	3	571 (20.8)		4.14			4.32		
	4	213 (7.8)		5.46	<0.001	0.751	5.77	<0.001	0.734

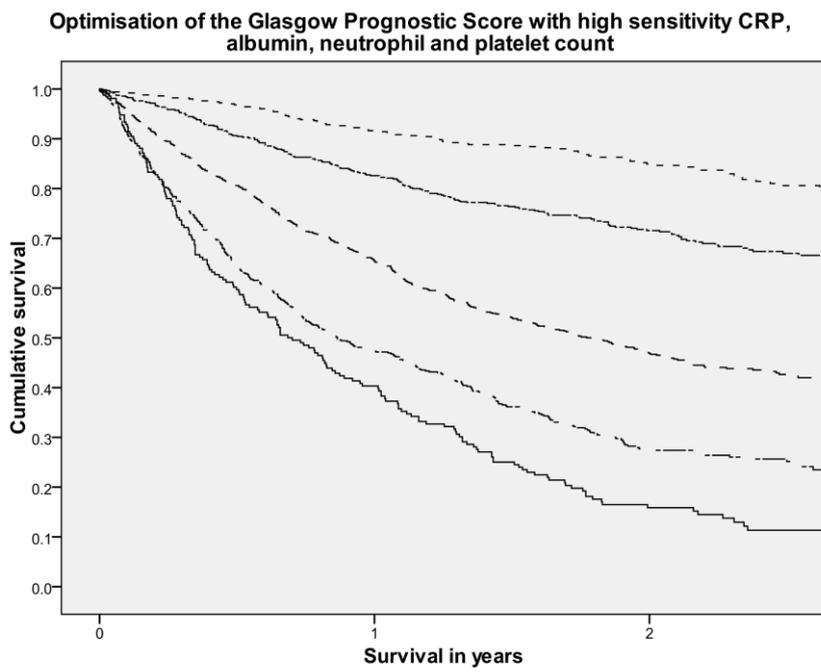
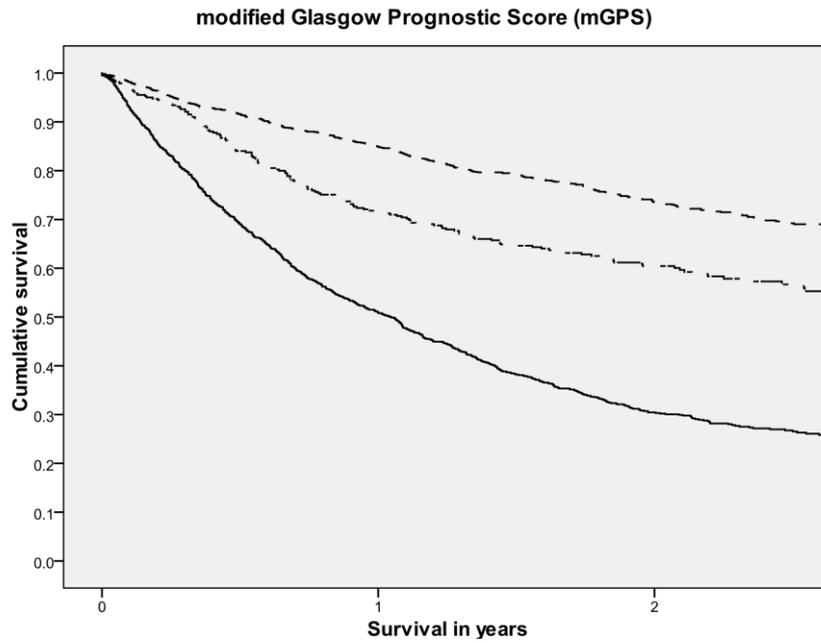


Figure 8.1. The relationship between the mGPS (0-top, small dash line; 1-middle, large dash line; 2-bottom, solid line) and the optimisation of the Glasgow Prognostic Score with high sensitivity CRP, albumin, neutrophil and platelet count (0-top, small dash line; 1-upper dot and dash line, 2-middle large dash line, 3-lower separated dot and dash line, 4-bottom, solid line) and survival.

CHAPTER 9. SYSTEMIC INFLAMMATION-BASED SCORE PREDICTS ALL-CAUSE, CARDIOVASCULAR AND CEREBROVASCULAR MORTALITY

9.1. ABSTRACT

Introduction: Markers of the systemic inflammatory response have been shown to be prognostic of survival in patients with cancer. The aim of the present chapter was to examine the prognostic relationship between C-reactive protein, albumin, neutrophil, lymphocyte and platelet counts with all-cause, cancer, cardiovascular and cerebrovascular mortality.

Methods: Patients (n=160,481) who had an incidental blood sample taken between 2000 and 2008 were studied for the prognostic value of C-reactive protein (>10mg/l), albumin, neutrophil, lymphocyte and platelet counts. Also, patients (n=52,091) sampled following the introduction of high sensitivity C-reactive protein (>3mg/l) measurements were studied. A combination of these markers were used to make cumulative inflammation-based scores.

Results: In all patients (n=160,481) C-reactive protein (>10mg/l) (HR 2.71, p<0.001), albumin (>35mg/l) (HR 3.68, p<0.001) and neutrophil counts (HR 2.18, p<0.001) were independently predictive of all-cause mortality. These associations were also observed in cancer, cardiovascular and cerebrovascular mortality before and after the introduction of high sensitivity C-reactive protein measurements (>3mg/l) (n=52,091). A combination of high sensitivity C-reactive protein (>3mg/l), albumin and neutrophil count predicted all-cause (HR 7.37, p<0.001, AUC 0.723), cancer (HR 9.32, p<0.001, AUC 0.731), cardiovascular (HR 4.03, p<0.001, AUC 0.650) and cerebrovascular (HR 3.10, p<0.001, AUC 0.623) mortality.

Conclusion: The results of the present study showed that an inflammation-based prognostic score, combining high sensitivity C-reactive protein, albumin and neutrophil counts is prognostic of all-cause mortality.

9.2. INTRODUCTION

As has been detailed in previous Chapters, there is now good evidence that inflammation based prognostic scores have independent prognostic value in patients with a variety of cancers. In Chapter 8, constituents of these scores, including high-sensitivity C-reactive protein ($>3\text{mg/l}$), albumin ($<35\text{g/l}$), neutrophils ($>7.5 \times 10^9$) and platelets ($>400 \times 10^9$), were combined to form the optimised Glasgow Prognostic Score (oGPS) that improved the prognostic value of the established Glasgow Prognostic Score [Proctor et al., 2013]. Furthermore, Chapter 1 detailed the well established relationship between certain markers of systemic inflammation with atherosclerotic and all-cause mortality, with more recent studies demonstrating similar correlations [Zacho et al., 2010; Phillips et al., 2011; Zamani et al., 2013]. However, it remains to be determined whether such markers, combined in an inflammation-based score, can reliably predict outcome and what the optimal constituents of this score would be.

Therefore, the aim of the present Chapter was to examine the relationship between markers of the systemic immune/inflammatory response and all cause, cancer, cardiovascular and cerebrovascular mortality in a large incidental cohort.

9.3. MATERIALS AND METHODS

From the cohort previously described in Chapter 4, patients in North Glasgow, who had a single blood sample taken for C-reactive protein and albumin were considered [Proctor et al., 2010]. Only those patients who also had a differential white cell count available, including lymphocyte, neutrophil and platelet count were included. Patients were sampled incidentally between January 2000 and November 2008. The sample size was originally based on similar work investigating the association between C-reactive protein and all-cause

survival [Marsik et al., 2008]. If multiple samples had been taken during this time period, only the initial set was used. Only those patients undergoing blood sampling after the introduction of the routine measurement of high sensitivity C-reactive protein (1st of August 2006), were included in the analysis investigating the prognostic value of high sensitivity thresholds ($<5\text{mg/l}$). Prior to the introduction of high-sensitivity C-reactive protein measurements, the limit of detection was a concentration of less than 5mg/L and 0.05mg/l thereafter. Standardised thresholds for C-reactive protein ($>10\text{mg/l}$), white cell count ($>11\times 10^9/\text{l}$), neutrophil count ($>7.5\times 10^9/\text{l}$), lymphocyte count ($>3\times 10^9/\text{l}$) and platelet count ($>400\times 10^9/\text{l}$) were used. High sensitivity C-reactive protein threshold $>3\text{mg/l}$, previously reported to be associated with survival [Ridker, 2003; Ridker, 2008; Lee et al., 2011; Ko et al., 2012] and shown to be the optimal threshold in this cohort [Proctor et al., 2013], was also studied.

As the presence of malignancy is known to be associated with activation of the systemic immune/inflammatory response [Proctor et al., 2010], all patients with a diagnosed malignancy any time prior to or within 14 days of blood sample were corrected for. Whether the patient had been sampled as an inpatient or outpatient was also noted.

Patient outcome and mortality was established through linkage with the Information Services Division (ISD) for Scotland by exact matching surname, forename, sex and date of birth (Appendix 3). If no match on surname then Daitch-Mokotoff followed by NYSIIS soundex algorithms were employed. On remaining unmatched cases, forenames and surnames were switched followed by matching on previous surnames, reversing gender, using first initial opposed to full forename and finally on surname, gender and date of birth only. On all those not with exact matches a manual check on whether to include or exclude was taken. Of the 223,303 patients originally identified, 213,127 were successfully matched

through this linkage and 10 176 were not identified (5%). Of the matched patients, 160 481 patients had C-reactive protein and albumin as well as a complete differential white cell count. The presence of SMR01 records were used to determine whether a patient was sampled during a hospital admission, the length of hospital stay and whether the patient had died. Hospital admission was defined as having been sampled during an continuous inpatient stay. SMRO6 records containing International Classification of Disease (ICD) codes were used to identify those with a diagnosis of cancer prior to, or within 14 days of blood sampling, as previously described [Proctor et al., 2010]. Cancer related mortality was defined as ICD10 codes C00 - C97, cardiovascular (ischaemic heart disease) related mortality was defined as ICD10 codes I20 - I25 and cerebrovascular disease related mortality was defined as ICD10 codes I60 - I69. At the time of data collection, the ISD held complete SMR01 data on registered deaths until May 2011. Survival was calculated from time of blood sample to date of death or censor date (31st of May 2011). Patients were excluded if under the age of 16 years, demographic details were incomplete or if returned ISD fields were inaccurate or incomplete. Overall patient numbers were different in this Chapter, when compared to previous, as the inclusion criteria and matching process differed for ISD compared to the Scottish Cancer Registry linkage.

Deprivation was measured with the Scottish Index of Multiple Deprivation (SIMD) as detailed in Chapter 1 [Proctor et al., 2010]. Ethical approval was granted for the present study by the Research Ethics Committee, North Glasgow NHS Trust.

Statistical analysis was performed as follows. Cox proportional hazard model was used to analyse the relationship between patient demographics, hospital admission, the presence of cancer, C-reactive protein, albumin, neutrophil, lymphocyte and platelet counts with all-cause mortality. Furthermore, cause specific mortality for those dying of cancer,

cardiovascular and cerebrovascular disease were also examined by individually entering these as the primary outcomes into the model.

In those patients with a high sensitivity C-reactive protein measurement, the prognostic value of the combination of C-reactive protein, albumin and neutrophils, adjusted for age, sex, deprivation, hospital admission and the presence of cancer, was analysed using the Cox proportional hazard model and the Area Under the receiver operating characteristic Curve (AUC). During this analysis the proportionality assumptions in the Cox model were explored using a log minus log visual inspection and were found to be satisfactory. The relationship between the optimised Glasgow Prognostic Score and survival was assessed using the Kaplan-Meier log rank. A p value of <0.05 was considered to have statistical significance in the analysis. Analysis was performed using SPSS software (SPSS, Chicago, IL, USA).

9.4. RESULTS

From GIOS cohort of 223,303 patients originally described [Proctor et al., 2010], 209,148 (94%) were matched with a unique patient record in the ISD dataset. Of this group 160,481 fitted the inclusion criteria and had a differential white cell count available, including neutrophil, lymphocyte and platelet counts. Most patients were under 65 years of age (n=103,779, 65%) and female (n=85,308, 53%). Forty four percent of patients (n=71,156) were SIMD 5 (most deprived). Fifty four percent (n=86,339) of patients were sampled during an inpatient stay. Following the introduction of high sensitivity C-reactive protein, 52,091 patients with high sensitivity C-reactive protein, albumin, neutrophil, lymphocyte and platelet counts, were present.

On follow-up of all patients, there were 42,242 deaths in total of which 13,176 (31%) were cancer related, 6076 (14%) were cardiovascular related and 3638 (9%) were cerebrovascular related. The minimum follow-up from blood sample for all patients was 31 months and the maximum 134 months (median follow-up 69 months for survivors). The relationship between patient demographics, hospital admission, history of cancer, C-reactive protein (>10mg/l), albumin, neutrophil, lymphocyte and platelet counts and survival (n=160,481) is shown in Table 9.1. Increasing age, male gender, increasing deprivation, hospital admission, history of cancer, high C-reactive protein (>10mg/l), neutrophil and platelet counts, as well as low albumin and lymphocyte counts were independently associated with an increase in all-cause mortality (all $p < 0.001$). Increasing age, male gender, history of cancer, high C-reactive protein (>10mg/l), neutrophil and platelet counts as well as low albumin and lymphocytes were independently associated with an increase in cancer specific mortality (all $p < 0.001$). Increasing age, male gender, increasing deprivation, hospital admission, history of cancer, high C-reactive protein (>10mg/l), neutrophil and platelet counts as well as low albumin was independently associated with an increase in cardiovascular mortality (all $p \leq 0.001$). Increasing age, hospital admission, no history of cancer, high C-reactive protein (>10mg/l), neutrophil count as well as low albumin was independently associated with an increase in cerebrovascular mortality (all $p < 0.001$). The results of Table 9.1 show that activation of the immune/inflammatory response, as evidenced by an elevated C-reactive protein, predicts all-cause mortality but was most strongly associated with death from cancer (HR 1.85) when compared with cardiovascular (HR 1.54) or cerebrovascular (HR 1.31) disease.

The relationship between patient demographics, hospital admission, history of cancer, C-reactive protein (>3mg/l), albumin, neutrophil, lymphocyte and platelet counts and survival

in patients with a differential white cell count following the introduction of high sensitivity C-reactive protein (n=52,091) is shown in Table 9.2. Increasing age, male gender, hospital admission, history of cancer, high C-reactive protein (>3mg/l), neutrophil and platelet counts as well as low albumin and lymphocyte counts were independently associated with an increase in all-cause mortality (all $p<0.001$). Increasing age, male gender, history of cancer, high C-reactive protein (>3mg/l), neutrophil and platelet counts as well as low albumin and lymphocyte counts were independently associated with an increase in cancer mortality (all $p<0.001$). Increasing age, male gender, hospital admission, history of cancer, high C-reactive protein (>3mg/l) and neutrophil counts as well as low albumin was independently associated with an increase in cardiovascular mortality (all $p<0.001$). Increasing age, hospital admission, history of cancer, high C-reactive protein (>3mg/l) and neutrophil counts as well as low albumin was independently associated with an increase in cerebrovascular mortality (all $p<0.05$). The results of Table 9.2 show that activation of the immune/inflammatory response, as evidenced by an elevated C-reactive protein (>3mg/l), predicts all-cause mortality but was most strongly associated with death from cancer (HR 1.81) when compared with cardiovascular (HR 1.66) or cerebrovascular (HR 1.25) disease.

As C-reactive protein, albumin and neutrophils were consistently predictive of all-cause, cancer, cardiovascular and cerebrovascular specific mortality, these were taken forward to examine the prognostic value of their combination. The relationship between cumulative markers of the systemic immune/inflammatory response (including high sensitivity C-reactive protein, albumin and neutrophil count) and mortality in an incidental cohort (n=52,091) is shown in Table 9.3. The model was built up in a similar manner as previously demonstrated [Proctor et al., 2013] with the area under the curve representing the full model for each outcome. All combinations of these systemic inflammatory markers were

predictive of all-cause, cancer, cardiovascular and cerebrovascular specific survival (all $p < 0.01$) independent of age, sex deprivation, hospital admission and the presence of cancer (Figure 9.1). The Kaplan-Meier plot, for a score of 3, shows initial steep curves that subsequently do not diverge. This suggests that systemic inflammation is particularly predictive of early all-cause mortality. The areas under the curve represent the full model for each of the three outcomes.

9.5. DISCUSSION

The results of the present Chapter show that, in a large incidentally sampled cohort of patients ($n=160,481$), a number of routine objective markers of the systemic immune/inflammatory response examined were independently associated with all-cause mortality. Furthermore, of these C-reactive protein, albumin and neutrophils were confirmed to have independent prognostic value not only in patients with cancer but also in patients with cardiovascular and cerebrovascular disease. While these variables do not predict cause of death, there are correlations between lethal pathological processes, such as cancer and atherosclerotic disease, and activation of the systemic immune/inflammatory response. These results implicate the activation of the systemic immune/inflammatory response as a key and common factor in the shortened survival in the most common lethal disease states.

The results of the present Chapter are consistent with previous work using individual markers of the systemic immune/inflammatory response to predict all-cause mortality [Goldwasser and Feldman, 1997; Gillum et al., 2005; Marsik et al., 2008; Grimm et al., 2009; Zacho et al., 2010] and goes further to demonstrate that the combination of C-reactive protein, albumin, neutrophils can further improve the prognostic value of the systemic immune/inflammatory response. Furthermore, these associations remained following

adjustment for hospital admission and a previous diagnosis of cancer, events likely to be associated with raised inflammatory markers and reduced survival.

With reference to cancer, there has been growing interest in the role of markers of the systemic immune/inflammatory response in predicting outcome [McMillan, 2012; Guthrie et al., 2013] and the clinical utility of combining such markers with tumour based factors to provide improved patient counselling and personalised treatment is becoming increasingly recognised [Gakis et al., 2011; Lamb et al., 2012; Inoue et al., 2013; McMillan, 2013]. The present results provide a framework for the further development of a paradigm that uses both patient and tumour based objective factors to assess patient prognosis.

With reference to cardiovascular and cerebrovascular disease the prognostic value of C-reactive protein is well established [Kaptoge et al., 2012]. Indeed, high sensitivity C-reactive protein has been recommended for use in cardiovascular risk stratification in moderate risk, asymptomatic patients [Greenland et al., 2010]. Therefore, the present results also provide further evidence for the clinical utility of systemic inflammation-based scoring systems in predicting outcome in cardiovascular and cerebrovascular disease. Indeed, this approach with C-reactive protein and albumin as core factors has recently been validated in the NHANES III cohort [Van Hemelrijck et al., 2012].

The present Chapter shows that activation of the immune/inflammatory response predicts all-cause mortality, predominantly within the first year following sample, with the strongest association in cancer, followed by cardiovascular and cerebrovascular disease. The basis of the independent prognostic value of C-reactive protein, albumin and neutrophil count over all disease states (and platelets in cancer) is not clear and likely to be complex. However, it is likely that these factors are associated with aspects of the immune/inflammatory response and in particular the innate immune/inflammatory response.

Therefore, it may be hypothesised that the complementary prognostic value of these factors reflects an upregulation of the innate immune/inflammatory response and is likely to identify those with an ongoing pathological process who may die early as a consequence. The consistency of observations across different tumour types and now different disease states suggests that this hypothesis is worthy of further study.

In the present Chapter, the cohort has a number of limitations. The patients were selected on the basis that measurements of C-reactive protein, albumin and a differential white cell count had been performed and were therefore not necessarily representative of all patients treated in the North Glasgow area. It is also recognised that patients may have concurrent morbidity causing a rise in their C-reactive protein and derangement of their albumin and other haematological parameters.

In summary, a number of markers of the systemic immune/inflammatory response, reflecting an optimised Glasgow Prognostic Score, have prognostic utility not only in cancer, but also in cardiovascular and cerebrovascular disease. Further work to validate such findings in other large cohorts is warranted.

Table 9.1. The relationship between patient demographics, markers of the systemic immune/inflammatory response and mortality

		n (%) n=160 481	All-cause mortality Deaths = 42,242				Cancer mortality Deaths = 13,176				Cardiovascular mortality Deaths = 6076				Cerebrovascular mortality Deaths = 3638			
			Univariate		Multivariate		Univariate		Multivariate		Univariate		Multivariate		Univariate		Multivariate	
			HR	p-value	HR	p-value	HR	p-value	HR	p-value	HR	p-value	HR	p-value	HR	p-value	HR	p-value
Age	<65/	103,779 (65)	1		1		1		1		1		1		1		1	
	65-74	26,989(17)	4.00	<0.001	3.31	<0.001	3.76	<0.001	2.21	<0.001	5.80	<0.001	5.64	<0.001	8.45	<0.001	8.18	<0.001
	>75	29,716 (18)	8.27	<0.001	6.51	<0.001	4.36	<0.001	2.41	<0.001	12.58	<0.001	12.36	<0.001	28.41	<0.001	26.10	<0.001
Sex	Male	75,173 (47)	1		1		1		1		1		1		1		1	
	Female	85,308 (53)	0.86	<0.001	0.77	<0.001	0.76	<0.001	0.70	<0.001	0.72	<0.001	0.63	<0.001	1.18	<0.001	0.94	0.232
SIMD	1	21,362 (13)	1		1		1		1		1		1		1		1	
	2	17,778 (11)	1.03	0.216	1.11	<0.001	1.03	0.459			1.10	0.104	1.21	0.001	0.84	0.012		
	3	20,432 (13)	1.16	<0.001	1.20	<0.001	1.18	<0.001			1.18	0.002	1.28	<0.001	0.91	0.139		
	4	29,753 (19)	1.28	<0.001	1.30	<0.001	1.23	<0.001			1.42	<0.001	1.47	<0.001	1.00	0.986		
	5	71,156 (44)	1.25	<0.001	1.36	<0.001	1.00	0.882			1.32	<0.001	1.45	<0.001	1.16	0.004		
Hospital Admission	No	74,142 (46)	1		1		1		1		1		1		1		1	
	Yes	86,339 (54)	1.75	<0.001	1.21	<0.001	1.55	<0.001	1.00	0.861	1.81	<0.001	1.33	<0.001	2.00	<0.001	1.49	<0.001
History of Cancer	None	139,080 (87)	1		1		1		1		1		1		1		1	
	Present	21,401 (13)	3.40	<0.001	2.13	<0.001	12.01	<0.001	8.94	<0.001	1.45	<0.001	0.84	<0.001	1.58	<0.001	0.84	<0.001
C-reactive protein	<10mg/l	94,544 (59)	1		1		1		1		1		1		1		1	
	>10mg/l	65,937 (41)	2.71	<0.001	1.60	<0.001	3.16	<0.001	1.85	<0.001	2.33	<0.001	1.54	<0.001	2.17	<0.001	1.31	<0.001
Albumin	>35mg/l	135,087 (84)	1		1		1		1		1		1		1		1	
	<35mg/l	25,394 (16)	3.68	<0.001	2.02	<0.001	4.25	<0.001	2.08	<0.001	2.36	<0.001	1.33	<0.001	2.54	<0.001	1.41	<0.001
Neutrophil count	<7.5×10 ⁹ /l	127,736 (80)	1		1		1		1		1		1		1		1	
	>7.5×10 ⁹ /l	32,745 (20)	2.18	<0.001	1.49	<0.001	2.51	<0.001	1.41	<0.001	1.89	<0.001	1.51	<0.001	1.89	<0.001	1.52	<0.001
Lymphocyte count	>3×10 ⁹ /l	12,542 (8)	1		1		1		1		1		1		1		1	
	<3×10 ⁹ /l	147,939 (92)	1.87	<0.001	1.21	<0.001	1.84	<0.001	1.19	<0.001	1.58	<0.001	0.97	0.564	2.00	<0.001	1.11	0.272
Platelet count	<400×10 ⁹ /l	143,431 (89)	1		1		1		1		1		1		1		1	
	>400×10 ⁹ /l	17,050 (11)	1.64	<0.001	1.07	<0.001	2.03	<0.001	1.39	<0.001	1.17	<0.001	0.81	<0.001	1.51	<0.001	1.03	0.584

Table 9.2. The relationship between patient demographics, markers of the systemic immune/inflammatory response (including high sensitivity C-reactive protein) and mortality

		n (%) n=52 091	All-cause mortality Deaths = 8243				Cancer mortality Deaths = 3480				Cardiovascular mortality Deaths = 929				Cerebrovascular mortality Deaths = 553			
			Univariate		Multivariate		Univariate		Multivariate		Univariate		Multivariate		Univariate		Multivariate	
			HR	p-value	HR	p-value	HR	p-value	HR	p-value	HR	p-value	HR	p-value	HR	p-value	HR	p-value
Age	<65/	36,834 (71)	1		1		1		1		1		1		1		1	
	65-74	7277 (14)	4.28	<0.001	2.95	<0.001	4.36	<0.001	2.12	<0.001	6.26	<0.001	5.73	<0.001	7.75	<0.001	7.39	<0.001
	>75	7880 (15)	8.10	<0.001	5.24	<0.001	4.82	<0.001	2.20	<0.001	14.36	<0.001	13.18	<0.001	24.31	<0.001	22.20	<0.001
Sex	Male	25,411 (49)	1		1		1		1		1		1		1		1	
	Female	26,680 (51)	0.87	<0.001	0.77	<0.001	0.86	<0.001	0.76	<0.001	0.72	<0.001	0.61	<0.001	1.18	0.058		
SIMD	1	6943 (13)	1				1				1				1			
	2	5972 (12)	1.07	0.141			1.02	0.768			1.36	0.024			0.92	0.639		
	3	6749 (13)	1.16	0.001			1.19	0.004			1.23	0.133			0.81	0.199		
	4	9583 (18)	1.17	<0.001			1.07	0.275			1.58	<0.001			0.87	0.355		
	5	22 844 (44)	1.06	0.129			0.83	<0.001			1.21	0.092			0.96	0.777		
Hospital Admission	No	23,616 (45)	1		1		1		1		1		1		1		1	
	Yes	28,475 (55)	1.76	<0.001	1.16	<0.001	1.56	<0.001	0.96	0.229	1.82	<0.001	1.29	<0.001	2.31	<0.001	1.74	<0.001
History of Cancer	None	20,917 (40)	1		1		1		1		1		1		1		1	
	Present	31,174 (60)	4.74	<0.001	2.49	<0.001	12.75	<0.001	8.00	<0.001	1.57	<0.001	0.74	0.002	1.78	<0.001	0.77	0.032
C-reactive protein	<3mg/l	20,917 (40)	1		1		1		1		1		1		1		1	
	>3mg/l	31,174 (60)	3.05	<0.001	1.55	<0.001	3.88	<0.001	1.81	<0.001	2.78	<0.001	1.66	<0.001	2.10	<0.001	1.25	0.032
Albumin	>35mg/l	41,010 (79)	1		1		1		1		1		1		1		1	
	<35mg/l	11,081 (21)	5.19	<0.001	2.68	<0.001	6.40	<0.001	2.95	<0.001	3.57	<0.001	1.87	<0.001	2.69	<0.001	1.28	0.010
Neutrophil count	<7.5×10 ⁹ /l	41,617 (80)	1		1		1		1		1		1		1		1	
	>7.5×10 ⁹ /l	10,474 (20)	2.41	<0.001	1.63	<0.001	2.66	<0.001	1.47	<0.001	1.80	<0.001	1.49	<0.001	2.14	<0.001	1.89	<0.001
Lymphocyte count	>3×10 ⁹ /l	5391 (10)	1		1		1		1		1		1		1		1	
	<3×10 ⁹ /l	46,700 (90)	2.06	<0.001	1.25	<0.001	2.42	<0.001	1.42	<0.001	1.37	0.008	0.80	0.083	1.78	0.001	1.02	0.992
Platelet count	<400×10 ⁹ /l	47,160 (91)	1		1		1		1		1		1		1		1	
	>400×10 ⁹ /l	4931 (9)	2.05	<0.001	1.16	<0.001	2.70	<0.001	1.54	<0.001	1.39	0.001	0.82	0.055	1.66	<0.001	1.01	0.950

Table 9.3. The relationship between markers of the systemic immune/inflammatory response (including high sensitivity C-reactive protein) and mortality: Adjusted for age, sex, deprivation, hospital admission and the presence of cancer

	n (%)	All-cause mortality Deaths = 8243			Cancer mortality Deaths = 3480			Cardiovascular mortality Deaths =929			Cerebrovascular mortality Deaths =553			
		HR	p-value	AUC	HR	p-value	AUC	HR	p-value	AUC	HR	p-value	AUC	
		n=52,091												
	0	26,603 (51)	1			1			1			1		
CRP > 10mg/l	1	13,734 (26)	1.92	<0.001		1.99	<0.001		1.66	<0.001		1.53	<0.001	
+Albumin >35g/L	2	8635 (17)	3.45	<0.001		4.27	<0.001		2.67	<0.001		1.76	<0.001	
+Neutrophils >7.5×10 ⁹ /l	3	3119 (6)	6.45	<0.001	0.720	7.95	<0.001	0.728	3.49	<0.001	0.645	2.50	<0.001	0.612
	0	17,542 (34)	1			1			1			1		
hs-CRP >3mg/l	1	19,880 (38)	1.76	<0.001		1.94	<0.001		1.62	<0.001		1.40	0.006	
+ Albumin >35g/L	2	11,158 (21)	3.58	<0.001		4.44	<0.001		2.92	<0.001		1.90	<0.001	
+ Neutrophils>7.5×10 ⁹ /l	3	3511 (7)	7.37	<0.001	0.723	9.32	<0.001	0.731	4.03	<0.001	0.650	3.10	<0.001	0.623

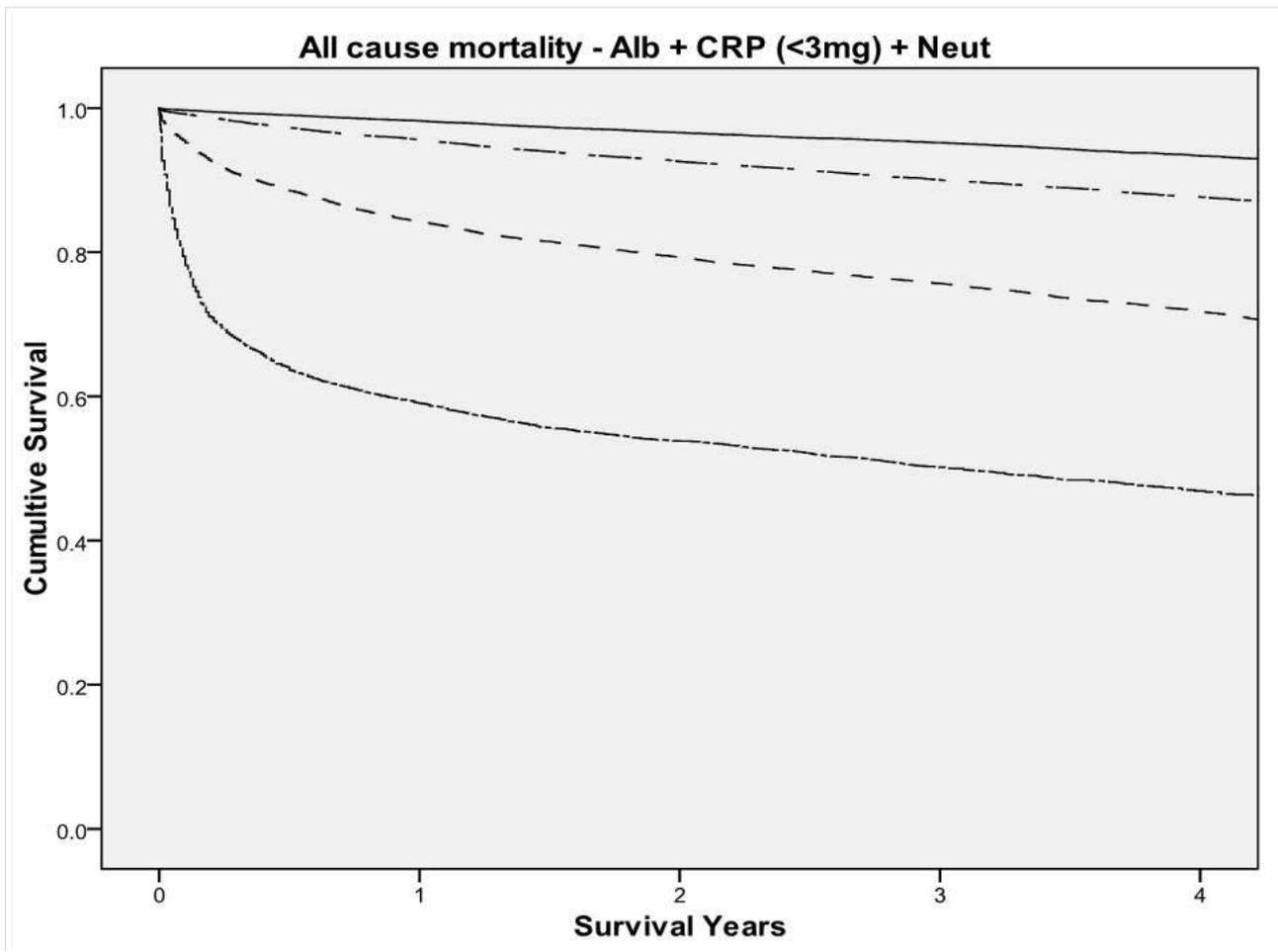


Figure 9.1. The relationship between cumulative markers of the systemic immune/inflammatory response and all-cause mortality (including high sensitivity C-reactive protein, albumin and neutrophil count) and survival (0-top, solid line; 1-upper separated dot and dash line, 2-middle large dash line, 3-lower close dot and dash line, n=52,091)

10.0. CONCLUSION

At the beginning of this period of research, as detailed in Chapter 1, it was clear that inflammation played an integral role in the pathogenesis and progression of malignancy and atherosclerotic disease. It had also been shown that activation of the systemic immune/inflammatory response, as measured by an objective systemic inflammatory-based score, was associated with reduced survival in particular cohorts with cancer, cardiovascular disease and stroke. However, it was unclear as to whether this was a universal phenomenon affecting all cancer types and what the optimal serum constituents of an inflammation based prognostic score would be for patients with cancer. It had also not been determined whether the link between inflammation and survival in arteriosclerotic disease was measurable by a similar inflammation-based score. The aim of this body of work was to address these issues in a single cohort of patients.

In Chapter 4, the relationships between the presence of cancer, serum levels of markers of systemic inflammation (mGPS) and other routinely sampled biochemical variables were investigated. Indeed, people with cancer were more likely to have an elevated mGPS and derangement of serum biochemical markers, both factors that had previously been associated with mortality. Interestingly, there were also significant interrelationships between markers of the systemic immune/inflammatory response and biochemical variables, suggesting that inflammation may be the underlying pathophysiological mechanism. It was also demonstrated that systemic inflammation (mGPS) across tumour types differed with the proportion of non-inflamed patients correlating with nationally reported five-year survival rates.

In Chapter 5, the relationship between the mGPS, biological parameters, tumour site and survival in patients with malignancy was investigated. It was discovered that serum C-reactive protein, albumin, liver function tests and calcium were all shown to have independent prognostic value in patients with a recent history of malignancy. Moreover, the relationship between the mGPS and survival was independent of tumour site and thus provided evidence that the systemic immune/inflammatory response is a universal marker of poor outcome in patients with malignancy. It was indicated that further work investigating the potential of combining a measure of the host immune/inflammatory response with conventional TNM staging would be worthwhile.

In Chapter 6, a comparison of the prognostic value of commonly reported systemic inflammation-based prognostic scores was carried out. The mGPS, NLR, PLR, PI and PNI were all shown to be associated with survival in patients with a recent diagnosis of cancer and it was argued that these scores were essentially measuring a common pathological process. Scores incorporating C-reactive protein were identified as having superior predictive nature and it was proposed that the widely validated mGPS should be used in the routine assessment of all patients with cancer. It was also noted that the addition of a white cell count to the mGPS may improve its prognostic value and further work on this issue was recommended.

In Chapter 7, the prognostic value of a derived systemic inflammation-based score (derived Neutrophil to Lymphocyte Ratio) using a white cell and neutrophil count widely available in chemotherapy trial databases was evaluated. The derived Neutrophil to Lymphocyte Ratio (neutrophil count/ white cell count - neutrophil count) had comparable prognostic value to the NLR and was shown to be predictive independent of tumour site. It was suggested that large oncology datasets be analysed, using the derived Neutrophil to

Lymphocyte Ratio, to assess further the clinical utility of this systemic inflammation-based prognostic score.

In Chapter 8, the aim was to examine whether it was possible to improve the prognostic value of the mGPS by including other markers of the systemic inflammatory response. The addition of a high sensitivity C-reactive protein measurement, neutrophil and platelet counts were found to enhance the predictive value (AUC) of the Glasgow Prognostic Score as well as improving the differentiation of patients at different levels of risk. These results were consistent with the concept that the systemic inflammatory response in patients with cancer was primarily due to an upregulation of the innate immune/ inflammatory response.

Given the universal utility of inflammation-based prognostic scores in patients with cancer, it was of interest as to whether a comparable score would be associated with all-cause, cardiovascular and cerebrovascular survival mortality. In Chapter 9, the prognostic value of C-reactive protein and albumin together with neutrophil, lymphocyte and platelet counts were investigated in a large unselected cohort. It was concluded that a similar systemic inflammation-based prognostic score to that used in cancer, containing high sensitivity C-reactive protein, albumin and neutrophils, was also independently associated with all-cause, cardiovascular and cerebrovascular mortality.

There are several areas for future study and potential clinical application of the inflammation-based scores presented in this body of work. Biomarkers are a topic of current interest as they can not only play an important role in disease detection and risk stratification, but can also assist in the personalisation of treatment [La Thangue and Kerr, 2011]. Biomarkers offer advantages over traditional clinical, radiological and pathological prognostic modalities given their objective nature [Strimbu and Tavel, 2010]. Furthermore it

has been suggested that the optimal biomarker, as is the case with the scores presented in the current body of work, would accurately predict outcome and could assist in the tailoring of appropriate treatment while being universally available and financially viable [Paoletti et al., 2004].

With regards to cancer screening, it has been suggested that genetic profiling may assist in the identification of high risk groups that will benefit from intensive surveillance [Dunlop et al., 2013]. However, to date genetic variants associated with an increased risk of colorectal cancer have not been shown to have a detrimental effect on survival [Tenesa et al., 2010]. It is of significance, when considered with the results of the current work, that Tao and co-workers have reported C-reactive protein levels in conjunction with Faecal Occult Blood Tests to improve the sensitivity and specificity of colonoscopy for colorectal cancer screening [Tao et al., 2012]. This clearly raises the possibility that the addition of an inflammation-based prognostic score may help improve this further. When screening for cardiovascular disease it has been recommended that C-reactive protein is also used to identify those at risk as well as help determine who will benefit from Statin therapy [Greenland et al., 2010; Kaptoge et al., 2012]. Interestingly, in those treated, a reduction in C-reactive protein concentrations has also been shown to correlate with improved outcomes independent of cholesterol reduction [Rao and Milbrandt, 2010; Tsai et al., 2012]. This raises the possibility that inflammation-based scores, including a sensitive C-reactive protein measurement, may have clinical utility in the screening and risk stratification of patients with atherosclerotic disease.

As the present body of work has shown, systemic inflammation-based prognostic scores have the potential to predict outcome in patients with cancer. A large number of other predictive biochemical biomarkers have been suggested, including carbonic anhydrase IX in

clear cell renal cancer [Stillebroer et al., 2010] and IMP3 in colorectal cancer [Lin et al., 2013]. Work has also been carried out investigating tumour indicators of survival in cancers with evidence demonstrating particular genetic signatures to be associated with survival [Ogino et al., 2009]. However, these tests are highly specialised with limited clinical availability and significant financial cost. Indeed, Kerr and Midgley stated:

The genetic model for colorectal (CRC) tumourigenesis proposal by Fearon and Vogelstein two decades ago, although broadly correct, has yielded to a much more complex and multifaceted paradigm. This has led to the clinical community being tantalized, but somewhat lacking in practical examples of biomarkers that can be applied for obvious patient benefit. There is, of course, the example of seeming resistance to anti-epidermal growth factor receptor (EGFR) antibody therapy in tumours carrying mutant K-ras, which has gained clinical traction, but there are no other biomarkers in CRC that are routinely incorporated into any treatment guidelines or algorithms. Many candidates have come and gone (p53, thymidylate synthase, loss of heterozygosity of 18q, to name but a few), lost in the fog of underpowered studies, technical variation in means of measurement, and absence of validating data sets with adequate controls [Kerr and Midgley, 2010].

It therefore follows that the simple, widely available, systemic inflammation-based scores presented in the current work may be a superior alternative.

Another potential use for systemic inflammation-based prognostic scores is to assist in the identification of patients that will benefit from particular treatment modalities. For example, anti-inflammatory chemotherapeutic agents, such as aspirin and cyclooxygenase-2 (COX-2) inhibitors, have been suggested to reduce risk of cancer as well as the subsequent

development of metastasis [Rothwell et al., 2012; Chan and Detering, 2013]. It is however worth noting that while Cox-2 inhibitors have a selective anti-inflammatory action, they also inhibit the production of prostacyclin, a prostaglandin that reduces platelet aggregation and arterial vasodilatation. This therefore allows the unchallenged promotion of platelet aggregation and vasoconstriction exerted by COX-1 induces thromboxane A₂, and a subsequent increased incidence of heart attack and stroke [Niederberger et al., 2004].

Non-steroidal anti-inflammatory drugs have also been shown to play a potential role in patients with advanced malignancy and cancer cachexia [Solheim et al., 2013]. It is possible that inflammation-based prognostic scores may be useful in identifying those who are systemically inflamed and would benefit from treatment, as well as measure the potential response [McMillan, 2013]. It has also been suggested that, in colorectal cancer patients with stage II disease, an inflammation-based prognostic score (GPS) could help identify patients who will gain a survival benefit from adjuvant chemotherapy [Toiyama et al., 2011]. Similarly, in patients with incurable cancer cachexia, the GPS has been able to identify those that would benefit from supportive parenteral nutrition [Bozzetti et al., 2014]. Also, the NLR has been shown to select those with metastatic prostate cancer that will benefit from Abiraterone hormonal therapy [Leibowitz-Amit et al., 2014].

There is also a growing pool of evidence regarding the use of markers of the systemic immune/inflammatory response when determining who will tolerate potentially toxic systemic therapies, such as palliative chemotherapy. In patients with advanced NSCLC, hypoalbuminaemia has been associated with anaemia, neuropathy, appetite loss, nausea and fatigue [Arrieta et al., 2010]. Furthermore, Gioulbasanis and co-workers demonstrated an association with mucositis, neurotoxicity, neutropenia, requirement for treatment termination and chemotherapy-related toxic deaths in patients with a raised inflammatory status (GPS)

receiving palliative chemotherapy for lung cancer [Gioulbasanis et al., 2012]. Similarly, in patients with advanced nasopharyngeal cancer, the presence of a non-normalising elevated C-reactive protein level after chemotherapy, has been associated with a poor outcome [Xia et al., 2013]. Also, patients receiving trans-arterial chemoembolisation for unresectable hepatocellular carcinoma were found to have worse outcomes in the presence of an activated immune/inflammatory response [Pinato and Sharma, 2012]. The current results show that inflammation-based prognostic scores have universal predictive value across all tumour types studied. They also offer an objective measurement of a patient's physical state, where performance status and physician opinion may vary. Moreover, activation of the systemic immune/inflammatory response has been shown to correlate with the majority of cancer symptoms [Laird et al., 2013] and accordingly, a standardised inflammation-based score, may assist traditional modalities in selecting patients who will benefit from further oncological therapies.

Another potential use for inflammation-based prognostic scores would be to monitor response to treatment. It has been suggested that targeted, immunomodulating therapies may confer a survival advantage and there is ongoing work to investigate such therapies, including anti-angiogenic TNF- α [Balkwill and Mantovani, 2010; Sasi et al., 2012]. It also been put forward been suggested that biomarkers are required to identify those appropriate for such targeted therapies and it is possible that inflammation-based scores may have a role to play in this process [Clarke et al., 2011]. Indeed, Ruxolitinib, a selective Janus kinase inhibitor (JAK1 and JAK2), has recently been shown to offer a survival advantage in patients with metastatic pancreatic cancer, but only in those patients with evidence of an active systemic inflammation as measured by a mGPS of 1 or 2 [Hurwitz H et al., 2014]. Moreover, there is also

some evidence that similar immune/inflammatory modification may be employed in cardiovascular disease [Frangogiannis, 2012].

The current piece of work gives an insight into the particular aspect of the immune/inflammatory response that is detrimental in cancer. Of the prognostic indices examined, it was noted that those associated with a reduced survival, including C-reactive protein as well as neutrophil and platelet counts, were predominantly linked to the innate immune response. Indeed, it is becoming clear that up-regulation of the innate response is detrimental as it facilitates angiogenesis and subsequent tumour dissemination [Tazzyman et al., 2013] and prognostic scores focusing on constituents of the innate immune response could be of future benefit.

In summary, this piece of research has demonstrated that a variety of inflammation-based prognostic scores, which essentially measure the same process, to be of universal predictive value in patients with cancer, cardiovascular disease and stroke. It also indicates that the basis of the relationship between the systemic immune/inflammatory response and outcome, particularly in cancer, is primarily due to activation of the innate immune/inflammatory response rather than the down regulation of the adaptive immune/inflammatory response.

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APPENDIX 1: GIOS Pathology Dataset

Inflammation Outcome Study - North of Glasgow Biochemical, Haematological and Pathological linkage with creation of GIOS database

Document date: 10th October 2009

(1.) Dataset (ProcExtract 1-6.exe) containing all primary single samples of C-reactive protein, albumin and Calcium between 1st January 2000 and 1st November 2008 in North Glasgow Hospital laboratories - Including Glasgow Royal Infirmary, Stobhill hospital, Gartnavel hospital and the Western Infirmary -extracted from TELEPATH systems with coded algorithm. Other variables extracted where available. 269 959 records extracted with 26 328 having temporary emergency hospital numbers not compatible with cross linkage.

(2.) Dataset (comp_histo_proc_extract 1-6.exe) containing all pathological diagnosis with malignant SNOMED codes in North Glasgow Hospital laboratories - Including Glasgow Royal Infirmary, Stobhill hospital, Gartnavel hospital and the Western Infirmary - extracted from telepath systems with coded algorithm.

(3.) Data cross linked to provide dataset with 223 303 patient (conforming to aforementioned criteria in Chapter 3) identifiers including hospital number gender, DOB, and postcode subsequent check referencing of >500 random samples to ensure accuracy.

(4.) SNOMED Clinical Terms (www.nlm.nih.gov/research/umls/Snomed/snomed_main) converted to ICD codes, subsequent clinical terms and then manual combination of morphology and topography terms to meaningful cancer diagnosis.

(5.) Converted to anonymised password-protected .zip file with the following fields:

Acquired fields:

- * ID at time of anonymisation
- * DOB date of birth
- * SEX gender
- * POSTCODE of address at time of blood sample
- * DATEOFREQUEST of blood test
- * Universal required variables: CRP (C-reactive Protein), ALB (Albumin), CCAL(Calcium)
- * Biochemical variables where available: BIL (Bilirubin), ALP (Alkaline Phosphatase), AST (aspartate transaminase), ALT (alanine transaminase) and γ -glutamyl transferase (GGT)
- * Haematological variables were available: WCC (White Cell), NEUT (Neutrophil), LYMPH (Lymphocyte), PLT (Platelet count)
- * DATEOFSAMPLE of neoplastic pathology diagnosis (1-4 where present)
- * Sample SNOMED topography (CATSITE) and morphology (CATTYPE) codes for each individual sample 1-4

Calculated fields:

- * AGE
- * SIMD 2006 Quintiles
- * CARSTAIRS Index

- * Blood sample "normal" classification depending on aforementioned range classification (Chapter3)
- * Inflammatory score mGPS
- * CATYPE1 type of primary malignancy and CASITE1 site of primary malignancy
- * NO presence of multiple malignancy or METYPEmetastatic malignancy
- * DIFFDATEDAYS difference in days between diagnosis of malignancy and blood test
- * CAGROUP broad group of malignancy

See attached file: *GIOS Dataset.SAV*

APPENDIX 2: GIOS Cancer Registry Dataset

Inflammation Outcome Study - North of Glasgow Biochemical, Haematological and Scottish Cancer Registry linkage to create GIOS database of patients with cancer and follow-up

Document date: 13th September 2010 with update linkage 12th July 2011

(1.) GIOS Dataset.SAV containing identifying variables as well as all primary single samples of C-reactive protein, albumin and Calcium between 1st January 2000 and 1st November 2008 in North Glasgow Hospital laboratories - Including Glasgow Royal Infirmary, Stobhill hospital, Gartnavel hospital and the Western Infirmary - as described in Appendix 1.

(2.) 223 303 records cross linked with Scottish Cancer Registry Data base held at West of Scotland Cancer Surveillance Unit, Faculty of Medicine, University of Glasgow, Glasgow G12 8RZ. Using exact matches of patients' forename, surname and date of birth followed by Daitch-Mokotoff and NYSIIS soundex algorithms were employed. On remaining unmatched cases, forenames and surnames were switched followed by matching on previous surnames, reversing gender, using first initial opposed to full forename and finally on surname gender and date of birth only. On all those not with exact matches a manual check on whether to include or exclude was taken. All unmatched cases were reviewed for manual match.

(3.) Anonymised data set returned to provide 21 668 patient conforming to aforementioned criteria in Chapter 4 with a various mix of ICD 9/10/Oncology codes (www.who.int/classifications/icd/en/) for diagnosis and death where appropriate. A selection of matched cases were reviewed for manual quality check (>500). ICD code manual combination of morphology and topography terms to meaningful cancer diagnosis and cause of death.

(5.) Converted to anonymised password-protected .zip file with the following fields:

Acquired fields:

- * ID anonymisation
- * AGEATREQ age at blood sample request
- * SEX gender
- * SIMD as GIOS dataset
- * CARSTAIRS Index as GIOS dataset
- * Universal required variables: CRP (C-reactive Protein), ALB (Albumin), CCAL(Calcium) as GIOS dataset
- * Biochemical variables where available: BIL (Bilirubin), ALP (Alkaline Phosphatase), AST (aspartate transaminase), ALT (alanine transaminase) and γ -glutamyl transferase (GGT) as GIOS dataset
- * Haematological variables were available: WCC (White Cell), NEUT (Neutrophil), LYMPH (Lymphocyte), PLT (Platelet count) as GIOS dataset
- * SITEICD10, SITEICDO, TYPEICDO regarding cancer diagnosis
- * DAYSINCIDENCE from blood sample to cancer incidence
- * SURVIVALREQ from cancer incidence to death or censor date
- * DESCCAUSE the primary cause of death
- * DUKES classification for colorectal cancer patients

Calculated fields:

- * Blood sample "normal" classification depending on aforementioned range classification (Chapter3)
- * Inflammatory score mGPS, GPS, NLR, PLR, PI
- * DESCICD10, DESCICDO3, DESCTYPEICDO2, DESCSITEICDO2 descriptions of cancer type for each code supplied.
- * TUMOURGROUPCR the categorical group of tumour type as detailed in Chapter 4.
- * DESCDEATHCAUSE the description of the ICD code of death
- * DEATHCAT the categorical cause of death

See attached file: *GIOS Cancer Registry Dataset.SAV*

APPENDIX 3: GIOS ISD dataset

Inflammation Outcome Study - ISD Linkage

Document date: 19th August 2011

(1.) GIOS Dataset.SAV containing identifying variables as well as all primary single samples of C-reactive protein, albumin and Calcium between 1st January 2000 and 1st November 2008 in North Glasgow Hospital laboratories - Including Glasgow Royal Infirmary, Stobhill hospital, Gartnavel hospital and the Western Infirmary - as described in Appendix 1.

(2.) 223 303 records cross linked with ISD, NHS National Services Scotland, Gyle Square, 1 South Gyle Crescent, Edinburgh, EH12 9EB, by exact matching surname, forename, sex and date of birth. If no match on surname then Daitch-Mokotoff followed by NYSIIS soundex algorithms were employed. On remaining unmatched cases, forenames and surnames were switched followed by matching on previous surnames, reversing gender, using first initial opposed to full forename and finally on surname gender and date of birth only. On all those not with exact matches a manual check on whether to include or exclude was taken. Successful match of 209 148 cases.

(3.) Anonymised data set returned to provide patient conforming to aforementioned criteria in Chapter 4 with a various mix of ICD 10 (www.who.int/classifications/icd/en/) for death where appropriate. ICD code manual combination of morphology and topography terms to meaningful cause of death.

(5.) Converted to anonymised password-protected .zip file with the following fields:

Acquired fields:

- * ID anonymisation
- * AGEATREQ age at blood sample request
- * SEX gender
- * SIMD as GIOS dataset
- * Universal required variables: CRP (C-reactive Protein), ALB (Albumin), CCAL(Calcium) as GIOS dataset
- * Biochemical variables where available: BIL (Bilirubin), ALP (Alkaline Phosphatase), AST (aspartate transaminase), ALT (alanine transaminase) and γ -glutamyl transferase (GGT) as GIOS dataset
- * Haematological variables were available: WCC (White Cell), NEUT (Neutrophil), LYMPH (Lymphocyte), PLT (Platelet count) as GIOS dataset
- * ISD_10_CODE of cause of death
- * ISD_CIS NUMBER of days continuous in patient hospital stay
- * ISD_ADMTYPE at time of blood sample emergency vs elective
- * ISD_DAYSFROMADMTOREQUEST days from admission to blood sample
- * ISD_INFECTIVECAUSE as primary reason of admission
- * ISD_SURVIVALFROMREQ from cancer incidence to death or censor date

Calculated fields:

- * Blood sample "normal" classification depending on aforementioned range classification (Chapter3)
- * Inflammatory score mGPS, GPS, NLR, PLR, PI
- * DESC_ISD_10 descriptions of cancer type for each code supplied.

See attached file: *GIOS ISD Dataset.SAV*

13.0 LIST OF ABBREVIATIONS

Abbreviation	Definition
ACS	Acute coronary syndrome
Alb	Albumin
Alk phos	Alkaline phosphatase
ALT	Alanine transaminase
AST	Aspartate transaminase
AUC	Area under the curve (ROC)
Bili	Bilirubin
Ca ²⁺	Adjusted calcium
CABG	Coronary Artery Bypass Graft
CI	Confidence interval
DNA	Deoxyribonucleic acid
dNLR	derived Neutrophil Lymphocyte Ratio
ECOG	Eastern Cooperative Oncology Group
GGT	γ -glutamyl transferase
g/l	Grams per litre
GPS	Glasgow Prognostic Score
HR	Hazard Ratio
hs-CRP	High sensitivity C-reactive protein
ICD	International Classification of Diseases
IL	Interleukin
ISD	Information Services Division for Scotland
LDH	Low density lipoprotein
LFT	Liver function test
mg/l	Milligrams per litre
mGPS	modified Glasgow Prognostic Score
mmol/l	millimoles per litre
NLR	Neutrophil Lymphocyte Ratio
NS	Not significant
oGPS	optimised Glasgow Prognostic Score
PCA	Percutaneous Coronary Angioplasty
PI	Prognostic Index
PLR	Platelet Lymphocyte Ratio
PNI	Prognostic Nutritional Index
p-value	Probability value
ROC	Receiver operating characteristic
SCR	Scottish Cancer Registry
SIMD	Scottish Index of Multiple Deprivation
SNOMED CT	Systematized Nomenclature of Medicine – Clinical Terms
STEMI	ST elevation myocardial infarction
TNM	Tumour Node Metastasis
U/l	Units per litre
μ mol/l	micromoles/litre
WBC/WCC	White blood cell count