

**DEVELOPMENT AND APPLICATION OF A NOVEL  
METHOD TO DETERMINE LARGE VERY LOW  
DENSITY LIPOPROTEIN (VLDL<sub>1</sub>) KINETICS**

by

**Iqbal A.R. AlShayji**

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# Author's Declaration

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Unless otherwise indicated by acknowledgment or reference to published literature, the presented work in this thesis is the author's own and has not been submitted for a degree at another institution.

**IQBAL ALSHAYJI** \_\_\_\_\_ Date \_\_\_\_\_

The findings of some of the studies have been published as follows:

## Published Papers

**Al-Shayji, I.A.R.**, Gill, J.M.R., Cooney, J., Siddiqui, S., & Caslake, M.J. (2007). Development of a novel method to determine very low density lipoprotein kinetics. *Journal of Lipid Research* 48: 2086-2095.

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*To my Mother,  
with the utmost love, respect and appreciation.*

*The more I know, the more I realise I don't know.*

# Table of Contents

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<b>LIST OF ABBREVIATIONS</b>	<b>XI</b>
<b>LIST OF TABLES</b>	<b>XIII</b>
<b>LIST OF FIGURES</b>	<b>XIV</b>
<b>ABSTRACT</b>	<b>XIX</b>
<b>1. INTRODUCTION AND LITERATURE REVIEW</b>	<b>1</b>
<b>1.1 Introduction</b>	<b>1</b>
<b>1.2 Lipoprotein Metabolism</b>	<b>1</b>
1.2.1 Classification and Properties of Lipoproteins	1
1.2.2 Exogenous Lipoprotein Metabolism	4
1.2.3 Endogenous Lipoprotein Metabolism	8
1.2.4 HDL Metabolism	9
1.2.5 Lipoprotein Lipase	10
1.2.6 Integration of TRL Metabolism	13
<b>1.3 Hepatic Assembly and Secretion of VLDL</b>	<b>15</b>
1.3.1 Apolipoprotein B Structure	15
1.3.2 Assembly of VLDL	16
<b>1.4 Regulation of VLDL Production</b>	<b>17</b>
1.4.1 Role of Fatty Acids	17
1.4.2 Role of Insulin	19
<b>1.5 Defective TRL Metabolism and Atherogenic Dyslipidaemia</b>	<b>20</b>
1.5.1 Atherogenic Lipoprotein Phenotype	21
1.5.2 Elevated Levels of VLDL <sub>1</sub> , Insulin Resistance and Obesity	21
1.5.3 VLDL <sub>1</sub> and the Generation of Atherogenic Dyslipidaemia	22
<b>1.6 Effect of Exercise on TRL Metabolism</b>	<b>25</b>
1.6.1 Exercise as a Therapy for Dyslipidaemia	25
1.6.2 Impact of Endurance Training	25
1.6.3 Impact of a Single Exercise Session	26
1.6.4 Energy Expenditure and Energy Deficit	27
1.6.5 Intermittent versus Continuous Exercise	28
1.6.6 Potential Mechanisms Responsible for the Exercise-Induced TG Reduction	28
	<b>VI</b>

1.6.6.1	Effect of Exercise on Chylomicron Metabolism	29
1.6.6.2	Effect of Exercise on VLDL Metabolism	29
1.6.7	Effect of ApoE Phenotype	31
<b>1.7</b>	<b>Measurements of TRL Kinetics</b>	<b>32</b>
1.7.1	Using Stable-Isotopes to Measure VLDL <sub>1</sub> and VLDL <sub>2</sub> Kinetics	33
1.7.2	Chylomicron Kinetics	36
1.7.2.1	ApoB-48 Labelling with Stable-Isotopes	36
1.7.2.2	Retinol Palmitate	37
1.7.2.3	Chylomicron-Like (Lipid) Emulsions	37
1.7.2.4	Intravenous Fat Tolerance Test	39
<b>1.8</b>	<b>Summary and Objectives</b>	<b>39</b>
<b>2.</b>	<b>GENERAL METHODS</b>	<b>41</b>
<b>2.1</b>	<b>Subject Recruitment</b>	<b>41</b>
<b>2.2</b>	<b>Anthropometry</b>	<b>42</b>
2.2.1	Height and Weight	42
2.2.2	Waist and Hip Circumferences	42
<b>2.3</b>	<b>Blood Sampling</b>	<b>42</b>
<b>2.4</b>	<b>Blood Samples Analysis</b>	<b>43</b>
2.4.1	Separation of VLDL <sub>1</sub> and VLDL <sub>2</sub> Lipoprotein Fractions	43
2.4.2	Spectrophotometric Assays	44
2.4.3	Insulin ELISA	45
2.4.4	Apolipoprotein E Phenotyping	45
2.4.5	Total Protein Determination (Lowry Method)	45
2.4.6	Accuracy and Precision of Assays	46
<b>2.5</b>	<b>Bruce Protocol</b>	<b>46</b>
<b>2.6</b>	<b>Submaximal Exercise Test</b>	<b>46</b>
<b>2.7</b>	<b>Monitoring of Heart Rate</b>	<b>47</b>
<b>2.8</b>	<b>Measurement of Oxygen Uptake and Carbon Dioxide Production</b>	<b>47</b>
<b>2.9</b>	<b>Apolipoprotein B Measurements in VLDL<sub>1</sub> and VLDL<sub>2</sub> Fractions – Comparison of a Manual and an Automated Method</b>	<b>49</b>
2.9.1	Introduction	49
2.9.2	Materials and Methods	50
2.9.2.1	Specimens and Separation of VLDL <sub>1</sub> and VLDL <sub>2</sub> fractions	50
2.9.2.2	Isopropanol Precipitation Method (Egusa <i>et al.</i> , 1983)	50
2.9.2.3	Immunoturbidimetric Method	50



2.9.2.4	Calculating ApoB %CV for Each Method	51
2.9.3	Results	51
2.9.4	Discussion	56
<b>2.10</b>	<b>Glucose Pilot Study</b>	<b>57</b>
2.10.1	Subjects and Methods	57
2.10.2	Protocol I	58
2.10.3	Protocol II	59
2.10.4	Protocol III	59
2.10.5	Conclusion	60
<b>3.</b>	<b>DEVELOPMENT OF A NOVEL METHOD TO DETERMINE VLDL<sub>1</sub> KINETICS</b>	<b>64</b>
<b>3.1</b>	<b>Introduction</b>	<b>64</b>
<b>3.2</b>	<b>Material and Methods</b>	<b>65</b>
3.2.1	Subjects	65
3.2.2	Intravenous Intralipid Test	66
3.2.3	Increasing the Intralipid Infusion Rate	67
3.2.4	Intralipid (Sf > 400) Separation from Whole Plasma	69
3.2.5	VLDL <sub>1</sub> and VLDL <sub>2</sub> Separation	69
3.2.6	Fasting Plasma Analysis	69
3.2.7	Correction for Glycerol	70
3.2.8	Kinetic Data Calculations	70
3.2.8.1	Calculating VLDL <sub>1</sub> -TG and VLDL <sub>1</sub> -ApoB Production Rates	70
3.2.8.2	Calculating VLDL <sub>1</sub> -TG and VLDL <sub>1</sub> -ApoB Fractional Synthetic and Catabolic Rates	72
3.2.8.3	Calculating Intralipid-TG Clearance Rate	72
3.2.9	Intralipid Recovery	75
3.2.10	Statistical Analyses	77
<b>3.3</b>	<b>Results</b>	<b>77</b>
3.3.1	Plasma, Intralipid, VLDL <sub>1</sub> - and VLDL <sub>2</sub> -TG Concentrations During and Post-Infusion	77
3.3.2	VLDL <sub>1</sub> - and VLDL <sub>2</sub> -ApoB Concentrations During Infusion	78
3.3.3	Kinetic Data	81
3.3.3.1	VLDL <sub>1</sub> -TG and VLDL <sub>1</sub> -ApoB Production Rates	81
3.3.3.2	VLDL <sub>1</sub> -TG and VLDL <sub>1</sub> -ApoB Fractional Synthetic and Catabolic Rates	81
3.3.3.3	Intralipid-TG Clearance Rate	81
3.3.4	Effect of Increasing the Intralipid Infusion Rate	83

3.3.5	Intralipid Recovery	83
3.3.6	Interrelationships between VLDL <sub>1</sub> -TG and VLDL <sub>1</sub> -apoB Production and TG Clearance	86
3.3.7	Relationships between Kinetic Variables and Subjects Characteristics	86
3.3.8	Relationships between Kinetic Variables and ALT Concentrations	87
<b>3.4</b>	<b>Discussion</b>	<b>91</b>
<b>4.</b>	<b>EFFECTS OF ACUTE HYPERINSULINAEMIA AND HYPERGLYCAEMIA ON VLDL<sub>1</sub>-TG AND VLDL<sub>1</sub>-APOB PRODUCTION AND INTRALIPID-TG CLEARANCE IN NORMOGLYCAEMIC SUBJECTS</b>	<b>97</b>
<b>4.1</b>	<b>Introduction</b>	<b>97</b>
<b>4.2</b>	<b>Subjects and Methods</b>	<b>99</b>
4.2.1	Subjects	99
4.2.2	Study Design	99
4.2.3	Intralipid Infusion	101
4.2.4	Analytical Methods	101
4.2.5	Calculations and Statistical Analyses	101
<b>4.3</b>	<b>Results</b>	<b>103</b>
4.3.1	Fasting Plasma Concentrations	103
4.3.2	Glucose, Insulin and NEFA Concentrations during and Post-Infusion	103
4.3.3	The Change in Production and Clearance Rates	105
4.3.4	VLDL <sub>1</sub> Particle Size	106
4.3.5	Predictors of the Change in Production and Clearance Rates	106
<b>4.4</b>	<b>Discussion</b>	<b>110</b>
<b>5.</b>	<b>EFFECTS OF MODERATE EXERCISE ON VLDL<sub>1</sub> KINETICS IN OVERWEIGHT/OBESE MIDDLE-AGED MEN</b>	<b>113</b>
<b>5.1</b>	<b>Introduction</b>	<b>113</b>
<b>5.2</b>	<b>Materials and Methods</b>	<b>115</b>
5.2.1	Subjects	115
5.2.2	Study Design	115
5.2.3	Treadmill Walk	116
5.2.4	Intralipid Infusion	116
5.2.5	Analytical Methods	116
5.2.6	Calculations and Statistics	117
5.2.7	Power Calculations	118

<b>5.3 Results</b>	<b>119</b>
5.3.1 Treadmill Walk	119
5.3.2 Fasting Concentrations in the Control and Exercise Trials	119
5.3.3 TRL Kinetics in Responses to Exercise	120
5.3.4 VLDL <sub>1</sub> Compositional Responses to Exercise	120
5.3.5 Predictors of the Exercise-Induced Changes in TRL Kinetics	123
<b>5.4 Discussion</b>	<b>125</b>
 <b>6. GENERAL DISCUSSION</b>	 <b>130</b>
<hr/>	
<b>6.1 Summary</b>	<b>130</b>
<b>6.2 Limitations of the Intralipid Method</b>	<b>134</b>
<b>6.3 Future Experiments and Applications of the Method</b>	<b>135</b>
<b>6.4 Conclusion</b>	<b>136</b>
 <b>7. REFERENCES</b>	 <b>137</b>
<hr/>	
 <b>8. APPENDICES</b>	 <b>163</b>
<hr/>	
Appendix A1: Subject Information Sheet and Consent Form – Chapters 3 & 4	164
Appendix A2: Subject Information Sheet and Consent Form - Chapter 5	169
Appendix B1: Health Screen Questionnaire – Chapters 3 & 4	176
Appendix B2: Health Screen Questionnaire – Chapter 5	178
Appendix C: Kinetic Results For All Subjects According To ApoE Phenotypes	180

# List of Abbreviations

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<b>Apo</b>	Apolipoprotein
<b>ABCA1</b>	ATP-binding cassette A1
<b>ALT</b>	Alanine aminotransferase
<b>ARF-1</b>	ADP ribosylation factor 1
<b>CABG</b>	Coronary artery bypass graft
<b>CE</b>	Cholesteryl esters
<b>CETP</b>	Cholesteryl ester transfer protein
<b>CHD</b>	Coronary heart disease
<b>CHOL</b>	Cholesterol
<b>CM</b>	Chylomicrons
<b>CR</b>	Clearance rate
<b>CV</b>	Coefficient of variation
<b>CVD</b>	Cardiovascular disease
<b>DNL</b>	<i>de novo</i> lipogenesis
<b>ER</b>	Endoplasmic reticulum
<b>ERGIC</b>	Endoplasmic reticulum-Golgi intermediate compartment
<b>Exp</b>	Exponential decay
<b>FA</b>	Fatty acid
<b>FC</b>	Free cholesterol
<b>FCR</b>	Fractional catabolic rate
<b>FSR</b>	Fractional synthetic rate
<b>GC-MS</b>	Gas chromatography mass spectrometry
<b>HDL</b>	High density lipoprotein
<b>HL</b>	Hepatic lipase
<b>HOMA<sub>IR</sub></b>	Homeostasis model assessment-estimated insulin resistance
<b>HSPG</b>	Heparan sulfate proteoglycans
<b>IDL</b>	Intermediate density lipoprotein
<b>IL</b>	Intralipid
<b>IVFTT</b>	Intravenous fat tolerance test
<b>LCAT</b>	Lecithin cholesterol acyltransferase

<b>LDL</b>	Low density lipoprotein
<b>LDLR</b>	LDL receptor
<b>LLTP</b>	Large lipid transfer protein
<b>LPL</b>	Lipoprotein lipase
<b>LRP</b>	LDL receptor-related protein
<b>MI</b>	Myocardial infarction
<b>MTP</b>	Microsomal triglyceride transfer protein
<b>MW</b>	Molecular weight
<b>NEFA</b>	Non-esterified fatty acid
<b>PL</b>	Phospholipids
<b>PLTP</b>	Phospholipids transfer protein
<b>PR</b>	Production rate
<b>QC</b>	Quality control
<b>SAAM II</b>	Simulation, Analysis And Modeling II
<b>SD</b>	Standard deviation
<b>sdLDL</b>	Small dense low density lipoprotein
<b>SEM</b>	Standard error of the mean
<b>S<sub>f</sub></b>	Svedberg floatation rate
<b>SR-BI</b>	Scavenger receptor type B class I
<b>SS</b>	Steady State
<b>TG</b>	Triglycerides
<b>TRL</b>	Triglyceride-rich lipoprotein
<b>VLDL</b>	Very low density lipoprotein
<b>VLDLR</b>	VLDL receptor
<b><math>\dot{V}O_{2\max}</math></b>	Rate of maximal oxygen uptake

# List of Tables

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## CHAPTER 1

Table 1.1: Composition and characteristics of the human plasma lipoproteins (Gurr <i>et al.</i> , 2002; Ginsberg <i>et al.</i> , 2005; Packard & Shepherd, 1997).....	2
Table 1.2: Characteristics of the major apolipoproteins (Gurr <i>et al.</i> , 2002; Ginsberg <i>et al.</i> , 2005).....	3
Table 1.3: Characteristics of tracers [modified from (Packard, 1995)]. .....	34

## CHAPTER 2

Table 2.1: Various times and speeds required for the separation of lipoprotein fractions using Beckman L8-60M Ultracentrifuge and SW 40 rotor.....	44
Table 2.2: Calculated mean, SD and CV for apoB concentrations measured 10 times for the same sample using the two methods .....	52
Table 2.3: Characteristics of the two female subjects who participated in Protocol I .....	58

## CHAPTER 3

Table 3.1: Subjects' physical and metabolic characteristics (n=10).....	66
Table 3.2: Fasting concentrations, apoE phenotypes and individual lipoprotein kinetic parameters calculated using the 'Intralipid method' (0.1 g.kg <sup>-1</sup> .h <sup>-1</sup> infusion dose) (n=10). .....	82
Table 3.3: Correlations between VLDL <sub>1</sub> -apoB and -TG production rates (expressed in mg.kg <sup>-1</sup> .d <sup>-1</sup> ) and subjects' characteristics.....	87
Table 3.4: Comparison of values for VLDL <sub>1</sub> -TG and VLDL <sub>1</sub> -apoB production rates (range) calculated in the present study (Intralipid Method) and in previously published studies using the stable-isotope method. ....	96

## CHAPTER 4

Table 4.1: Subjects' physical characteristics and mean fasting concentrations (n=8).....	100
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## CHAPTER 5

Table 5.1: Characteristics of the participants (n=12).....	115
Table 5.2: Plasma and VLDL <sub>1</sub> concentrations in the fasted state .....	121
Table 5.3: Exercise-induced compositional changes to circulating VLDL <sub>1</sub> particles.....	123

# List of Figures

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## CHAPTER 1

Figure 1.1: Chylomicron Metabolism.....	6
Figure 1.2: Model of possible hepatic uptake of TRL remnants.....	7
Figure 1.3: Overview of the general aspects of endogenous lipoprotein metabolism..	12
Figure 1.4: TRL metabolism in the [A] fasted and [B] postprandial state.....	14
Figure 1.5: The Intracellular assembly of VLDL.....	18
Figure 1.6: Schematic diagram of the mechanisms involved in the formation of atherogenic dyslipidaemia. ....	24
Figure 1.7: Schematic diagram of the stable-isotope laboratory protocol used to determine the kinetics of VLDL <sub>1</sub> - and VLDL <sub>2</sub> -TG and -apoB.....	35
Figure 1.8: A diagram of the mathematical multicompartmental model (using SAAM II software) used to simultaneously determine the kinetics of apoB and TG in VLDL <sub>1</sub> and VLDL <sub>2</sub> after a bolus injection of stable isotopes D <sub>3</sub> -leucine and D <sub>5</sub> -glycerol [Figure from (Adiels <i>et al.</i> , 2005a)].....	36

## CHAPTER 2

Figure 2.1: A schematic diagram of a 4-stage submaximal exercise test performed for subjects in Chapter 5. The blue block represents an expired air sample and heart rate measurements.....	47
Figure 2.2: ApoB concentrations (mg.dl <sup>-1</sup> , with linear-regression lines of ‘best-fit’ and equation of the line) in [A] VLDL <sub>1</sub> and [B] VLDL <sub>2</sub> fractions measured using an automated immunoturbidimetric method and by a manual isopropanol method with the line of equality. (n = 260) .....	53
Figure 2.3: A scatterplot of the percent difference in apoB concentrations (mg.dl <sup>-1</sup> ) between the two methods (isopropanol minus immunoturbidimetry) against their mean in [A] VLDL <sub>1</sub> and [B] VLDL <sub>2</sub> , with 95% limits of agreement (mean ± 2SD): VLDL <sub>1</sub> (0.8 ± 110.2%) and VLDL <sub>2</sub> (9.1 ± 66.4%), n = 260. ....	54
Figure 2.4: A scatterplot of the percent difference (isopropanol minus immunoturbidimetry) for apoB concentrations > 3.0 mg.dl <sup>-1</sup> against the mean of the two methods in [A] VLDL <sub>1</sub> and [B] VLDL <sub>2</sub> , with 95% limits of agreement (mean ± 2SD): VLDL <sub>1</sub> (20.2 ± 53.5%, n = 65) and VLDL <sub>2</sub> (17.7 ± 35.3%, n = 85). ....	55

Figure 2.5: [A] glucose, [B] insulin and [C] NEFA response (n = 2) for the glucose pilot - Protocol I: 30% (w/v) glucose (15 g in 50 ml) every 15 min for 2 h. The solid lines represent subject 1, while the dotted lines represent subject 2. ....	61
Figure 2.6: [A] glucose, [B] insulin and [C] NEFA response (n = 1) for the glucose pilot - Protocol II: a bolus drink of 20% (w/v) of glucose (30 g in 150 ml) followed by a 20% glucose drink (5 g in 25 ml) every 10 min for 2 h. ....	62
Figure 2.7: [A] glucose, [B] insulin and [C] NEFA response (n = 1) for the glucose pilot - Protocol III: a bolus glucose drink of 30 g (in 100 ml) followed by 15 g of glucose (in 25 ml) every 15 min for 4 hours. ....	63

### CHAPTER 3

Figure 3.1: A picture of a subject during an Intralipid trial.) .....	68
Figure 3.2: Changes in [A] VLDL <sub>1</sub> -TG and [B] VLDL <sub>1</sub> -apoB pools (mg) over time in a female subject No. 3 (55 yr, 84.5 kg) during infusion of 10% Intralipid (0.1 g.kg <sup>-1</sup> .h <sup>-1</sup> ). An initial bolus dose of 20% Intralipid (0.1 g.kg <sup>-1</sup> body mass) was given at 0 min. ....	71
Figure 3.3: Intralipid-TG concentrations (mmol.l <sup>-1</sup> ) [A] during infusion and [B] post-infusion of 0.1 g.kg <sup>-1</sup> .h <sup>-1</sup> of 10% Intralipid for a female subject (23 y, 52.0 kg). An initial bolus dose of 20% Intralipid (0.1 g.kg <sup>-1</sup> body mass) was given at 0 min. The Y-axis error bars are ± 13.8% (i.e. the 95% confidence interval for the measured Intralipid-TG concentration). ....	74
Figure 3.4: Schematic diagram of the 'Intralipid Method' protocol .....	76
Figure 3.5: Plasma-, Intralipid-, VLDL <sub>1</sub> - and VLDL <sub>2</sub> -TG concentrations (mmol.l <sup>-1</sup> ) [A] during infusion of 0.1 g.kg <sup>-1</sup> .h <sup>-1</sup> of 10% Intralipid and [B] for 75 min post-infusion. An initial bolus dose of 20% Intralipid (0.1 g.kg <sup>-1</sup> body mass) was given at 0 min. N = 10, values are mean ± SEM. ....	78
Figure 3.6: An example of a full 135min-Intralipid trial during development of the method. [Top] Plasma-, Intralipid-, VLDL <sub>1</sub> - and VLDL <sub>2</sub> -TG and [Bottom] VLDL <sub>1</sub> - and VLDL <sub>2</sub> -apoB concentrations during and post-infusion for a female subject (34 y, 55.2 kg). A bolus dose of 20% Intralipid (0.1 g.kg <sup>-1</sup> body mass) was injected at zero min, followed immediately by a constant infusion of 10% Intralipid (0.1 g.kg <sup>-1</sup> .h <sup>-1</sup> ) for 120 min. The dotted line represents the end of infusion. VLDL <sub>1</sub> kinetics for this subject were calculated from the first 75 min of infusion: VLDL <sub>1</sub> -TG and VLDL <sub>1</sub> -apoB production rates were 711.2 and 20.7 mg.h <sup>-1</sup> and fractional catabolic rates were 56.3 and 20.0 pools.d <sup>-1</sup> , respectively. Intralipid-TG clearance rate was calculated only from its exponential decay post-infusion as 146.9 pools.d <sup>-1</sup> , as no steady state was achieved during infusion. ....	79



- Figure 3.7: VLDL<sub>1</sub>-, VLDL<sub>2</sub>- and total VLDL-apoB concentrations (mg.dl<sup>-1</sup>) during infusion of 10% Intralipid (0.1 g.kg<sup>-1</sup>.h<sup>-1</sup>). An initial bolus dose of 20% Intralipid (0.1 g.kg<sup>-1</sup> body mass) was given at 0 min. N = 10, values are mean ± SEM. .... 80
- Figure 3.8: VLDL<sub>1</sub>- and VLDL<sub>2</sub>-TG/apoB ratios during infusion of 10% Intralipid (0.1 g.kg<sup>-1</sup>.h<sup>-1</sup>). An initial bolus dose of 20% Intralipid (0.1 g.kg<sup>-1</sup> body mass) was given at 0 min. N = 10, values are mean ± SEM. .... 80
- Figure 3.9: [A] An example of Intralipid separation in the High dose of Intralipid infusion (0.2 g.kg<sup>-1</sup>.h<sup>-1</sup>). Intralipid separation is done by overlaying 2 ml of plasma with 4 ml of density solution 1.006 g.ml<sup>-1</sup> and centrifuged at 10,000 rpm for 30 min. [B] A magnification of the bottom of the tube showing a thin layer of possible contamination of Intralipid at the top of plasma due to the high concentration of Intralipid. .... 84
- Figure 3.10: Scatterplots (with the line of equality) illustrating the agreement between [A] VLDL<sub>1</sub>-TG FSR (pools.d<sup>-1</sup>) and [B] Ln VLDL<sub>1</sub>-apoB FSR between the Low (0.1 g.kg<sup>-1</sup>.h<sup>-1</sup>) and High (0.2 g.kg<sup>-1</sup>.h<sup>-1</sup>) doses. FSR is equivalent to FCR in the fasted state. N = 5, *r* and *P* values are for Pearson product-moment correlations between variables. VLDL<sub>1</sub>-apoB FSR was expressed in pools.d<sup>-1</sup> prior to log-transformation. .... 85
- Figure 3.11: Scatterplots (with linear-regression lines of ‘best-fit’) illustrating the interrelationships between [A] Ln Intralipid-TG clearance rate and Ln VLDL<sub>1</sub>-apoB production rate, [B] Ln Intralipid-TG clearance rate and Ln VLDL<sub>1</sub>-TG production rate, and [C] Ln VLDL<sub>1</sub>-apoB and Ln VLDL<sub>1</sub>-TG production rates. N = 10, *r* and *P* values are for Pearson product-moment correlations between variables. Production rates and Intralipid-TG clearance rate were expressed in mg.h<sup>-1</sup> and pools.d<sup>-1</sup> prior to log-transformation, respectively. .... 88
- Figure 3.12: Scatterplots (with linear-regression lines of ‘best-fit’) illustrating the relationships between the kinetic variables [Ln VLDL<sub>1</sub>-TG production rate (left), Ln VLDL<sub>1</sub>-apoB production rate (middle) and Ln Intralipid-TG clearance rate (right)] and subjects’ characteristics: waist circumference (top), Ln fasting TG concentrations (middle) and Ln HOMA-estimated insulin resistance (HOMA<sub>IR</sub>) (bottom). N = 10, *r* and *p* values are for Pearson product-moment correlations between variables. .... 89
- Figure 3.13: Scatterplots (with linear-regression lines of ‘best-fit’) illustrating the relationships between serum ALT concentration (mU.l<sup>-1</sup>) and [A] Ln VLDL<sub>1</sub>-apoB production rate, [B] Ln Intralipid-TG clearance rate and [C] Ln VLDL<sub>1</sub>-apoB production rate. N = 9, *r* and *P* values are for Pearson product-moment correlations between variables. .... 90

## CHAPTER 4

- Figure 4.1: Study Protocol. In the Glucose trial a bolus drink of 30 g of glucose in 120 ml water was given at -60 mins, followed by 10 g of glucose in 40 ml of water at 15-min intervals thereafter. In the Control trial, equivalent volumes of water were given at the same intervals. .... 100
- Figure 4.2: Mean [A] glucose, [B] insulin and [C] NEFA concentrations before and during Intralipid infusion (left panel) and post-infusion (right panel) between the Control (●) and Glucose (○) trials. The dotted line represents the beginning of a bolus Intralipid dose of  $0.1 \text{ g.kg}^{-1}$  followed immediately by a constant Intralipid infusion of  $0.1 \text{ g.kg}^{-1}.\text{h}^{-1}$ . Significant differences between the two trials were tested using two-way ANOVA with (\*) representing  $P < 0.05$  and (\*\*)  $P < 0.001$ . .... 104
- Figure 4.3: Individual (according to apoE phenotype) and mean (bars) [A] VLDL<sub>1</sub>-TG and [B] VLDL<sub>1</sub>-apoB production rates and [C] Intralipid-TG clearance rate in the Control and Glucose trials ( $n = 8$ ). (#)  $P = 0.05$ , (\*)  $P < 0.05$ . .... 108
- Figure 4.4: Scatterplots (with linear-regression lines of ‘best-fit’) illustrating the relationships between the change (Control minus Glucose) in [A] NEFA concentrations at 0 min (just before the start of the Intralipid infusion) and the change in VLDL<sub>1</sub>-TG (left) and VLDL<sub>1</sub>-apoB production rates (right); between [B] serum ALT concentrations and the change in VLDL<sub>1</sub>-TG (left) and VLDL<sub>1</sub>-apoB production rates (right); and [C] HOMA<sub>IR</sub> and the change in Intralipid-TG clearance rate.  $N = 8$ ,  $r$  and  $P$  values are for Pearson product-moment correlations between variables. .... 109

## CHAPTER 5

- Figure 5.1: The mean percentage change in VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB production rates (PR) and fractional catabolic rates (FCR) and Intralipid (IL)-TG clearance rate (CR) in response to a single session of 120-min brisk walking at  $\sim 50\% \text{ VO}_{2\text{max}}$ . (\*  $P < 0.05$ , \*\*  $P < 0.001$ ). .... 121
- Figure 5.2: Individual (according to apoE phenotype) and mean (bars) VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB production rates (top panel) and fractional catabolic rates (FCR, middle panel) and Intralipid-TG clearance rate (bottom) in response to a single session of 120-min brisk walking at  $\sim 50\% \text{ VO}_{2\text{max}}$  (\*  $P < 0.05$ , \*\*  $P < 0.001$  for the difference in mean response between the Control and Exercise trials). .... 122
- Figure 5.3: Scattergrams indicating the relationships between the exercise-induced changes (Exercise minus Control) in [A] Intralipid-TG and VLDL<sub>1</sub>-TG FCR (pools. $\text{d}^{-1}$ ) and [B] VLDL<sub>1</sub>-apoB FCR (pools. $\text{d}^{-1}$ ) and HOMA<sub>IR</sub>. .... 124

Figure 5.4: Possible mechanisms involved in moderate exercise-induced reduction in VLDL<sub>1</sub>. Control: once in the circulation, the TG content of the newly secreted VLDL<sub>1</sub> particle from the liver, and consequently its size, are being reduced rapidly by hydrolysis by lipoprotein lipase (LPL) and the action of cholesteryl ester transfer protein (CETP), which promotes the transfer of TG from VLDL (and LDL) in exchange for cholesteryl ester (CE) from HDL. Exercise: A prior session of moderate exercise significantly reduces VLDL<sub>1</sub> concentration by increasing its clearance from plasma, possibly by compositional changes to the VLDL<sub>1</sub> particle which render it bigger in size and more TG-enriched, thereby increase its affinity for LPL-mediated lipolysis. The bigger particles are either produced directly by the liver and/or reduced CETP activity. Also, increased LPL activity is likely to play a role..... 129

## CHAPTER 8

Figure 8.1: Production rates of [A] VLDL<sub>1</sub>-TG and [B] VLDL<sub>1</sub>-apoB and [C] Intralipid-TG clearance rate for all subjects (n = 23) according to apoE phenotypes: E2/E3 (n = 3), E3/E3 (n = 11) and E4+ (n = 9).  $P > 0.05$  for differences between phenotypes. .... 180

# Abstract

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High concentrations of large very low density lipoproteins (VLDL<sub>1</sub>) give rise to atherogenic dyslipidaemia, which is usually associated with insulin resistant conditions (e.g. obesity) and increases the risk for cardiovascular disease (CVD). Isotopic tracer methods for determining VLDL<sub>1</sub> kinetics are costly, time-consuming, labour intensive and need experience and skill to calculate the kinetic parameters. The aim of this thesis was to develop a simpler and cost-effective method of obtaining triglyceride-rich lipoproteins (TRL) kinetic data, based on the fact that chylomicrons (CM) or CM-like particles (e.g. Intralipid) compete with large VLDL<sub>1</sub> for the same lipoprotein lipase (LPL)-mediated catalytic pathway. From this method, it was possible to determine VLDL<sub>1</sub>-triglyceride (TG) and -apolipoprotein (apo) B production rates and the Intralipid-TG clearance rate (as a surrogate measure of CM clearance). Kinetic data obtained from this method agreed with values and relationships obtained from stable isotope methods. The protocol is relatively quick, inexpensive, and transferable to non-specialist laboratories.

As a first application, the 'Intralipid method' was used to investigate the effects of hyperinsulinaemia and hyperglycaemia due to glucose ingestion on VLDL<sub>1</sub>-TG and -apoB production rates and Intralipid-TG clearance rate. This showed that hepatic VLDL<sub>1</sub> production is suppressed in response to hyperinsulinaemia and that the change in Intralipid-TG clearance rate with hyperinsulinaemia correlated significantly with HOMA-estimated insulin resistance (HOMA<sub>IR</sub>). In addition, alanine aminotransferase (ALT) concentrations (a marker for liver fat), within normal range, predicted the extent of hepatic VLDL<sub>1</sub> suppression.

Secondly, the Intralipid method was used to investigate the mechanisms responsible for the hypotriglyceridaemic effect of a moderate exercise session (120 min walking at 50%  $\dot{V}O_{2max}$ ) in overweight/obese middle-aged men; the section of the population at high risk of CVD in whom exercise-for-health guidelines are targeted. This showed that the exercise-induced reduction in plasma TG was due to increased VLDL<sub>1</sub>-TG and -apoB clearance, rather than decreased hepatic production. Exercise

also increased Intralipid-TG clearance rate, but to a lesser extent than VLDL<sub>1</sub>, suggesting an increased affinity of VLDL<sub>1</sub> for LPL-mediated lipolysis post-exercise.

Taken together, the Intralipid method is a relatively simple, safe and cost-effective method to determine in VLDL<sub>1</sub>-TG and -apoB production rates and Intralipid-TG clearance rates. It is also sensitive enough to detect physiological changes in TRL kinetics.

# 1. Introduction and Literature Review

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## 1.1 Introduction

This chapter aims to establish a scientific rationale for the experimental studies described in this thesis. It starts with a brief overview of lipoprotein classification and metabolism. This is followed by an in-depth review of very low density lipoprotein (VLDL) metabolism and regulation, with special emphasis on the effects of elevated concentrations of VLDL in the formation of atherogenic dyslipidaemia. In addition, the effects of hyperinsulinaemia and moderate exercise on VLDL production are also discussed. Finally, methods used for measuring VLDL kinetics are reviewed.

## 1.2 Lipoprotein Metabolism

### 1.2.1 Classification and Properties of Lipoproteins

Lipids, being insoluble in water, need to be combined with proteins for transport in the blood. The resulting particles are *lipoproteins* and the protein moieties are known as *apolipoproteins (apo)* (Gurr *et al.*, 2002). The Lipoprotein particle consists of a hydrophobic core of triglycerides (TG) and cholesteryl esters (CE), surrounded by a monolayer surface of the amphipathic (both hydrophilic and hydrophobic) phospholipids (PL), small amounts of free cholesterol and proteins (Ginsberg *et al.*, 2005). Lipoproteins differ according to the ratio of lipid to protein within the particle as well as the proportions of lipids, they form a continuum of particles varying in composition, size, density and function (Gurr *et al.*, 2002; Ginsberg *et al.*, 2005). It is convenient, therefore, to classify plasma lipoproteins into different density classes and to separate and isolate them by ultracentrifugation (Gurr *et al.*, 2002). These are: chylomicrons (CM), very low density lipoproteins (VLDL), intermediate density lipoproteins (IDL), low density lipoproteins (LDL) and high density lipoproteins (HDL) (**Table 1.1**). It should be noted, however, that these classes are not homogenous and there is a wide variety of particle size and chemical composition within each class. The major apolipoproteins are described in **Table 1.2**.

**Table 1.1: Composition and characteristics of the human plasma lipoproteins (Gurr *et al.*, 2002; Ginsberg *et al.*, 2005; Packard & Shepherd, 1997)**

Class	Density (g.ml <sup>-1</sup> )	Svedberg Floatation Rate (S <sub>f</sub> )	Diameter (nm)	Apolipoproteins	Function	Composition (%)			
						Protein	TG	CHOL*	PL
<b>Chylomicrons</b>	<0.95	>400	80-1000	AI, AII, AIV, B-48, CI, CII, CIII, E	Transport of dietary fat	2	80-95	2-7	3-9
<b>VLDL</b>	0.95-1.006	20-400	30-80	B-100, CI, CII, CIII, E	Transport of endogenous fat	8	55-80	5-15	10-20
<b>IDL</b>	1.006-1.019	12-20	25-35	B-100, CI, CII, CIII, E	-	19	20-50	20-40	15-25
<b>LDL</b>	1.019-1.063	0-12	18-25	B-100	Transport of CHOL to periphery	22	5-15	40-50	20-25
<b>HDL</b>	1.063-1.210	—	5-12	AI, AII, AIV, CI, CII, CIII, D, E	Reverse transport of CHOL	40	5-10	15-25	20-30

\* Free and esterified cholesterol (CHOL).

**Table 1.2: Characteristics of the major apolipoproteins (Gurr *et al.*, 2002; Ginsberg *et al.*, 2005)**

<b>Apolipoprotein</b>	<b>MW</b>	<b>Major site of synthesis</b>	<b>Function</b>
apo A-I	28 000	Liver, intestine	Structural component of HDL; activates LCAT
apo A-II	17 000	Liver, intestine	May inhibit HL activity; inhibits AI/LCAT
apo A-IV	44 500	Intestine	Activates LCAT; possibly facilitates transfer of apos between HDL and chylomicrons
apo A-V	39 000		Associated with lower TG levels; facilitates LPL
apo B-48	241 000	Intestine	Structural component of chylomicrons
apo B-100	513 000	Liver	Necessary for secretion of VLDL from liver; structural protein of VLDL, IDL and LDL; ligand for the LDL receptor
apo C-I	6 600	Liver	Activates LCAT; may inhibit hepatic uptake of chylomicrons and VLDL remnants; may inhibit CETP
apo C-II	8 800	Liver	Activates LPL (essential cofactor)
apo C-III	8 800	Liver, intestine	Inhibits LPL and hepatic uptake of chylomicrons and VLDL remnants
apo E	34 000	Liver (60-80%), other tissues including adipose tissue	Ligand for LDL receptor, LDL receptor-related protein and proteoglycans



Lipoproteins can also be classified according to their major lipids content into TG-rich lipoproteins (TRL), which include chylomicrons and VLDL, and cholesterol-rich lipoproteins, which include LDL and HDL (Sparks & Sparks, 1994).

Lipoprotein metabolism is a complex process to which new insights are continually being added. TRL metabolism is divided into the exogenous and endogenous lipid transport pathways. The exogenous pathway involves the delivery of dietary lipids (cholesterol, TG and PL) from the small intestine to the liver and peripheral tissues, while the endogenous pathway involves the delivery of lipids from the liver to peripheral tissue. As far as cholesterol metabolism is concerned, the exogenous and endogenous pathways are collectively known as forward cholesterol transport, whereas HDL metabolism is also known as the reverse cholesterol transport as it removes cholesterol from peripheral tissues.

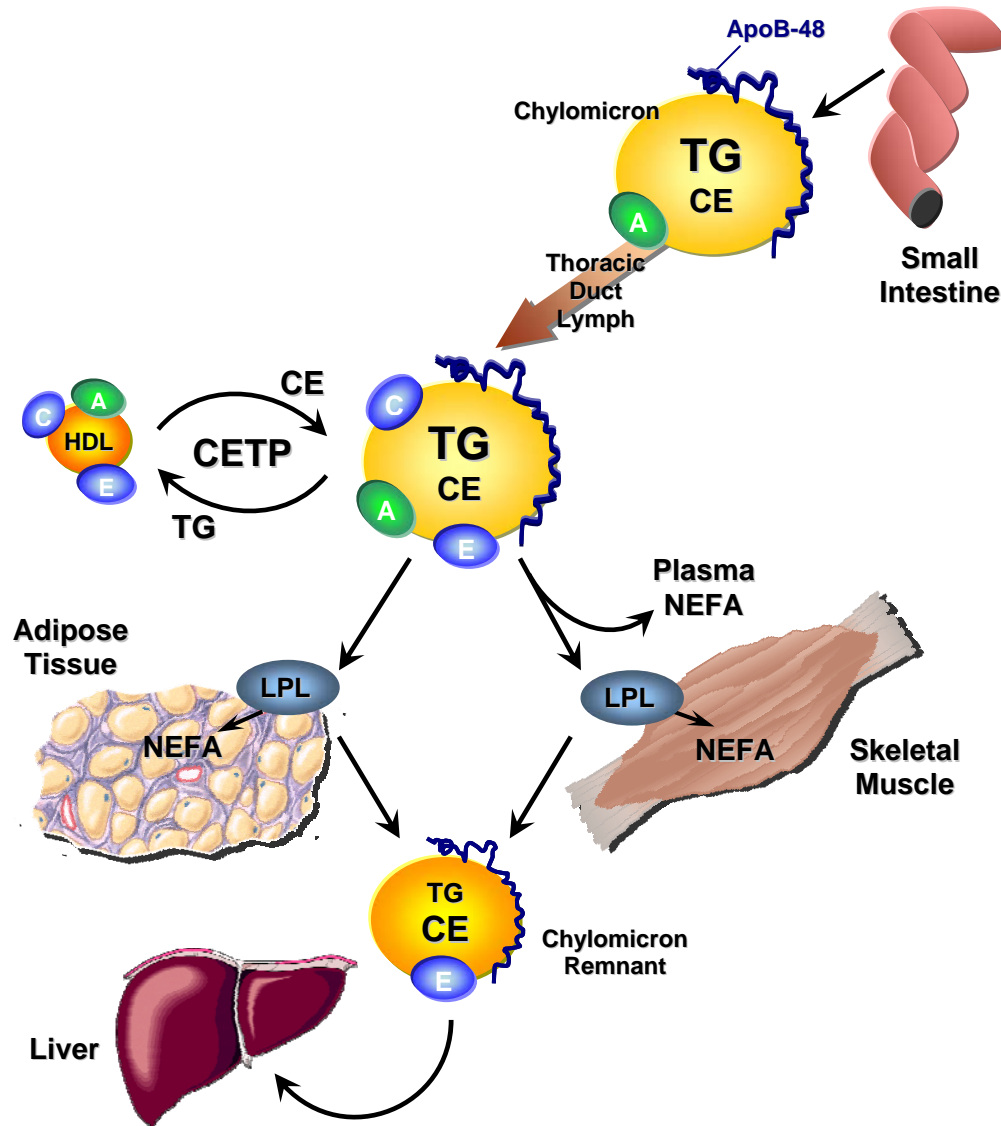
### **1.2.2 Exogenous Lipoprotein Metabolism**

After ingestion of a meal, dietary TG and cholesterol are absorbed into the small intestine cells where they are packed into nascent apoB-48-containing chylomicrons (Ginsberg *et al.*, 2005) with a Svedberg flotation rate ( $S_f$ ) of  $>400$  (Gurr *et al.*, 2002) (**Figure 1.1**). The newly synthesised chylomicrons enter the circulation through the lymphatic system and via the superior vena cava, where they acquire apoC-II, apoC-III and apoE (Ginsberg *et al.*, 2005). Chylomicrons also contain apoA-I and apoA-IV (Alaupovic, 1991). The TG content of chylomicrons is then hydrolysed by lipoprotein lipase (LPL) ([section 1.2.5](#)) situated on the capillary endothelium of predominantly adipose and muscle tissues (Fielding & Frayn, 1998; Goldberg, 1996). While apoC-II acts as activator for LPL (Olivecrona & Beisiegel, 1997), both apoC-III (Wang *et al.*, 1985) and apoE (Jong *et al.*, 1997) act to inhibit lipolysis by LPL. However, it is the balance of apoC-II and apoC-III that determines, in part, the efficiency with which LPL hydrolyses chylomicron TG (Ginsberg *et al.*, 2005; Chan *et al.*, 2008). The released non-esterified fatty acids (NEFA) from chylomicrons are either taken up by adipose tissues, re-incorporated into TG and stored, or by muscle tissue where they can be used for energy (Ginsberg *et al.*, 2005). Some dietary NEFA will also ‘spillover’ into the circulation (Barrows *et al.*, 2005). As a result of lipolysis, a new particle called ‘chylomicron remnant’ is formed with some attached LPL molecules (Saxena *et al.*, 1989). However, it should be noted that the majority

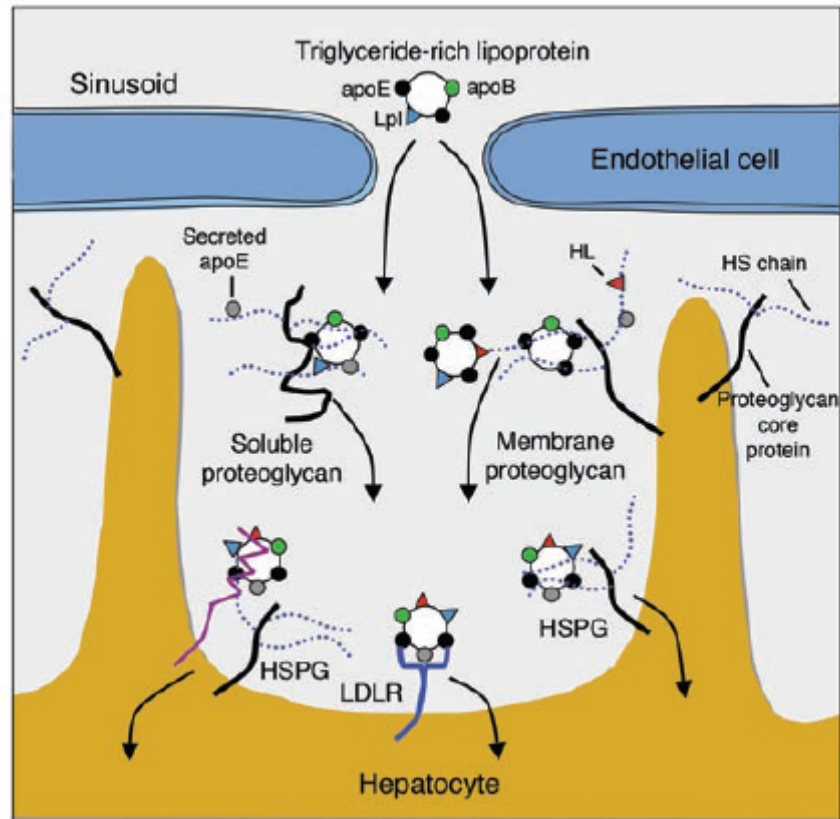
of chylomicrons do not form small chylomicron remnants, rather they show signs of being marginated to the vascular endothelium and removed as very large chylomicron remnants long before they reach the size of VLDL (i.e.  $S_f$  60-400 or the  $S_f$  20-60 range) (Karpe *et al.*, 1997). Chylomicron remnants are relatively enriched in CE, both diet- and HDL-derived, which is transferred in exchange for TG, mediated by cholesteryl ester transfer protein (CETP) (Cooper, 1997; Ko *et al.*, 1994; Chung *et al.*, 2004). They are also enriched in apoE and apoB and depleted of apoA-I and apoCs (Cooper, 1997; Ginsberg *et al.*, 2005). The normal physiological function of chylomicron remnants is to return bile cholesterol to the liver by an enterohepatic circulation (Redgrave, 2004), while its residual TG content is an important source of hepatic fatty acids (FA), accounting for ~73% of newly synthesised hepatic VLDL in mice (Jung *et al.*, 1999).

Hepatic uptake of remnant particles (**Figure 1.2**) is thought to start when they are sequestered in the liver perisinusoidal space (space of Disse), where they undergo further processing by hepatic lipase (HL) and LPL (both of which may remain associated with the particle) and acquire apoE. LPL, HL and apoE can potentially serve as ligands for a group of hepatic scavenger receptors, which endocytoses the particles, leading to their lysosomal catabolism. Endocytosis of remnant particles occurs via the apoB/apoE-recognizing LDL receptor (LDLR) or LDL receptor-related protein (LRP) in association with heparan sulfate proteoglycans (HSPGs) or independently by proteoglycans that are secreted into the space of Disse (MacArthur *et al.*, 2007; Havel & Hamilton, 2004; Mahley & Ji, 1999).

Another member of the LDLR family that functions as a peripheral apoE-recognising remnant lipoprotein receptor is the VLDL receptor (VLDLR). It is expressed abundantly in FA-active tissues (heart, skeletal muscle and adipose) and binds TRLs but not LDL (Takahashi *et al.*, 2004). It is likely that VLDLR functions in concert with LPL and has been reported to play a major role in chylomicron metabolism by enhancing LPL-mediated TG hydrolysis in mice (Goudriaan *et al.*, 2004).



**Figure 1.1: Chylomicron Metabolism.** Dietary triglycerides (TG) and cholesteryl esters (CE) are absorbed into the small intestine and packed as chylomicrons, which also contain apoB-48 and apoA-I proteins. Chylomicrons are secreted into the circulation through the lymphatic system, where they acquire apoCs and E (apoC-II acts as an activator for the lipoprotein lipase enzyme (LPL) and apoE is needed for receptor recognition by the liver). Chylomicron-TG is hydrolysed by LPL situated on the capillary endothelium of mainly adipose tissues and skeletal muscles, providing the underlying tissue with non-esterified fatty acids (NEFA). Some dietary NEFA may ‘spillover’ in the circulation during hydrolysis by LPL. Chylomicron-TG is also removed from the particle in exchange for CE from high density lipoprotein (HDL). This is catalysed by the action of cholesteryl ester transfer protein (CETP). The result is a slightly smaller and CE-enriched particle called ‘chylomicron remnant’. It is also depleted in apoCs and apoA-I. The majority of chylomicron remnants are removed from plasma long before they reach the size of VLDL particles range. Remnant particles are taken up by the liver by a number of receptors (see **Figure 1.2**).



**Figure 1.2: Model of possible hepatic uptake of TRL remnants.** Hepatocytes and endothelial cells produce membrane-bound heparan-sulfate (HS) proteoglycans (HSPGs) and secrete proteoglycans into the space of Disse. After lipolytic processing of lipoproteins in the circulation by lipoprotein lipase (LPL; blue triangles), apoE-enriched (black circles) remnant lipoproteins enter the space of Disse through fenestrations in the endothelium. Remnant lipoproteins are thought to be sequestered near the hepatocyte cell surface via apoE-HS binding or lipase-HS bridging on secreted HSPGs. Lipoproteins are further processed in the space of Disse by transfer of soluble apoE (grey circles) and by hepatic lipase (HL; red triangles) bound via HS. ApoE, HL, and LPL can potentially serve as ligands of TRLs. Endocytosis of lipoprotein particles occurs via LDL receptor (LDLR; blue) or LDL receptor-related protein (LRP; purple) in association with HSPGs or independently by proteoglycans. [Figure and legend reproduced with permission from *Journal of Clinical Investigation* (MacArthur *et al.*, 2007).]

### 1.2.3 Endogenous Lipoprotein Metabolism

The major bulk of endogenous plasma TG is carried by VLDL, which is continually synthesised and secreted by the liver (Packard & Shepherd, 1997). The VLDL particle contains one molecule of apoB-100 (Elovson *et al.*, 1988) and approximately 5-25,000 TG molecules in its core (Bjorkegren *et al.*, 1998). It also contains apoCs and apoE (Packard & Shepherd, 1997). There are at least two major subclasses of VLDL particles: large TG-rich VLDL<sub>1</sub> (S<sub>f</sub> 60-400) and smaller, more dense VLDL<sub>2</sub> (S<sub>f</sub> 20-60), which have more cholesterol and a lower ratio of apoCs and apoE to apoB (Packard & Shepherd, 1997; Packard *et al.*, 1984). In normolipidaemic subjects, about 75% of the variation in plasma TG seems to be determined by the VLDL<sub>1</sub> concentration (Tan *et al.*, 1995). Once in the circulation, the TG content of VLDL<sub>1</sub> is rapidly removed by LPL-mediated hydrolysis (Karpe *et al.*, 2007) and the action of CETP, which facilitates the transfer of TG from TRLs (mainly VLDL) to HDL in exchange for CE from HDL to apoB-containing lipoproteins (VLDL, IDL and LDL) (Stein & Stein, 2005; Ko *et al.*, 1994). As VLDL<sub>1</sub> particles become progressively depleted in TG, they may either attain the same size and TG content as VLDL<sub>2</sub> particles, or continue its delipidation to form IDL. This is further aided by the lipolytic action of HL, which is also involved in the hydrolysis of IDL-TG. As a result, LDL appears as the terminal particle (Packard *et al.*, 1984). During the transition of VLDL to IDL and IDL to LDL, a portion of the surface PL, apoCs and apoE are transferred to HDL (Verges, 2005).

Approximately 10% of VLDL<sub>1</sub> particles undergo complete delipidation to LDL, while the remaining cease delipidation in the VLDL or IDL density range forming remnant particles that persist in the circulation for considerable periods before being removed from plasma (Packard & Shepherd, 1997; Packard *et al.*, 1984). On the other hand, almost all VLDL<sub>2</sub> particles will be effectively delipidated to LDL (Gaw *et al.*, 1995; Packard *et al.*, 1984). Similarly, not all IDL particles are delipidated to LDL, some are cleared by the liver (Packard & Shepherd, 1997) along with TRL remnant particles, possibly by the clearance pathway as chylomicron remnants (see **Figure 1.2**).

LDL is the main cholesterol-carrying lipoprotein in plasma with each particle containing one molecule of apoB-100. Generally about 65% of the total plasma

cholesterol is carried in LDL in man (Wang & Briggs, 2004) and its major role is to supply tissues with cholesterol, which is paramount for every cell for the synthesis and maintenance of cell membranes. LDL particles are removed from plasma by the LDLR with LDL<sub>2</sub> particle exhibiting the highest affinity (Packard & Shepherd, 1997). While 70% of the LDLRs are located on hepatic cells, 30% are located on the other cells of the body (Verges, 2005).

#### **1.2.4 HDL Metabolism**

The movement of lipids via chylomicrons, VLDL, IDL and LDL provides the peripheral cells with FA (fuel) and cholesterol (necessary for synthesis and maintenance of cell membranes). While this ‘forward’ movement of TG is balanced by the metabolic consumption of NEFA as a source of fuel, there is no analogous (i.e. catabolic) mechanism for the metabolic disposal of cholesterol. This is accomplished by ‘reverse’ cholesterol transport by HDL particles (Tulenکو & Sumner, 2002). Over the last several decades, epidemiological studies have revealed that levels of LDL-cholesterol are directly associated with risks of CVD, while an inverse relationship existed between HDL-cholesterol and the risk of the disease (Wang & Briggs, 2004).

HDL particles are synthesised and secreted, mainly by the liver, as nascent or lipid-poor disc-shaped particles (pre-β HDL) containing apoA-I and small amount of PL (Wang & Briggs, 2004; Tulenko & Sumner, 2002). This lipid-poor apoA-I avidly absorbs free cholesterol and PL from peripheral cells mediated by ATP-binding cassette A1 (ABCA1) transporter on the ‘donor’ cells plasma membrane (Wang & Briggs, 2004). The free cholesterol, being amphipathic, is absorbed onto the surface of the small HDL, which is progressively esterified by lecithin cholesterol acyltransferase (LCAT; bound to HDL and activated by apo A-I) and stored in the core of HDL (Tulenکو & Sumner, 2002; Wang & Briggs, 2004). The continuous addition of cholesterol converts pre-β HDL to larger spherical HDL<sub>3</sub>. Together with the action of phospholipid transfer protein (PLTP) which transfers phospholipids to HDL (mostly HDL<sub>3</sub>) particles from VLDL/LDL (van Tol, 2002), HDL<sub>3</sub> is converted to the much larger HDL<sub>2</sub> particles (Wang & Briggs, 2004; Verges, 2005). This stage of cholesterol transfer from cells to the HDL<sub>3</sub> is mediated either by cell-surface receptors scavenger receptor class B type I (SR-BI) or passive diffusion, both distinct

from that mediated by ABCA1 (Wang & Briggs, 2004). In addition, HDL accumulates apoC-II and apoE from VLDL and IDL, which serve as a reservoir of apolipoproteins, especially apoC-II which is needed for the activation for LPL (Tulenکو & Sumner, 2002).

The TG content of HDL is hydrolysed by HL, while removal of cholesterol from HDL<sub>3</sub> particles occurs through CETP-mediated neutral lipid (TG and CE) exchange between HDL and apoB-containing lipoproteins. Because the bulk of cholesterol removal is accomplished by shuttling to these particles, their uptake by the liver is essential for the disposal of HDL cholesterol (Tulenکو & Sumner, 2002; Wang & Briggs, 2004). The degradation of the larger HDL<sub>2</sub> occurs directly by CE-selective uptake by the liver mediated by SR-B1. The apoA-I from the degradation is either recycled for new HDL formation or cleared by the cubulin, a receptor highly expressed in kidney and yolk sac (Wang & Briggs, 2004).

**Figure 1.3** summarises the endogenous lipoprotein metabolism.

### **1.2.5 Lipoprotein Lipase**

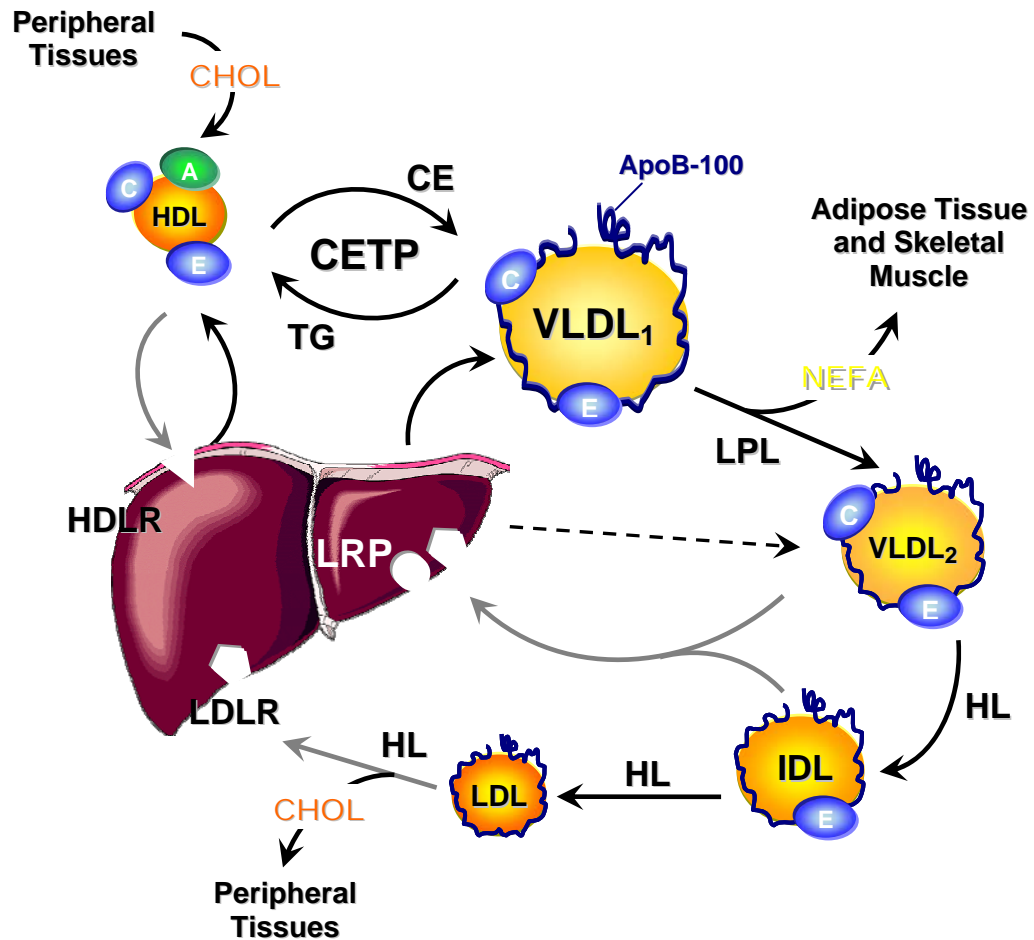
Lipoprotein lipase (*EC* 3.1.1.34) is a key enzyme for the hydrolysis of circulating TG (Fielding & Frayn, 1998; Preiss-Landl *et al.*, 2002). It is synthesized by parenchymal cells, then transferred to the luminal surface of endothelial cells, where it is bound to HSPG (Goldberg, 1996). LPL is distributed in a variety of tissues, with the highest concentrations occurring in adipose tissue and muscle (Wong & Schotz, 2002) [cardiac and skeletal (Preiss-Landl *et al.*, 2002)]. However, substantial LPL activity is also detectable in differential macrophages, brain, placenta, lung, spleen, pancreatic  $\beta$ -cells and steroidogenic tissue (Preiss-Landl *et al.*, 2002). LPL functions to supply the underlying tissue with FAs derived from the TG-rich core of circulating chylomicrons and VLDL (Olivecrona *et al.*, 1997). However, chylomicrons have been shown to be the preferred substrate for LPL (Potts *et al.*, 1991; Fisher *et al.*, 1995), as they may bind to LPL with an affinity of 50 times higher than VLDL (Xiang *et al.*, 1999). In addition, it has been shown that about forty LPL molecules may act on a TRL particle simultaneously to achieve the rates of TG hydrolysis observed (Scow & Olivecrona, 1977). After hydrolysis of TG, the TG-depleted particle detaches along with some LPL molecules that may dissociate from the

endothelium and leave attached to the remnant particle (Goldberg, 1996; Fielding & Frayn, 1998). There is a continual turnover of LPL at the endothelial site of action, with replacement of the dissociated LPL molecules by newly secreted molecules from within the tissue. LPL circulating attached to lipoprotein particles may play an important role in their eventual receptor-mediated uptake (Fielding & Frayn, 1998) (see **Figure 1.2**).

The apoC-II associated with the TRL particle acts as an activator of the enzyme (Goldberg, 1996; Olivecrona & Beisiegel, 1997) and apoE, which is a strong heparan-binding protein, anchors TRLs to the HSPG on the surface of the endothelial cells (Goldberg, 1996). In contrast, apoC-III is the principal plasma inhibitor of VLDL lipolysis; it inhibits LPL as well as interfering with remnant lipoprotein clearance (Shachter, 2001; Havel *et al.*, 1973).

The regulation of LPL is tissue-specific in such a way that its expression correlates highly with both the need for, and the uptake of, lipid fuels by the tissue. For instance, in the fed state, when energy is abundant, LPL is downregulated in skeletal muscle and heart and activated in white adipose tissue which facilitates the lipolysis of TRL-TG in the latter tissue, directing released NEFA for esterification and storage (Fielding & Frayn, 1998). On fasting, however, the situation is reversed, with suppression in adipose tissue and upregulation in muscle, increasing TRL-TG lipolysis in muscle, so that the resultant NEFAs are directed to the tissue in which they are needed as an oxidative fuel (Fielding & Frayn, 1998; Ruge *et al.*, 2005). Such changes in LPL expression are mediated predominantly through the action of hormones, such as insulin, glucocorticoid, and adrenaline (Mead *et al.*, 2002) as well as cytokines, fatty acids and glucose (Preiss-Landl *et al.*, 2002). Insulin is of particular interest because it plays a central role linking dyslipidaemia and insulin resistance, the defining feature of the metabolic syndrome (Howard, 1999). This will be discussed further in [section 1.5](#).





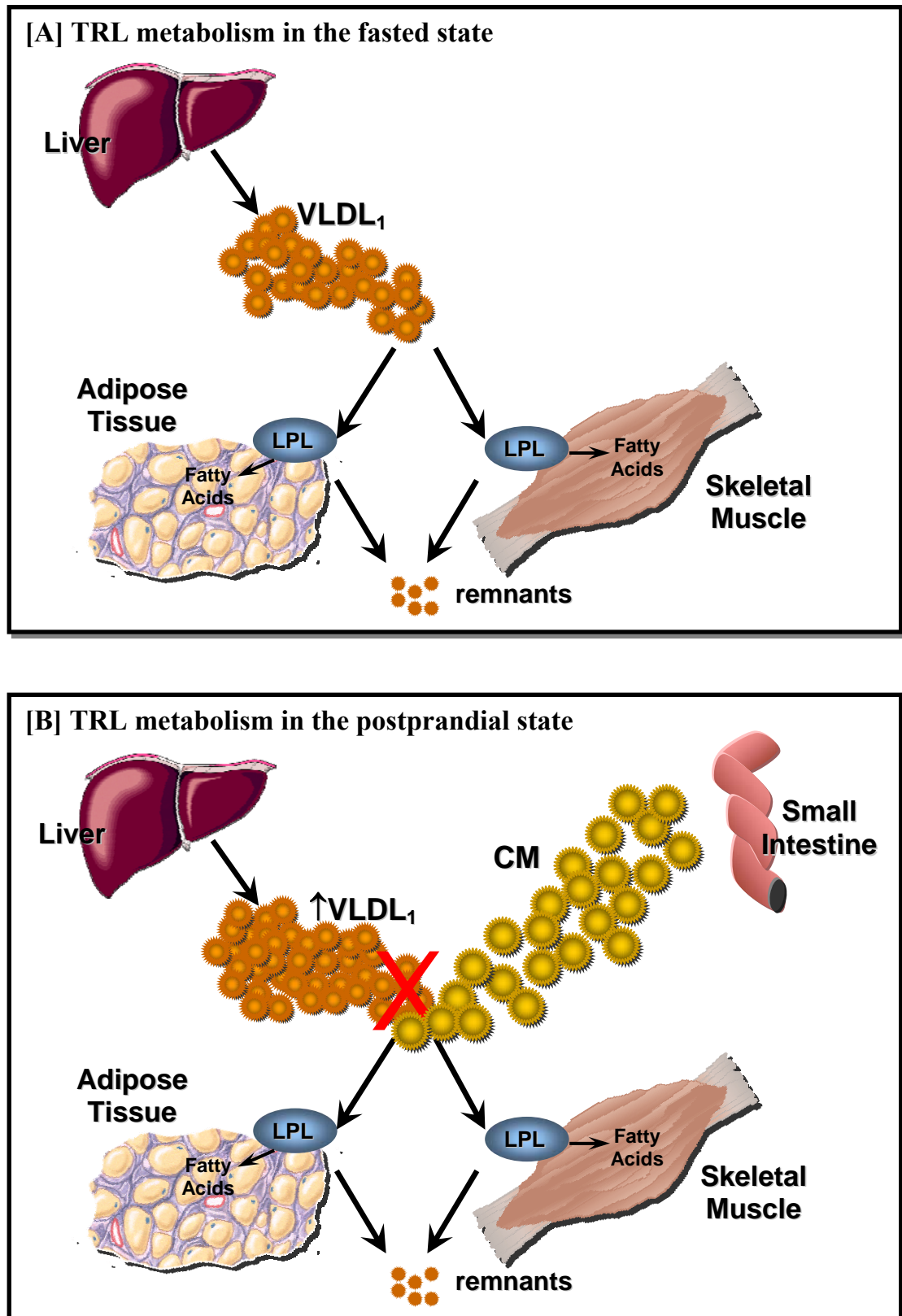
**Figure 1.3: Overview of the general aspects of endogenous lipoprotein metabolism.** The grey lines represent hepatic uptake of lipoproteins and lipoprotein remnants. The major bulk of endogenous triglycerides (TG) is secreted by the liver in the form of VLDL<sub>1</sub> particles, of which apoB-100 protein forms an integral part on the surface of the particle. Once in the circulation, TG in the core of VLDL<sub>1</sub> is hydrolysed by the action of lipoprotein lipase (LPL) which provides non-esterified fatty acids (NEFA) to adipose tissue and skeletal muscle. VLDL<sub>1</sub> also loses some of its TG content to HDL in exchange for cholesteryl esters (CE) via the action of cholesteryl ester transfer protein (CETP). The continual removal of TG from VLDL<sub>1</sub> gives rise to smaller VLDL<sub>2</sub>, which is also produced by the liver. VLDL<sub>2</sub> is either taken up by the liver or delipidated further by the action of hepatic lipase (HL) to form IDL then LDL. Unlike other apolipoproteins, apoB-100 stays with the particles as they are delipidated. LDL is the main cholesterol (CHOL)-carrying lipoprotein which provides cholesterol to peripheral tissues. HDL, on the other hand, is also secreted by the liver and acts as a 'reverse' cholesterol carrier, from peripheral tissues to the liver. LDL is taken up by the liver by the apoB/E recognising LDL receptor (LDLR). Remnant (smaller) lipoprotein particles are also taken up by the liver in a similar manner to chylomicron remnants (Figure 1.2) by hepatic receptors, such as LDLR and LDL receptor-related protein (LRP).

### 1.2.6 Integration of TRL Metabolism

In the fasted state, the continually secreted VLDL (particularly the larger VLDL<sub>1</sub>) particles by the liver are the major substrates for LPL (Fisher *et al.*, 1995), providing endogenous fat to peripheral cells. However, after ingestion of a meal, the newly formed chylomicrons, being the preferred substrate for LPL (Potts *et al.*, 1991; Fisher *et al.*, 1995), will compete with VLDL<sub>1</sub> for the same LPL-mediated catalytic pathway (Brunzell *et al.*, 1973; Karpe *et al.*, 1993a). As the percent TG hydrolysed by LPL in TRL subfractions was chylomicrons > VLDL<sub>1</sub> > VLDL<sub>2</sub>, it was proposed that increasing the size and TG content of a lipoprotein particle increases its susceptibility to hydrolysis by LPL (Fisher *et al.*, 1995). Another possible explanation for this preferential lipolysis of chylomicrons compared with VLDL might be due to transient changes in VLDL<sub>1</sub> composition in the postprandial state. Both VLDL<sub>1</sub> and VLDL<sub>2</sub> are enriched in cholesterol, apoE, apoC-I and apoC-III in the postprandial state, but depleted in apoC-II (Bjorkegren *et al.*, 1997). While apoC-II acts as activator for LPL (Olivecrona & Beisiegel, 1997), both apoC-III (Chan *et al.*, 2008) and apoE (Jong *et al.*, 1997) act to inhibit lipolysis by LPL.

As a result of this competition, the delipidation of VLDL<sub>1</sub> particles is delayed and their concentration is increased as they accumulate in plasma after the appearance of chylomicrons (Karpe *et al.*, 1993a) or chylomicron-like particles (Karpe & Hultin, 1995; Bjorkegren *et al.*, 1996) (**Figure 1.4**). Indeed, the postprandial increase in the TRL particle number is mainly accounted for by VLDL, particularly VLDL<sub>1</sub> (Karpe *et al.*, 1993a). VLDL<sub>2</sub> particles, on the other hand, are not affected by this competition and, in fact, their concentrations are often lowered or unchanged postprandially (Bjorkegren *et al.*, 1996; Karpe *et al.*, 2007). This is because, firstly, less VLDL<sub>1</sub> is converted into VLDL<sub>2</sub> and, secondly, VLDL<sub>2</sub> is not a particularly good substrate for LPL (Fisher *et al.*, 1995; Karpe *et al.*, 2007).

This competition and the preferential lipolysis by LPL between chylomicron-like particles and VLDL<sub>1</sub> form the basis of work described in this thesis.



**Figure 1.4: TRL metabolism in the [A] fasted and [B] postprandial state.** VLDL<sub>1</sub> and chylomicrons (CM) are both cleared by lipoprotein lipase (LPL), situated on the capillary endothelium of adipose tissue and skeletal muscle. CMs are the preferred substrate for LPL. Thus, the presence of CM (or CM-like particles) inhibits VLDL<sub>1</sub> clearance by this pathway, causing VLDL<sub>1</sub> to accumulate in the circulation.

### 1.3 Hepatic Assembly and Secretion of VLDL

The assembly of VLDL in the liver involves a number of complex processes which are still not fully understood. VLDL assembly is believed to take place in two major steps within two different compartments of the cell, rough endoplasmic reticulum (ER) and Golgi apparatus, and involves three different particles, pre-VLDL, VLDL<sub>2</sub> and VLDL<sub>1</sub> (Olofsson & Borén, 2005). It commences during the synthesis of apoB which is an integral part of the VLDL structure.

#### 1.3.1 Apolipoprotein B Structure

Unlike other plasma lipoproteins, apoB is an amphipathic high molecular weight glycoprotein which is insoluble in aqueous solutions (Segrest *et al.*, 2001; Kane, 1983). Because it binds irreversibly to TRLs and LDL, it is not transferable between lipoproteins (Kane, 1983; Davidson & Shelness, 2000). There are two forms of apoB in mammals, which are encoded by the same gene (Davidson & Shelness, 2000). The full-length apoB, apoB-100, consists of 4536 amino acids, whereas apoB-48 corresponds to 48% of the protein from the N-terminal (Davidson & Shelness, 2000). ApoB-48 is produced by a site-specific cytidine-to-uridine RNA editing reaction which converts a glutamine codon to a stop codon causing translational termination of (intestinal) apoB mRNA at residue 2152 (Davidson & Shelness, 2000; Olofsson & Borén, 2005). In humans, apoB-100 is expressed in the liver, forming VLDL, while apoB-48 is expressed in the small intestine, forming chylomicrons.

The apoB protein has a pentapartite structure consisting of one globular N-terminal structure, followed by two amphipathic  $\beta$ -sheets alternating with two amphipathic  $\alpha$ -helical domains (NH<sub>2</sub>- $\beta\alpha_1$ - $\beta_1$ - $\alpha_2$ - $\beta_2$ - $\alpha_3$ -COOH) (Segrest *et al.*, 2001). The  $\beta\alpha_1$  domain (the N-terminal 1000 residues) is a globular structure that bears structural homology with the lipid-binding pocket of microsomal TG transfer protein (MTP) (an ER luminal protein with lipid transfer activity – see below) and the lamprey vitellogenin (an egg yolk lipoprotein) (Segrest *et al.*, 2001). Thus, apoB, MTP and vitellogenin are considered members of the same gene family collectively known as large lipid transfer proteins (LLTPs). Although vitellogenin was presumed to be the

ancestral member, recent evidence suggests that MTP may be the oldest LLTP family member (Shelness & Ledford, 2005).

The amphipathic  $\beta$ -sheet domains of apoB interact irreversibly with the neutral lipid core, while the  $\alpha$ -helical domains, which are similar to those found in soluble apolipoproteins, may form strong but reversible binding to the lipoprotein surface (Davidson & Shelness, 2000). The  $\beta_2$ -region contains the LDL receptor-binding domain, needed for binding and hepatic endocytosis of LDL particles, and resembles the receptor binding domain of apoE, another LDLR ligand. As this domain is in the truncated C-terminal region of apoB-48, chylomicrons depend on apoE to bind to the LDLR (Segrest *et al.*, 2001).

### **1.3.2 Assembly of VLDL**

The intracellular assembly of VLDL (**Figure 1.5**) starts in the rough ER during the biosynthesis of apoB, which is lipidated co-translationally; i.e. while the C-terminal portion of apoB is still being synthesised on the ribose of the ER, the N-terminal portion is translocated across the ER and is assembled as a small lipoprotein particle (Olofsson & Borén, 2005; Dashti *et al.*, 2007). ApoB interacts co-translationally with MTP, which catalyses the addition of lipids [in the order of TG > CE > diacylglycerol > cholesterol > PL (Rava *et al.*, 2005)] to the growing apoB molecule. This results in the formation of a primordial HDL-sized lipoprotein particle, pre-VLDL, which is retained in the cell by interaction with chaperones and other ER proteins (Olofsson & Borén, 2005; Shelness & Sellers, 2001). It has been recently shown that MTP is not required for the initiation of this step as the first N-terminal 1000 amino acid residue (apoB:1000) has the capacity to fold forming a PL-rich 'lipid-pocket', independently of the structural requirement and lipid-transfer activity of MTP (Manchekar *et al.*, 2004; Dashti *et al.*, 2007). It is currently unclear at which point in VLDL assembly that MTP is required. In addition to its lipid transfer activity, MTP facilitates the co-translational translocation of apoB across the ER lumen (Blasiole *et al.*, 2007) and prevents its degradation as apoB will be sorted to proteasomal degradation if it misfolds or is underlipidated (Fisher & Ginsberg, 2002; Hussain *et al.*, 2003). Pre-VLDL is lipidated further to become VLDL<sub>2</sub>, or sorted to degradation. This conversion of pre-VLDL to VLDL<sub>2</sub> may explain why the MTP

activity is needed for the secretion of apoB-100 after the translation is completed (Rustaeus *et al.*, 1998; Olofsson & Borén, 2005).

The apoB-100 containing VLDL<sub>2</sub> particle exits the ER at specific exit sites carried by two vesicle proteins: a GTPase referred to as SAR1 and a coat protein called coatamer protein II (COPII) (Gusarova *et al.*, 2003; Olofsson & Borén, 2005), which fuse to become ER-Golgi intermediate compartment (ERGIC). It is converted to *bona fide* VLDL<sub>1</sub> by post-translational acquisition of TG in the Golgi apparatus (Shelness & Sellers, 2001; Olofsson & Borén, 2005). This lipidation differs from that which gives rise to VLDL<sub>2</sub> and requires that apoB has reached the size of apoB-48. In addition, it is dependent on ADP ribosylation factor 1 (ARF-1), a small GTP binding protein which plays a role in the membrane trafficking between the ER and the Golgi apparatus (Asp *et al.*, 2000; Asp *et al.*, 2005). It has been suggested that this ‘maturation’ step of VLDL<sub>1</sub> may involve the formation of lipid droplets in the lumen of the secretory pathway, which fuse with apoB to form VLDL<sub>1</sub> (Olofsson & Borén, 2005). Further, this conversion of VLDL<sub>2</sub> to VLDL<sub>1</sub> seems an additional step for TG secretion from the liver as it is not necessary for the secretion of apoB (Olofsson & Borén, 2005). The liver has been shown to secrete both VLDL<sub>1</sub> and VLDL<sub>2</sub> particles (Packard *et al.*, 2000). Interestingly, this stepwise assembly and secretion of VLDL explains the time delay between the biosynthesis of apoB-100 and the major addition of lipids to the VLDL<sub>1</sub> particles, which was estimated as ~15 min (Adiels *et al.*, 2005a).

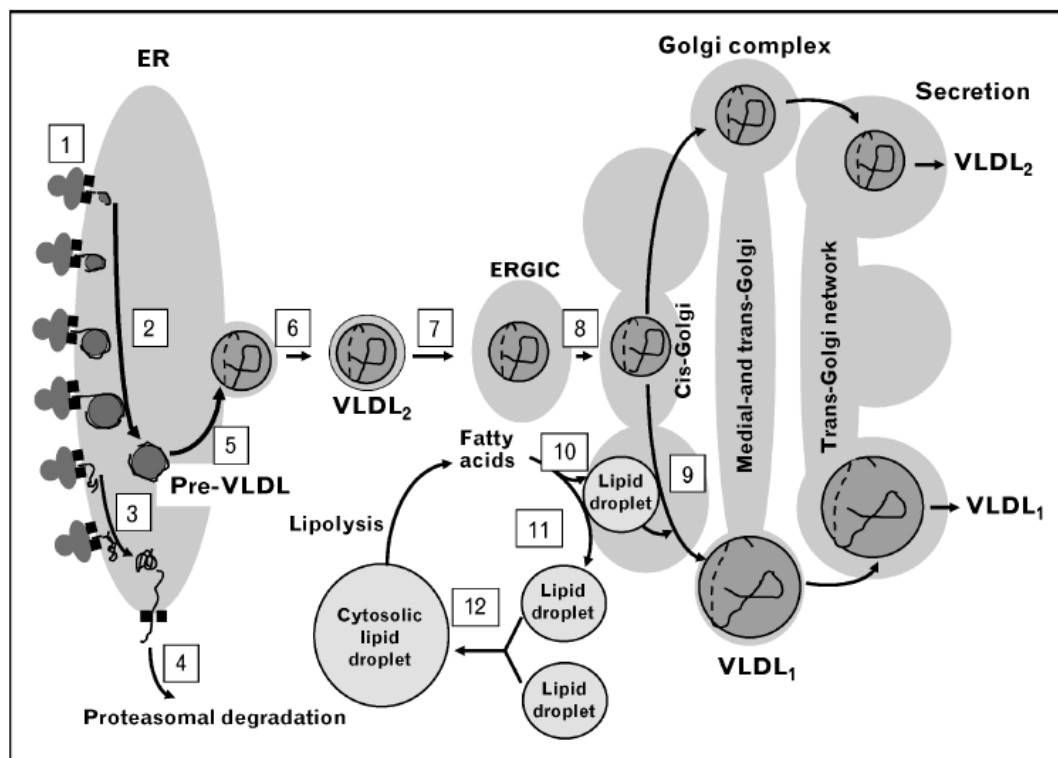
## **1.4 Regulation of VLDL Production**

The secretion of VLDL has been shown to be mainly dependent on fatty acids and insulin.

### **1.4.1 Role of Fatty Acids**

The liver has the capacity to store TG in order to accommodate plasma fatty acids, which are surplus to immediate energy requirements, thereby neutralising any potential ‘lipotoxicity’ to peripheral tissue. There are three potential sources of fatty acids which enter the hepatic TG storage pool: *de novo* lipogenesis (DNL), plasma NEFA [originating from adipose tissue and ‘spillover’ from dietary fat lipolysis (Barrows *et al.*, 2005)] and remnant lipoproteins (Gibbons *et al.*, 2000). A healthy

liver uses different sources of FAs in the fasted and fed states (Barrows & Parks, 2006). It is estimated that during fasting, ~77% of VLDL-TG is derived from recycling adipose FAs and ~4% from DNL. With feeding, 44% come from adipose FAs, 15% from uptake of chylomicron remnants, 10% from dietary NEFA spillover into plasma and 8% from DNL (Barrows & Parks, 2006), with dietary FAs appearing in VLDL within 90 min of food intake (Heath *et al.*, 2003).



**Figure 1.5: The Intracellular assembly of VLDL.** (Numbers in parentheses refer to corresponding numbers in the figure.) The assembly process starts in the rough endoplasmic reticulum (ER) by the biosynthesis and concomitant (cotranslational) translocation of apoB-100 to the lumen of this organelle (1). ApoB-100 interacts cotranslationally with the microsomal triglyceride transfer protein (MTP) and is lipidated to form a primordial particle (pre-VLDL) (2). Alternatively, apoB-100 fails to be lipidated and misfolds resulting to degradation (3). The protein is unfolded and retracted to the cytosol, ubiquitinated, and sorted to proteasomal degradation (3 and 4). Pre-VLDL is converted to VLDL<sub>2</sub> late in the ER compartment (5). VLDL<sub>2</sub> exits the ER at specific exit sites of this organelle by Sar1/Cop II vesicles (6), which fuse to become the ER-Golgi intermediate compartment (ERGIC) (7). ERGIC fuses with Cis-Golgi (8). In the Golgi apparatus, VLDL<sub>2</sub> is converted to VLDL<sub>1</sub> by the addition of a bulk load of triglycerides (9). This lipidation process differs from that which gives rise to pre-VLDL and VLDL<sub>2</sub>. The formation of VLDL<sub>1</sub> may involve the formation of a lipid droplet in the lumen of the secretory pathway (10), the mechanism of which may follow that of cytosolic lipid droplets (11 and 12). The formation of the cytosolic droplets also involves a fusion step (12). [Figure and legend with permission from *Current Opinion in Lipidology* (Adiels *et al.*, 2006a).]

It has been suggested that this hepatic TG pool is metabolically active with a rapid turnover such that extracellular FAs entering the cell are either oxidised in the mitochondrion or esterified into TG in the ER and then transferred to the cytosolic storage pool. TG required for VLDL synthesis is recruited from this pool by a process of lipolysis to give FAs (Gibbons *et al.*, 2000). This may explain the findings of Malmstrom *et al.* (1999) who reported that acute NEFA flux is not accompanied by concomitant changes in the production of total VLDL-apoB (Malmstrom *et al.*, 1999). In fact, reducing NEFA concentrations using acipimox (an antilipolytic agent) caused a shift from VLDL<sub>1</sub> to VLDL<sub>2</sub> production, without affecting total VLDL-apoB production (Malmstrom *et al.*, 1998), suggesting an important role of fatty acid availability for the assembly and secretion of VLDL<sub>1</sub>.

Conversely, earlier studies which did not isolate VLDL subfractions indicate that acutely raising plasma NEFA concentrations could increase both VLDL-TG and VLDL-apoB production (Lewis *et al.*, 1995), thus suggesting a role for FAs in the stimulation of TG and apoB synthesis. This assumption is supported by animal studies, where chronic administration of intravenous albumin-bound oleic acid doubled hepatic secretion of apoB-100 (without changes in apoB mRNA levels), suggesting that FAs might act as both a stimulus for TG synthesis, thereby driving assembly of VLDL, and as a 'signal' for apoB assembly (Zhang *et al.*, 2004b). Indeed, increased liver fat has been shown to increase the secretion of VLDL<sub>1</sub>-apoB-100, and more – but not larger – VLDL<sub>1</sub> particles are secreted (Adiels *et al.*, 2006b).

#### **1.4.2 Role of Insulin**

Despite insulin's induction of lipogenesis in the liver, it acutely suppresses hepatic VLDL production in normal weight, healthy subjects (Lewis *et al.*, 1993) – specifically affecting the VLDL<sub>1</sub> subclass (Malmstrom *et al.*, 1998; Adiels *et al.*, 2007). This effect is partly attributed to its induced reduction of NEFA flux to the liver, thus decreasing substrate availability for VLDL assembly (Coppack *et al.*, 1994). However, other mechanisms are likely to play a role as this suppressive effect of insulin on VLDL<sub>1</sub> production persisted even when compared with the action of acipimox (which caused a similar reduction in NEFA) (Malmstrom *et al.*, 1998). Although these mechanisms are still elusive, some have been investigated. Insulin inhibits the secretion of apoB (Sparks & Sparks, 1994) and insulin signalling (via



phospho-inositide 3-kinase) inhibits the maturation phase of VLDL assembly by preventing bulk lipid transfer to a VLDL precursor, thus enhancing the degradation of apoB (Brown & Gibbons, 2001). In addition, it inhibits transcription (Lin *et al.*, 1995) and upregulation (Wolfrum & Stoffel, 2006) of the MTP gene, although it has been suggested that this is unlikely to account for the acute effect of insulin on VLDL production due to the long half-life of the MTP protein (4.4 days) (Lin *et al.*, 1995; Blasiolo *et al.*, 2007).

In contrast, chronic hyperinsulinaemia, caused by insulin resistance, is associated with increased production of VLDL (Malmstrom *et al.*, 1997a), specifically VLDL<sub>1</sub> (Adiels *et al.*, 2007). Possible mechanisms for this loss of the suppressive effect of insulin have been studied in a fructose-fed hamster model of insulin resistance (Taghibiglou *et al.*, 2000). These included increased cellular secretion of total and VLDL-TG, enhanced stability of nascent apoB, thereby reducing its post-translational degradation and increased MTP mass (Taghibiglou *et al.*, 2000). In addition, insulin resistance also leads to loss of the acute insulin-mediated inhibition of apoB secretion (Chirieac *et al.*, 2004). In humans, central obesity and an increased liver fat content, which are usually associated with insulin resistance conditions, have been associated with the lack of insulin-induced suppression of VLDL<sub>1</sub> production (Adiels *et al.*, 2007) (see below).

### **1.5 Defective TRL Metabolism and Atherogenic Dyslipidaemia**

The concept of atherogenicity of TG and TRL, especially in the postprandial state, was first introduced almost 30 years ago by Zilversmit (Zilversmit, 1979). Since then, an increasing body of evidence has implicated hypertriglyceridaemia as an independent risk factor for cardiovascular disease (CVD), particularly coronary heart disease (CHD) (Cullen, 2000). TRL are thought to have a role in the development and progression of atherosclerosis (Tanaka *et al.*, 2001; Cullen, 2003) as TRL remnants have been found in human aortic atherosclerotic plaques (Nakano *et al.*, 2008). In addition, elevated concentrations of VLDL<sub>1</sub> particles are believed to start a chain of metabolic events that generate an ‘atherogenic lipoprotein phenotype’, also known as atherogenic dyslipidaemia.

### **1.5.1 Atherogenic Lipoprotein Phenotype**

The atherogenic lipoprotein phenotype is a cluster of lipoprotein abnormalities associated with insulin resistant states (Ginsberg *et al.*, 2005), such as obesity (Bamba & Rader, 2007; Marsh, 2003; Chan *et al.*, 2004a), type 2 diabetes (Marsh, 2003; Taskinen, 2003) and the metabolic syndrome (Grundy, 2006; Kolovou *et al.*, 2005), and significantly contributes to increased risk of CVD (Avramoglu *et al.*, 2006; Watson *et al.*, 2003).

Such atherogenic dyslipidaemia manifests as increased concentrations of hepatic VLDL (causing hypertriglyceridaemia), small dense LDL (sdLDL), TRL remnants and low HDL concentrations (Grundy, 2006; Taskinen, 2003; Avramoglu *et al.*, 2006). Importantly, it is the excess hepatic production of VLDL<sub>1</sub> [the major determinant of plasma TG concentration (Hiukka *et al.*, 2005)] but not VLDL<sub>2</sub> particles (Adiels *et al.*, 2005b; Adiels *et al.*, 2006b) that is associated with insulin resistance, indicating that VLDL<sub>1</sub> and VLDL<sub>2</sub> are independently regulated (Gill *et al.*, 2004a; Packard & Shepherd, 1997).

### **1.5.2 Elevated Levels of VLDL<sub>1</sub>, Insulin Resistance and Obesity**

Hepatic overproduction of VLDL (particularly the large VLDL<sub>1</sub>) has been associated with insulin resistance (Gill *et al.*, 2004a; Johanson *et al.*, 2004) and related conditions such as diabetes (Adiels *et al.*, 2005b), the metabolic syndrome (Grundy, 2006) and obesity (Mittendorfer *et al.*, 2003a). Of particular interest, abdominal obesity which is characterised by excess visceral and/or deep upper body subcutaneous adipose tissue accumulation and is strongly associated with obesity, insulin resistance and atherogenic dyslipidaemia (Després, 2007; Sniderman *et al.*, 2007). Because visceral fat is more lipolytically active than subcutaneous fat *in vitro* (Mauriege *et al.*, 1987; Arner, 1998; Rebuffe-Scrive *et al.*, 1989), it has been hypothesised that visceral fat accumulation results in a markedly increased flux of NEFA via the portal vein to the liver (Bjorntorp, 1990; Chan *et al.*, 2004a), thereby increase substrate availability for hepatic VLDL formation (Kissebah *et al.*, 1976). However, Nielsen *et al.* (2004) demonstrated that although the release of NEFA from visceral fat depots into the portal vein increases with increasing visceral fat, it was much smaller than the amount derived from lipolysis of upper-body subcutaneous fat

(Nielsen *et al.*, 2004). Nevertheless, hepatic overproduction of VLDL<sub>1</sub> particles has been found to be driven by increased liver fat content in man (Adiels *et al.*, 2006b) and reduction in visceral adipose tissue is associated with decreased hepatic production of VLDL (Riches *et al.*, 1999). In addition, increased NEFA flux to the liver impairs hepatic extraction of insulin (Chan *et al.*, 2004a). Normally, insulin inhibits hepatic VLDL<sub>1</sub> production, in the fed state, but this effect is lost in insulin resistant conditions and high liver fat content, leading to increased hepatic VLDL<sub>1</sub> secretion (Adiels *et al.*, 2007).

Furthermore, the clearance of VLDL<sub>1</sub> and TRL particles and their remnants is impaired in insulin resistant conditions due to decreased LPL activity (Mead *et al.*, 2002), excess of apoC-III (an inhibitor of LPL) (Florez *et al.*, 2006) and failure to bind to LDLR and LRP (Packard, 2003) or suppression of LDLR (Mamo *et al.*, 2001).

### ***1.5.3 VLDL<sub>1</sub> and the Generation of Atherogenic Dyslipidaemia***

Whether due to hepatic overproduction or defective clearance, elevated levels of VLDL<sub>1</sub>, which give rise to a plasma TG concentration  $> 1.5 \text{ mmol.l}^{-1}$ , are believed to start a sequence of events that result in the atherogenic lipoprotein phenotype (Packard, 2003; Taskinen, 2003). Increased proportion of large LDL is a marker of efficient lipolysis of TG, which, in effect, will lower fasting and postprandial levels of TRL (Griffin, 1997). However, the presence of VLDL<sub>1</sub> gives rise to LDL particles ( $\beta$ LDL) with altered apoB-100 conformation which impedes binding of these particles to LDLR. In addition, impaired clearance of VLDL<sub>1</sub> particles (see above) will result in their accumulation in the circulation. As a result, VLDL<sub>1</sub> and  $\beta$ LDL particles have a prolonged residence time in the circulation, thereby increasing their likelihood of remodelling by CETP (Packard, 2003), which is increased in activity in insulin resistant states (Guerin *et al.*, 2001). Under these conditions, CETP promotes the acquisition of TG by LDL and HDL in exchange for CE from HDL to VLDL<sub>1</sub> and LDL (Packard, 2003; Guerin *et al.*, 2001). These TG-enriched LDL and HDL particles become favourable substrates for HL resulting in a reduction in particle size. HL also has an enhanced activity in obesity and insulin-resistant states (Deeb *et al.*, 2003). This gives rise to (1) sdLDL with a prolonged residence time ( $\sim 5$  days) compared with the larger LDL derived from VLDL<sub>2</sub> and IDL, which

has a residence time of ~2 days (Packard, 2003) and (2) small unstable HDL particles that are cleared rapidly from the circulation, resulting in low HDL cholesterol and apoA-I (Rashid *et al.*, 2003) and loss of its protective potential against CVD (Wang & Briggs, 2004). sdLDL is the most readily oxidised subfraction in the lipoprotein class (Tribble *et al.*, 1992), increasing its atherogenic potential. **Figure 1.6** shows a schematic diagram of the mechanisms involved in the formation of atherogenic dyslipidaemia.



## 1.6 Effect of Exercise on TRL Metabolism

### 1.6.1 Exercise as a Therapy for Dyslipidaemia

Due to the well established role of TRLs in CVD in progression of atherosclerosis described above, interventions that have the potential to reduce fasting and/or postprandial TG concentrations have been investigated as valuable tools for lowering CVD risk. Exercise has long emerged as an affordable, non-pharmaceutical method of improving lipoprotein metabolism. Regular exercise training has been shown to reduce total TG [as well as VLDL-TG (Kraus *et al.*, 2002)] concentrations, increase HDL cholesterol in adult men and women (Kraus *et al.*, 2002; Kodama *et al.*, 2007) and increase LDL particle size (Kraus *et al.*, 2002; Altena *et al.*, 2006). In addition, exercise improves insulin sensitivity in men and women (Gill *et al.*, 2002b), sedentary overweight/obese subjects (Houmard *et al.*, 2004), hypertriglyceridaemic men (Zhang *et al.*, 2006) and subjects with the metabolic syndrome (Zhang *et al.*, 2007) and type 2 diabetes (Alam *et al.*, 2004). Certainly, sedentary lifestyle has been associated with increased risk of obesity, metabolic deterioration, type 2 diabetes, CVD, cancer and all-cause mortality (Slentz *et al.*, 2007). In fact, a recent review on exercise training in patients with atheromatous CVD has found it to be a ‘true therapy’, reducing mortality by 25-35%, reducing clinical manifestations and complications (rhythm problems, thrombosis) and improving physical capacity, reintegration and quality of life (Casillas *et al.*, 2007). The following section focuses mainly on the effect of exercise on TRL metabolism.

### 1.6.2 Impact of Endurance Training

Compared with their untrained peers, endurance-trained men have reduced levels of postprandial lipaemia (Cohen *et al.*, 1989; Ziogas *et al.*, 1997; Merrill *et al.*, 1989). In addition, a period of endurance training has been shown to decrease fasting and/or postprandial TG concentrations in normolipidaemic subjects (Weintraub *et al.*, 1989), overweight men and women with mild-to-moderate dyslipidaemia (Kraus *et al.*, 2002) and older men and women (Halverstadt *et al.*, 2007). However, evidence suggests that this TG-lowering effect of exercise is due to short-term metabolic responses to recent exercise which is lost after a period of detraining. For example, there was no difference between fasting or postprandial TG concentrations between endurance trained and untrained young adults (Herd *et al.*, 2000) or untrained

middle-aged men (Tsetsonis *et al.*, 1997) after at least 2 days of inactivity, or after an exercise training programme when post-training measurements were made 60 h after the last exercise session (Herd *et al.*, 1998).

### **1.6.3 Impact of a Single Exercise Session**

An increasing body of evidence shows that a bout of exercise (e.g. brisk walking, jogging, cycling etc) reduces fasting and/or postprandial TG concentrations in different population groups including young adult men (Tsetsonis & Hardman, 1996b; Zhang *et al.*, 1998; Herd *et al.*, 2001) and women (Tsetsonis & Hardman, 1996b; Gill *et al.*, 2002a), middle-aged men (Gill *et al.*, 2001a; Gill *et al.*, 2001b) and women (Tsetsonis *et al.*, 1997), postmenopausal women (Gill & Hardman, 2000) and individuals with the metabolic syndrome (Zhang *et al.*, 2006) and type 2 diabetes (Tobin *et al.*, 2008), although responses are more heterogeneous in patients with type 2 diabetes (Gill *et al.*, 2007). The percentage reductions in plasma TG concentrations following exercise are broadly similar (15-25%) across these groups, however, subjects with higher TG concentrations (e.g. centrally obese men) had a greater decrease in postprandial TG compared with subjects with lower TG (Gill *et al.*, 2004b). Furthermore, a single session of exercise has shown to acutely decrease VLDL (Gill *et al.*, 2006) and VLDL-TG concentrations (Borsheim *et al.*, 1999; Magkos *et al.*, 2006; Morio *et al.*, 2004).

A session of exercise conducted before (Katsanos & Moffatt, 2004; Tsetsonis & Hardman, 1996b; Tsetsonis & Hardman, 1996a; Gill & Hardman, 2000; Gill *et al.*, 2001a; Gill *et al.*, 2001b) or after (Katsanos & Moffatt, 2004; Hardman & Aldred, 1995) ingestion of a fat meal has been reported to attenuate postprandial lipaemia. However, the weight of evidence (Zhang *et al.*, 2004a; Zhang *et al.*, 1998; Ferguson *et al.*, 1998; Borsheim *et al.*, 1999) suggests that the maximal TG-lowering effect of exercise appears to occur after a delay of approximately 12-18 h, rather than during or immediately post-exercise (Malkova & Gill, 2006). For example, Zhang and colleagues (Zhang *et al.*, 1998) reported a 21% greater reduction in postprandial TG when exercise was performed 12 h compared to 1 h prior to a meal and 49% lower than that performed after a meal. However, this TG-lowering effect seems to be 'short-lived' as exercising 24 h prior to the meal does not attenuate postprandial lipaemia (Zhang *et al.*, 2004a).

#### 1.6.4 Energy Expenditure and Energy Deficit

The energy expended during exercise is a key determinant of the TG-lowering effect: the higher the energy expenditure, the bigger the reduction in postprandial lipaemia (Petitt & Cureton, 2003). For instance, doubling energy expenditure by either doubling exercise intensity for the same duration [60 vs. 30% of maximal oxygen uptake ( $\dot{V}O_{2\max}$ ) for 90 min] (Tsetsonis & Hardman, 1996a) or by doubling exercise duration at the same intensity (120 vs. 60 min at 50%  $\dot{V}O_{2\max}$ ) (Gill *et al.*, 2002a) essentially doubles the exercise-induced reduction to lipaemia. In addition, postprandial TG concentrations are similarly reduced by different exercise settings (e.g. 180 min of walking at 30%  $\dot{V}O_{2\max}$  vs. 90 min at 60%  $\dot{V}O_{2\max}$ ) when the same amount of energy is expended (Tsetsonis & Hardman, 1996b). Similarly, a 2-h cycling at 60%  $\dot{V}O_{2\max}$  significantly decreased VLDL-TG and -apoB concentrations (Magkos *et al.*, 2006), whereas 1-h exercise of the same intensity had no effect (Magkos *et al.*, 2007). Of note, recently, Gormsen *et al.* (2006) reported a significant positive correlation between VLDL-TG production and resting energy expenditure, suggesting that it should be taken into account when VLDL-TG production comparisons between groups are made (Gormsen *et al.*, 2006).

Furthermore, the exercise-induced improvement in insulin sensitivity appears to be related to duration and intensity of exercise (Houmard *et al.*, 2004) and thus directly related to energy expenditure (Magkos *et al.*, 2008). It was proposed that exercise-induced changes in a homeostasis model assessment-estimated insulin resistance ( $HOMA_{IR}$ ) are curvilinearly related to exercise energy expenditure with a threshold of ~3.77 MJ (900 kcal) for improvements in  $HOMA_{IR}$  to be manifested (Magkos *et al.*, 2008). However, an inverse association was observed between the exercise-reduced changes in baseline (i.e. resting)  $HOMA_{IR}$ , suggesting that less exercise may be required to improve insulin sensitivity in insulin-resistant subjects than those with good insulin sensitivity at baseline (Magkos *et al.*, 2008).

The mechanisms by which exercise energy expenditure attenuates postprandial lipaemia are currently not known. Interestingly, dietary-induced energy deficit of similar magnitude to that induced by a bout of exercise session results in a much smaller TG reduction (Gill & Hardman, 2000). This implies that either the TG-



lowering effect of exercise is not dependent on energy deficit and/or that exercise- and dietary-induced energy deficits are not metabolically equivalent. However, a recent study investigating the effects of exercise, with or without energy replacement, on fasting and postprandial TG metabolism, demonstrated that the exercise-induced TG-lowering effect was only evident with an accompanying energy deficit (Burton *et al.*, 2008). This suggests that dietary-induced and exercise-induced energy deficits elicit different effects on postprandial metabolism. This may be related to specific body tissues in which the energy deficits occur as exercise induces quantitatively larger muscle and hepatic substrate deficits than energy intake restriction (Burton *et al.*, 2008).

### ***1.6.5 Intermittent versus Continuous Exercise***

Current exercise-for-health guidelines recommend the accumulation of physical activity throughout the day (Haskell *et al.*, 2007). Thus, it is important to understand whether small sessions of exercise spread throughout the day would be as beneficial in lowering TG concentrations as a single prolonged session of exercise. A number of studies have been conducted to investigate the effects of intermittent and continuous exercise, such as 90-min session *vs.* three 30-min sessions (Gill *et al.*, 1998), 30-min *vs.* three 10-min exercise session (Murphy *et al.*, 2000; Altena *et al.*, 2004) and even 30-min continuous exercise *vs.* ten 3-min bouts performed throughout the day (Miyashita *et al.*, 2006). All these studies consistently demonstrated that intermittent patterns of physical activity are effective in lowering TG concentrations as long as sufficient energy is expended, independently of the duration of the individual sessions.

### ***1.6.6 Potential Mechanisms Responsible for the Exercise-Induced TG Reduction***

Concentrations of TRLs in the circulation reflect the balance between rate of appearance and rate of clearance of intestinally-derived chylomicrons and hepatically-derived VLDL. Thus, the reduction in plasma TG concentration following exercise could be due to disturbance in the balance of appearance and clearance rates of chylomicrons and/or VLDL.

#### **1.6.6.1 Effect of Exercise on Chylomicron Metabolism**

The fact that maximal reduction in postprandial lipaemia is observed after a delay of number of hours, seems unlikely that the exercise-induced TG reduction is due to a reduction in gastrointestinal blood flow leading to potentially slower gut absorption and secretion of chylomicrons (Malkova & Gill, 2006). This is supported by three pieces of evidence: (1) the peak time for chylomicron-TG concentrations is not delayed post-exercise (Gill *et al.*, 2001a; Gill *et al.*, 2006), (2) gastric emptying time, evident by peak time of ingested paracetamol concentration, is unaffected by prior exercise (Gill *et al.*, 2001a; Gill *et al.*, 2001b) and (3) chylomicron particle number is not affected by prior exercise (James *et al.*, 2007).

There is clear evidence that endurance-trained individuals have high clearance rates of chylomicron-like lipid emulsions compared with untrained peers (Cohen *et al.*, 1989; Podl *et al.*, 1994), which is likely to reflect the increased post-heparin plasma LPL activity (reflecting overall LPL activity from all body tissues) observed in them (Podl *et al.*, 1994; Kantor *et al.*, 1984). In addition, muscle LPL activity has been reported to increase over 200% in response to intense exercise sessions lasting for hours (Lithell *et al.*, 1984; Sady *et al.*, 1986).

On other hand, recent studies investigating the effects of a more moderate exercise on LPL activity are equivocal. While some studies reported a significant increase in plasma (Zhang *et al.*, 2002), muscle and adipose tissue LPL activity after a moderate session of exercise (Perreault *et al.*, 2004), others reported no significant difference in muscle (Herd *et al.*, 2001) or post-heparin plasma LPL activity (Gill *et al.*, 2003) or muscle LPL mass (Magkos *et al.*, 2006) post-exercise. Interestingly, however, in the absence of an exercise-induced effect on muscle LPL mass, a significant ~20% increase in plasma LPL concentrations has been observed (Magkos *et al.*, 2006). Similarly, a significant correlation between LPL activity and the exercise-induced changes in fasting and postprandial TG has been observed without an effect on post-heparin LPL activity post-exercise (Gill *et al.*, 2003).

#### **1.6.6.2 Effect of Exercise on VLDL Metabolism**

Studies investigating the effect of moderate exercise on postprandial lipaemia demonstrated that the lipoprotein class most affected by prior exercise is VLDL

(Malkova *et al.*, 2000), specifically VLDL<sub>1</sub>, rather than chylomicrons (Gill *et al.*, 2001a; Gill *et al.*, 2006). However, little is known about the mechanisms that regulate VLDL concentrations (i.e. production and clearance rates) in response to moderate exercise. Certainly, there is no information available regarding the effect of exercise on large VLDL<sub>1</sub>.

Malkova *et al.* (2000) examined the influence of a prolonged session of prior exercise (running at ~60%  $\dot{V}O_{2max}$ ) on postprandial extraction of TG across the leg (Malkova *et al.*, 2000). Although they found no significant increase in total, chylomicron- and VLDL-TG uptake across the leg, TG clearance (defined as uptake divided by arterial concentration) – a marker for the efficiency of TG removal - was greater following exercise. However, as absolute VLDL-TG uptake did not differ significantly between control and exercise conditions, it is unclear whether the lower VLDL-TG concentrations following exercise were due to increase efficiency of removal or a lower VLDL production rate (Malkova *et al.*, 2000).

Following exercise, there is an increase in circulating NEFA (Burton *et al.*, 2008), which, theoretically, can increase hepatic VLDL secretion. However, this is not the case, as there is no correlation between the change in NEFA concentration and VLDL-TG following exercise (Borsheim *et al.*, 1999). This may be explained by the concurrent increase in 3-hydroxybutyrate following exercise (Malkova *et al.*, 2000; Burton *et al.*, 2008), which is a marker of hepatic fatty acid oxidation (Williamson & Whitelaw, 1978), suggesting a shift of the hepatic fatty acid flux towards  $\beta$ -oxidation and ketone body production and away from re-esterification and VLDL synthesis (Malkova & Gill, 2006).

Due to the fact that chylomicrons are the preferred substrate for LPL and their presence prevents VLDL clearance (Fisher *et al.*, 1995; Bjorkegren *et al.*, 1997), it has been hypothesised that moderate exercise likely to reduce hepatic VLDL production, rather than increase its clearance (Malkova & Gill, 2006). However, recent evidence suggests that a session of moderate exercise (e.g. walking or cycling for more than 1 h at 60%  $\dot{V}O_{2max}$ ) increases VLDL-TG clearance rates (Magkos *et*

*al.*, 2006; Tsekouras *et al.*, 2007) in healthy, lean, young men. In addition, Magkos *et al* also reported a decreased production of VLDL-apoB (reflecting the number of particles being produced) post-exercise (Magkos *et al.*, 2006), which may contribute to the exercise-induced reduction in VLDL concentrations. It should be noted, however, that these findings may not necessarily be applicable to overweight or centrally obese individuals (at which exercise-for-health guidelines are targeted), as this condition is (1) associated with hepatic overproduction of VLDL<sub>1</sub> due to their likely increased liver fat content (Adiels *et al.*, 2006b) and (2) impaired ability to suppress hepatic VLDL production due to their likely insulin resistance (Adiels *et al.*, 2007).

### **1.6.7 Effect of ApoE Phenotype**

ApoE is a 299-amino acid glycoprotein which is an integral surface component of TRLs and some subclasses of HDL (Hatters *et al.*, 2006). It primarily functions as a ligand for receptor-mediated uptake of TRLs, but also modulates the activity of LPL, LCAT and CETP (Leon *et al.*, 2004). The apoE gene has three alleles ( $\epsilon$ 2,  $\epsilon$ 3, and  $\epsilon$ 4) that give rise to 6 different phenotypes (E2/2, E2/3, E2/4, E3/3, E3/4, and E4/4) (Davignon *et al.*, 1988), with the E3 isoform being the most common (Hagberg *et al.*, 2000). ApoE polymorphism has been shown to have a substantial influence on plasma lipids and lipoproteins, which could be explained by a number of mechanisms (Kolovou & Anagnostopoulou, 2007): (1) receptor-binding affinities of different apoE-containing lipoproteins, (2) dietary fat clearance, (3) differences in the clearance of LDL-apoB and (4) efficiency of intestinal cholesterol absorption.

In the general population, the E2 isoform is associated with elevated levels of TG and apoE compared with the E3 isoform (Hagberg *et al.*, 2000), which is caused by impaired clearance of remnant particles bearing the apo E2 isoform probably due to defective receptor recognition (Havel *et al.*, 1980). Homozygous E2/2 carriers can develop type III hyperlipoproteinaemia, which is characterised by the accumulation of chylomicron and VLDL remnants in fasting plasma (Kolovou & Anagnostopoulou, 2007). On the other hand, the apo E4 isoform is typically associated with low TG levels compared with the apo E3 individuals (Davignon *et al.*, 1988). This is consistent with the fact that apo E4 subjects clear chylomicron

remnants into the liver more rapidly than E3/3 subjects and twice as fast as E3/2 subjects (Weintraub *et al.*, 1987).

In contrast, studies investigating the effects of apoE polymorphism on ( $\dot{V}O_{2\max}$ ), lipids and lipoproteins in response to endurance exercise training are conflicting. While a cohort study reported an improvement in lipid and lipoprotein profile in subjects homozygous for the E 3/3, despite a small reduction in  $\dot{V}O_{2\max}$ , compared with E 2/3 and E 3/4 (Thompson *et al.*, 2004), another reported no significant influence of apoE polymorphism on  $\dot{V}O_{2\max}$  in response to exercise (Leon *et al.*, 2004). Conversely, in a study with smaller subject number, increases in  $\dot{V}O_{2\max}$  (and HDL) in response to endurance exercise training, were in the order, apo E4 > apo E3 > apo E2 (Hagberg *et al.*, 1999).

However, to the best of the author's knowledge, there is only one study investigating the effect of apoE polymorphism on lipoprotein responses to moderate exercise. In a study including 38 men and 43 women, Gill *et al* (Gill *et al.*, 2002b) reported no significant differences between subjects possessing the E3/2, E3/3 and E4/3 phenotypes on the magnitude of the exercise-induced reduction in fasting or postprandial TG.

## 1.7 Measurements of TRL Kinetics

Lipoprotein metabolism is a complex system of tightly regulated and coordinated dynamic processes. Measurements of plasma lipid and lipoproteins concentrations provide useful, yet limited information – ‘a snapshot of the various processes’ (Packard, 1995), which do not reveal the dynamics of the system. This is achieved *in vivo* by the use of kinetic studies, in which labelled precursors are used to help characterise the metabolic pathways of lipoprotein metabolism (Packard, 1995; Barrett *et al.*, 2006). The system can be studied either in steady state (when the concentration of the substance of interest is constant) or in acute perturbation (non steady state, where a stimulus is applied and the return to steady state is monitored) (Packard, 1995).

There are basically two study protocols for investigating the metabolism of apoB-containing lipoproteins (i.e. VLDL<sub>1</sub>, VLDL<sub>2</sub>, IDL and LDL) (Demant & Packard,

1997). Lipoproteins can be tracer-labelled exogenously by isolating them from plasma, labelling them with radioactive substance (e.g. radioiodine). They are then re-injected into the donor subject and followed over time as they disappear from the plasma compartment. Alternatively, a stable isotope amino acid (e.g. D<sub>3</sub>-leucine or D<sub>5</sub>-phenylalanine) is injected, either as a bolus or as a primed constant infusion, and incorporated into the newly synthesised apoB protein serving as an endogenous label (**Table 1.3** shows the characteristics of each tracer and labelling method). In either case, multiple plasma samples are collected during the first 12 h of tracer injection and daily thereafter in the fasted state for up to 10-14 days. ApoB-containing lipoprotein fractions are then separated at each timepoint using cumulative gradient ultracentrifugation in salt solutions (Lindgren *et al.*, 1972) and apoB is isolated by selective precipitation (Egusa *et al.*, 1983). In the case of radioactive tracer studies, the specific activity is calculated by measuring the radioactive tracer in a scintillation counter, whereas in the stable-isotope tracer studies, tracer enrichment is determined by gas chromatography mass spectrometry (GC-MS) (Demant *et al.*, 1996). Multicompartmental modelling using the Simulation, Analysis And Modeling software (SAAM II; SAAM Institute, Seattle, WA) is used to calculate rates of production, delipidation and catabolism of apoB-containing lipoproteins (Packard *et al.*, 1995).

Although such methods provide valuable information about lipoprotein metabolism, they are difficult, time-consuming, expensive and labour-intensive. Furthermore, the laboratory manipulations are complex and require the use of specialised equipment (see below). [Comprehensive reviews of kinetic methods can be found by (Barrett *et al.*, 2006; Packard, 1995; Chan & Watts, 2006; Chan *et al.*, 2004b).]

### ***1.7.1 Using Stable-Isotopes to Measure VLDL<sub>1</sub> and VLDL<sub>2</sub> Kinetics***

Recently, Adiels and colleagues (Adiels *et al.*, 2005a) developed a multicompartmental model which permits the kinetics of apoB and TG to be assessed simultaneously in VLDL<sub>1</sub> and VLDL<sub>2</sub> fractions. The method involves administering D<sub>3</sub>-leucine and D<sub>5</sub>-glycerol as bolus injections in the fasted state. Several blood samples are taken before and for 8 h after the tracer injection while subjects are still fasting. Adiels *et al.* (Adiels *et al.*, 2005a) reported a significant linear correlation

between TG and apoB production in VLDL<sub>1</sub> and VLDL<sub>2</sub>, suggesting a coupling of the two processes governing the metabolism of these subfractions.

**Table 1.3: Characteristics of tracers [modified from (Packard, 1995)].**

Advantages and disadvantages of endogenous versus exogenous tracers	
Exogenous	Endogenous
Labels specific lipoprotein apos or lipoprotein subfractions	Labels all lipoprotein apolipoproteins
Labelled according to mass distribution	Labels according to production rate
Measures synthesis indirectly by inference from steady state calculation	Measures synthesis directly by tracer incorporation
Tracer does not recycle	Tracer may recycle
Urine data useful for assessing catabolism	Excretion data of little value
Radioactive versus stable-isotope endogenous tracers	
Radioactive	Stable isotope
Radiation hazard	Safe
Limited applicability	Applicable to all, including women and children
Limited repeatability	Can be repeated many times
Easy to measure	Difficult to measure with sufficient precision
Inexpensive apparatus and tracers	Capital cost high

For purposes of this thesis, it is essential to appreciate the complexity of the stable-isotope method, thus, **Figure 1.7** shows a schematic summary of the laboratory techniques involved, which were developed and are still used in the Vascular Biochemistry Department (Glasgow Royal Infirmary, University of Glasgow). Briefly, VLDL<sub>1</sub> and VLDL<sub>2</sub> fractions are separated from plasma by cumulative gradient ultracentrifugation (Lindgren *et al.*, 1972). ApoB protein is precipitated using absolute isopropanol and delipidated using ethanol-ether (Egusa *et al.*, 1983). Samples are then prepared (derivatised and fragmentised) for analysis of leucine and D<sub>3</sub>-leucine enrichment in hydrolysed apoB protein and plasma amino acids using the GC-MS (Demant *et al.*, 1994). In order to determine glycerol enrichment in TG, the isopropanol and ethanol-ether supernatants are treated with zeolite to precipitate PL and obtain TG. Glycerol is then extracted by saponification using potassium hydroxide and ethanol (Witter & Whitner, 1972), then derivatised and fragmented for enrichment determination by the GC-MS (Beylot *et al.*, 1987).

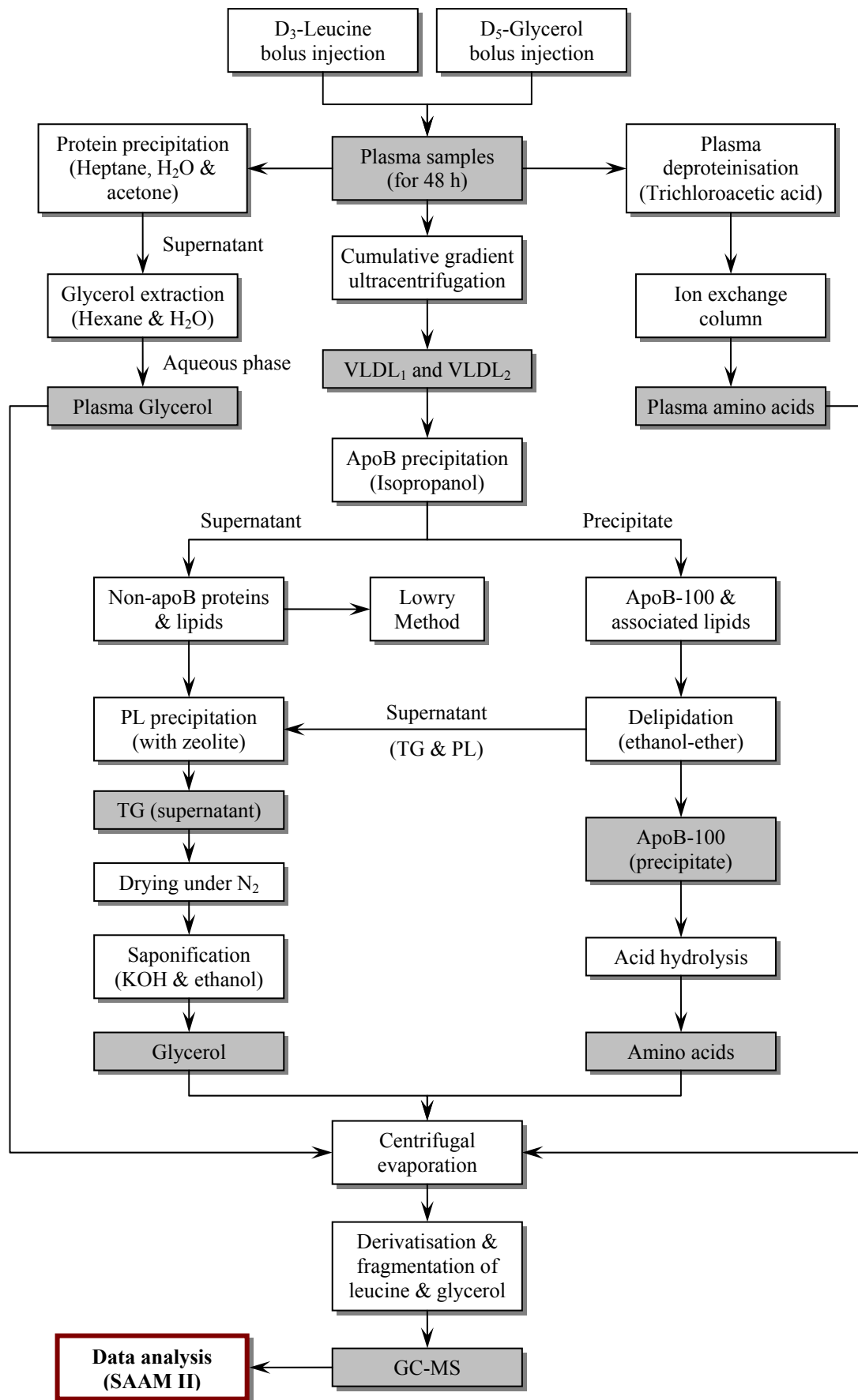
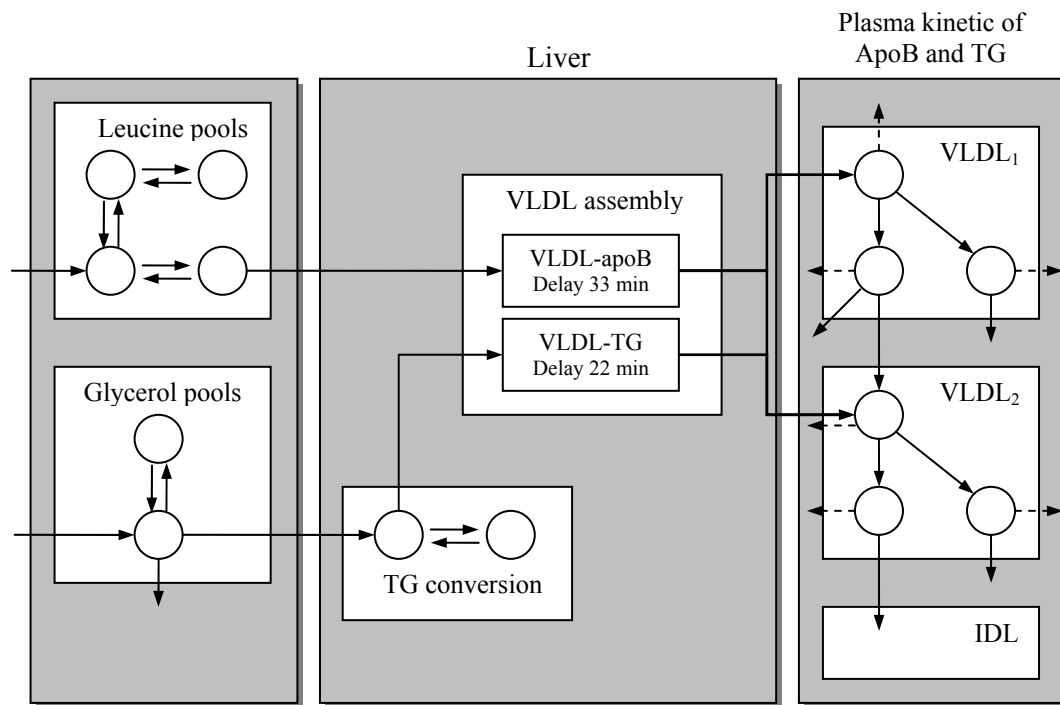


Figure 1.7: Schematic diagram of the stable-isotope laboratory protocol used to determine the kinetics of VLDL<sub>1</sub>- and VLDL<sub>2</sub>-TG and -apoB.



Based on the enrichment of leucine and glycerol in plasma, VLDL<sub>1</sub> and VLDL<sub>2</sub>, and the known injected amounts of labelled leucine and glycerol, the kinetic parameters are determined using the modelling software SAAM II as previously described (Adiels *et al.*, 2005a). **Figure 1.8** shows the mathematical multicompartmental model used to assess the kinetics of glycerol and leucine in plasma and lipoprotein fractions, which is based on the apoB model originally described by Packard *et al* (Packard *et al.*, 1995).



**Figure 1.8:** A diagram of the mathematical multicompartmental model (using SAAM II software) used to simultaneously determine the kinetics of apoB and TG in VLDL<sub>1</sub> and VLDL<sub>2</sub> after a bolus injection of stable isotopes D<sub>3</sub>-leucine and D<sub>5</sub>-glycerol [Figure from (Adiels *et al.*, 2005a)].

### 1.7.2 Chylomicron Kinetics

#### 1.7.2.1 ApoB-48 Labelling with Stable-Isotopes

Kinetics of apoB-48 containing lipoproteins are difficult with only a few studies reported using endogenously labelled stable-isotope tracer (primed constant infusion of [5,5,5-<sup>2</sup>H<sub>3</sub>] leucine) (Welty *et al.*, 1999; Welty *et al.*, 2004; Lichtenstein *et al.*, 1992; Hogue *et al.*, 2007). This is because in the fasting state, apoB-48 concentrations are generally too low for protein enrichment to be detected, whereas they vary postprandially (along with other lipoproteins) which makes interpretation

of tracer data very difficult (Barrett, 1998). However, these obstacles were first overcome by Lichtenstein *et al.* (1992) by providing small hourly feeds of identical composition to the subjects (over a 20 h period) to ensure an 'elevated' steady state of chylomicron concentrations. Recently, Bickerton *et al.* used a combination of stable isotope labeling to specifically label VLDL (using an intravenous infusion of [ $^2\text{H}_2$ ]palmitate) and chylomicrons (using a test meal containing [ $\text{U-}^{13}\text{C}$ ]palmitate) to quantify the TG extraction across human skeletal muscle and adipose tissue (Bickerton *et al.*, 2007). However, it was reported that rapid recirculation of the label leads to loss of chylomicron specificity (Heath *et al.*, 2003; Karpe *et al.*, 2007).

#### **1.7.2.2 Retinol Palmitate**

Another commonly used, non-stable isotope, method to measure chylomicron metabolism involves the ingestion of retinol with a fat load, which is esterified into retinyl palmitate and secreted into the intestinal lymph in the core of the chylomicron particle. The palmitate within the core remains with the particle as it undergoes lipolysis and becomes a remnant particle. Thus, production and clearance rates of chylomicrons can be determined by monitoring the concentrations of plasma palmitate or chylomicron-associated palmitate (Barrett, 1998). However, a major disadvantage of this approach is the exchange of core components between chylomicrons and other lipoproteins [e.g. 25% of retinyl palmitate may also be found in apoB-100 containing TRLs (Cohn *et al.*, 1993)].

#### **1.7.2.3 Chylomicron-Like (Lipid) Emulsions**

Intravenous lipid emulsions have been developed to supply patients with a balanced parenteral nutrition and to prevent or correct essential fatty acid deficiency. They are TG-rich particles that are modeled on the endogenous chylomicrons and consist of a TG core, traditionally derived from vegetable oils (e.g. soybean, safflower, coconut, olive). This TG is stabilised by a monolayer of PL derived from egg yolk (Olivecrona & Olivecrona, 1998; Carpentier & Dupont, 2000). However, unlike chylomicrons, lipid emulsions contain no cholesterol or apolipoproteins (Olivecrona & Olivecrona, 1998). Nevertheless, once in the circulation, these TG-rich particles rapidly acquire apoCs and apoE from plasma lipoproteins, mainly HDL (Iriyama & Carpentier, 1994); apoC-II is necessary for the action of LPL and apoE is important for receptor recognition.

Because the intravascular metabolism of these TG-rich particles resembles that of chylomicrons (Olivecrona & Olivecrona, 1998), they have been also used to study particular aspects of chylomicrons metabolism, both *in vitro* and *in vivo*. After binding of emulsion particles to LPL situated on the endothelial cells of adipose and muscle tissues, a substantial amount of the TG content of these particles is hydrolysed and the released NEFAs are either taken by the adjacent tissue or spilled into the circulation (Evans *et al.*, 1999; Carpentier & Dupont, 2000). However, evidence suggests that lipolysis and particle removal may be simultaneous rather than sequential mechanisms like chylomicrons; i.e. emulsion particles may disappear from blood before they have become TG-depleted remnants. This is because some remnants are not likely to leave the site of lipolysis (endothelial site) but be internalized together with HSPG/LPL or be delivered to adjacent specific lipoprotein receptors (Olivecrona & Olivecrona, 1998), including VLDLR (Carpentier & Dupont, 2000).

Intralipid® (soybean oil) and Liposyn® (safflower oil) are two widely used commercial lipid emulsions, with the major difference in composition is their  $\alpha$ -linolenic acid content. While Intralipid contains 8%, Liposyn has only 0.5% (Byrne, 1982). Intralipid emulsions have been used in intravenous fat tolerance tests to study TG clearance rate (see below). Recently, Park and colleagues (Park *et al.*, 2000; Park *et al.*, 2001) have reported a new method to measure chylomicrons kinetics by labelling Liposyn with radioactive tracers.

Redgrave and Maranhao (1985) prepared experimental lipid emulsions which, unlike the commercial ones, contain triolein, cholesteryl oleate, cholesterol and egg phosphatidyl-choline (Redgrave & Maranhao, 1985). The particles are usually smaller than those in commercial emulsions and they contain no PL vesicles (Olivecrona & Olivecrona, 1998). These emulsions have also been used in chylomicron kinetic studies, where the emulsion was double-labelled with radioactive cholesteryl esters and TG. After intravenous injection into the subjects, determination of the plasma decaying curves of the labeled lipids allows the 2-step metabolism of chylomicrons to be followed (Oliveira & Maranhao, 2002).

#### 1.7.2.4 Intravenous Fat Tolerance Test

The intravenous fat tolerance test (IVFTT) provides a simple measure of postprandial TG clearance. It was first introduced in the early 1960s by Carlson and Hallberg (Carlson & Hallberg, 1963) in dogs then in humans in 1965 (Hallberg, 1965). In 1972, Carlson and Rössner (Carlson & Rossner, 1972) provided an IVFTT protocol which involved the injection of a bolus dose of Intralipid (0.1 g TG per body mass) into a forearm vein, with blood being drawn from a contralateral antecubita vein at 5-min intervals for 40 min. The decline in plasma Intralipid-TG concentrations followed first-order kinetics (i.e. the clearance rate was directly proportional to the concentration). Initially, Carlson and Rössner measured Intralipid-TG concentrations indirectly in plasma by measuring plasma turbidity using nephelometry (Carlson & Rossner, 1972), and although Rössner later reported a good reproducibility of this method for up to 6 months (Rossner, 1982), Sady and colleagues found better reproducibility using enzymatic TG quantification (Sady *et al.*, 1986).

### 1.8 Summary and Objectives

This chapter attempted to briefly review current knowledge of lipid and lipoprotein metabolism, particularly TRLs. Special emphasis was placed on VLDL<sub>1</sub> assembly, metabolism, regulation and role in development of atherogenic dyslipidaemia. In addition, the role of exercise as a potential therapeutic option to reduce VLDL concentrations, in an attempt to improve the atherogenic lipoprotein phenotype, was discussed. Finally, the last section focused on the necessity of determining TRL kinetics; i.e. production and clearance rates, rather than concentrations, and the methods used to do so. Of note, despite the well-established heterogeneity of VLDL and the independent regulation of VLDL<sub>1</sub> and VLDL<sub>2</sub>, the majority of kinetic studies, especially those aimed at investigating TG kinetics, have focused on total VLDL, rather than VLDL<sub>1</sub> and VLDL<sub>2</sub> separately. This is likely due to the difficulty in determining TG kinetics in VLDL subfractions compared with total VLDL. Although this obstacle has been recently overcome, the method still employed the use of stable-isotope tracers. Although these methods are considered the ‘gold-standard’ in kinetic studies and have been providing valuable information for a better understanding of lipoprotein metabolism, they are not widely available to most

laboratories due to their complexity and high cost. Therefore, the aims of the present thesis are to develop and validate a relatively easy and cost-effective method of determining VLDL<sub>1</sub> kinetics and use it to investigate the effects of hyperglycaemia and hyperinsulinaemia as well as moderate exercise on VLDL<sub>1</sub> kinetics. This is important as elevated concentrations of VLDL<sub>1</sub>, rather than VLDL<sub>2</sub>, are believed to be responsible for the generation of atherogenic dyslipidaemia. In addition, it is essential to elucidate the mechanisms by which moderate exercise, a potential tool for lowering CVD risk, reduces plasma TG in obese/overweight middle-aged men; a typical population at which exercise-for-health guidelines are targeted.

## 2. General Methods

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This chapter is divided into three main parts. The first part describes the employed experimental procedures, many of which are common to several studies. The second part explains a study conducted to compare between a chemical manual reaction and an antibody-based automated method for apoB determination in VLDL fractions. Finally, the third part describes a pilot study conducted to determine the concentration and dose of a glucose drink needed to sustain a steady hyperinsulinaemic and hyperglycaemic state.

### 2.1 Subject Recruitment

Apparently healthy subjects were recruited from within the University of Glasgow, Glasgow Royal Infirmary and Greater Glasgow area by personal contact as well as by local advertising using posters, web-based and radio announcements. At least one week prior to the study, volunteers were interviewed and a subject information sheet about the study, including possible risks and discomforts (Appendices A1 & A2), was given and explained to them. Volunteers were also encouraged to ask any questions before signing an informed consent (Appendices A1 & A2) and screened according to a health screen questionnaire (Appendices B1 & B2). Their resting arterial blood pressure was measured and a venous blood sample was collected to test for their liver, renal, and thyroid functions. For the glucose and exercise studies described in [Chapters 4](#) and [5](#), subjects were also tested for fasting glucose. Common exclusion criteria were used as follows:

- a. A history of known cardiovascular disease or abnormalities, including established CHD (e.g. MI, stroke, CABG), acute illness or active, chronic systemic disease
- b. Uncontrolled hypertension ( $>160/90$  mmHg on anti-hypertensive medication)
- c. Taking any medication known to influence carbohydrate or lipid metabolism;
- d. Anaemia ( $\text{Hb} < 12 \text{ g.dl}^{-1}$  males,  $< 11 \text{ g.dl}^{-1}$  females)
- e. Current smoking (stopped for at least 6 months)
- f. Frank diabetes (fasting blood sugar  $\geq 7 \text{ mmol.l}^{-1}$ ) ([Chapters 4](#) and [5](#))

- g. Abnormal renal, liver or thyroid function tests
- h. Allergy to soybeans
- i. Taking part in another study (within the last 3 months)

Inclusion criteria were different for different studies. These will be discussed in [Chapters 4](#) and [5](#). All study protocols were approved by the Research Ethics Committee of the North Glasgow University Hospitals NHS Trust.

## **2.2 Anthropometry**

### **2.2.1 Height and Weight**

The heights and weights of subjects were measured using a weighing machine (D. Brash and Sons; Glasgow, UK) with an attached graduated metal plate. Subjects stood barefoot with their heels together and wearing light clothing. Height (m) and weight (kg) measurements were made to the nearest 0.1 unit.

### **2.2.2 Waist and Hip Circumferences**

Waist and hip circumferences were measured using a flexible, inelastic tape measure (Supralip®160, West-Germany). The waist circumference was measured on the horizontal plane midway between the costal margin and iliac crest with the abdominal muscles relaxed and the subject breathing shallowly. The hip measurement was made horizontally around the maximum circumference of the buttocks. The reported values are means of two to three measurements.

## **2.3 Blood Sampling**

Venous blood was obtained via an indwelling cannula (Biovalve, 18G/1.2 mm, Vygon, France) placed in an antecubital vein, to which a 10 cm three-way stopcock (Connecta Plus 3, BD, Sweden) was attached. The cannula was kept patent by flushing with non-heparinised saline solution (0.9% NaCl). Blood samples were collected directly in 10-ml tubes containing K<sub>3</sub>EDTA as an anticoagulant (BD Vacutainer Systems, Plymouth, UK). Blood samples were then placed immediately in ice and centrifuged (GS-6KR, Beckman Instruments, Inc, California, US) within 15-30 min of collection for 15 minutes at 3000 rpm and 4° C.

EDTA plasma was pipetted into aliquots of 200  $\mu\text{l}$  in 0.5 ml Eppendorf tubes (Treff Lab, Switzerland) and  $2 \times 350 \mu\text{l}$  in Apex tubes (2 ml, Alpha Laboratory Ltd, UK) and frozen immediately at  $-70^{\circ}\text{C}$ , for subsequent analysis of insulin, NEFA, glucose and lipid profile. The remaining EDTA plasma was used for lipoprotein separation. This was either started on the same day of blood collection or the next morning. In case of the latter, plasma was stored overnight at  $4^{\circ}\text{C}$ .

At the end of a successful lipoprotein separation, the remaining EDTA plasma was pipetted into aliquots of no less than 0.5 ml (to minimise any freeze drying effect) in 1.5 ml Eppendorf tubes (Treff Lab, Switzerland) and stored at  $-70^{\circ}\text{C}$ .

## 2.4 Blood Samples Analysis

Lipoprotein fractions and EDTA plasma were separated and analysed in the Vascular Biochemistry Department of Glasgow Royal Infirmary, University of Glasgow. The methods and techniques employed were taught to the author by researchers in the Vascular Biochemistry Department.

### 2.4.1 Separation of $\text{VLDL}_1$ and $\text{VLDL}_2$ Lipoprotein Fractions

$\text{VLDL}_1$  ( $S_f$  60-400) and  $\text{VLDL}_2$  ( $S_f$  20-60) fractions were isolated from plasma using a modification of the cumulative ultracentrifugation density gradient technique described by Lindgren *et al.* (Lindgren *et al.*, 1972).

The density of 2 ml of plasma was adjusted to  $d\ 1.118\ \text{g.ml}^{-1}$  by the addition of 0.341 g NaCl. This was carefully layered over a cushion of 0.5 ml  $d\ 1.182\ \text{g.ml}^{-1}$  solution in an ultraclear Beckman SW 40 ultracentrifugation tube (Beckman Instruments Inc., UK) which had been coated with polyvinyl alcohol (Holmquist, 1982); this allowed the solutions to gravity feed down the side of the tubes smoothly without disturbing the formation of the gradient. A discontinuous gradient was formed by overlaying  $d\ 1.0988\ \text{g.ml}^{-1}$  (1 ml),  $d\ 1.0860\ \text{g.ml}^{-1}$  (1 ml),  $d\ 1.0790\ \text{g.ml}^{-1}$  (2 ml),  $d\ 1.0722\ \text{g.ml}^{-1}$  (2 ml),  $d\ 1.0641\ \text{g.ml}^{-1}$  (2 ml) and finally  $d\ 1.0588\ \text{g.ml}^{-1}$  (2 ml). The density solutions were prepared from stock solutions  $1.006\ \text{g.ml}^{-1}$  and  $d\ 1.182\ \text{g.ml}^{-1}$  of NaBr in 0.195M NaCl, 0.001%  $\text{Na}_2\text{EDTA}$  and their densities were measured to 3 decimal places in a Paar Scientific densitometer (model DMA 35). The centrifugation was carried out using a Beckman SW 40 rotor (Beckman Instruments Inc., UK) in a



Beckman L8-60M ultracentrifuge for 1.38 h at 39K rpm and 23° C for separation of the VLDL<sub>1</sub> fraction. Rotors were decelerated without brakes and VLDL<sub>1</sub> was removed in the top 1 ml using a finely drawn glass Pasteur pipette. This volume was replaced by 1 ml of d 1.0588 g.ml<sup>-1</sup> and tubes were recapped and placed back in the centrifuge overnight for separation of VLDL<sub>2</sub>. Various run times and speeds required for VLDL<sub>2</sub> separation were previously calculated as shown in **Table 2.1**. At the end of the run, 0.5 ml of VLDL<sub>2</sub> fraction was aspirated in the same way as VLDL<sub>1</sub>. Fractions were kept in tightly capped 2 ml Apex tubes (Alpha Lab Ltd., UK) at 4° C for subsequent analysis.

**Table 2.1: Various times and speeds required for the separation of lipoprotein fractions using Beckman L8-60M Ultracentrifuge and SW 40 rotor.**

VLDL <sub>1</sub> ( $\omega^2t$ 9.81)		VLDL <sub>2</sub> ( $\omega^2t$ 2.12)	
Time (h)	RPM	Time (h)	RPM
1.38	39K	12.03	21.1K
		14.52	19K
		15.41	18.5K
		16.34	18K
		17.31	17.5K
		18.08	17.2K
		20.58	16K

#### 2.4.2 Spectrophotometric Assays

Plasma analyses were carried out using commercially available enzymatic colorimetric kits. Plasma glucose (Glucose hexokinase, Randox Laboratories Ltd, UK) and total and HDL cholesterol (CHOL and HDL-C, Roche Diagnostics, UK, respectively) were analysed in the fasted state. LDL cholesterol was calculated in the fasted state using the Friedewald equation [LDL cholesterol (mmol.l<sup>-1</sup>) = total cholesterol (mmol.l<sup>-1</sup>) – HDL cholesterol (mmol.l<sup>-1</sup>) – (TG (mmol.l<sup>-1</sup>)/2.2) (Friedewald *et al.*, 1972)]. TG (TG, Roche Diagnostics Limited, UK), NEFA (NEFA C, Wako Chemicals, USA) and glycerol (GLY, Randox Laboratories Ltd, UK) were analysed at all time points.

In [Chapter 5](#), PL (Phospholipids C, Wako Chemicals, USA) and FC (Free Cholesterol, Wako Chemicals USA) were measured in the VLDL<sub>1</sub> fraction in all time points. In addition, VLDL<sub>1</sub>-apoC-II, -apoC-III, and -apoE were also measured in the fasted state using commercial automated turbidimetric immunoassay kits, supplied by Wako Chemicals.

All the above analyses were made by an automated clinical chemistry analyser (ILab<sup>TM</sup> 600, Instrumentation Laboratory, USA) by Mrs. Josephine Cooney, Mrs. Elaine McDonald and Mrs. Elizabeth Murray (Department of Vascular Biochemistry, Glasgow Royal Infirmary).

Serum ALT concentrations were measured at screening (General Biochemistry Laboratory, Glasgow Royal Infirmary) using commercially available enzymatic colorimetric kit (Alanine Aminotransferase, Abbott Laboratories, USA).

### **2.4.3 Insulin ELISA**

Insulin was measured in freshly frozen EDTA plasma using commercially available ELISA kits (Mercodia AB, Uppsala, Sweden). The method is a solid phase two-site enzyme immunoassay. The CV for the assay was < 4%.

### **2.4.4 Apolipoprotein E Phenotyping**

ApoE phenotype was determined for each subject by Mrs Elizabeth Murray (Department of Vascular Biochemistry, Glasgow Royal Infirmary), by isoelectric focusing using Western blot techniques as described by Menzel (Menzel & Utermann, 1986) and Havekes (Havekes *et al.*, 1987).

### **2.4.5 Total Protein Determination (Lowry Method)**

Total protein content in lipoprotein fractions was measured using a modified Lowry method (Lowry *et al.*, 1951). This method involved the addition of 1 ml of Biuret reagent [100 ml of 2% Na<sub>2</sub>CO<sub>3</sub> in 0.1 M NaOH (w/v), 1 ml of 2% NaK Tartrate (w/v), 1 ml of 1% CuSO<sub>4</sub> (w/v), and 1 ml of 10% (w/v) sodium dodecyl sulphate (SDS)] to a total volume of 200 µl of sample (100 µl VLDL<sub>1</sub> + 100 µl of distilled water or 50 µl VLDL<sub>2</sub> + 150 µl distilled water). Samples were diluted or concentrated as

required. One hundred microlitre of 1:1 Folin Ciocalteu reagent (Sigma-Aldrich Company Ltd., Irvine, Scotland) was then added and mixed immediately. After incubation at room temperature for 30-60 min, the developed colour intensity was measured at an optical density of 750 nm using a Beckman DU 70 Spectrophotometer. The protein concentration was calculated using a standard curve of known concentrations. Inter-assay precision was checked using 2 levels (100 µl and 200 µl) of human and bovine quality control (QC) materials. The coefficients of variation (CVs) for the low QC were (human: 1.8%, bovine: 1.3%) and high QC were (human: 1.0%, bovine: 1.4%).

#### **2.4.6 Accuracy and Precision of Assays**

The accuracy and precision of the automated assays described [section 2.4.2](#), except for glycerol, were monitored using quality control sera (Wako QC abnormal and normal, Wako Chemicals GmbH, Denmark; distributed by Alpha Laboratories, UK). Glycerol quality control was obtained from Randox Laboratories Ltd, UK. In order to minimise intra-assay variation, all samples obtained for each subject were performed in the same analyser run for a given assay. The CVs for the assays were 2.9% for total cholesterol, 3.8% for TG, 2.8% for HDL-cholesterol, 5.2% for NEFA, 2.0% for glucose, 1.9% for glycerol, 2.2% for free cholesterol, and 3.8% for PL.

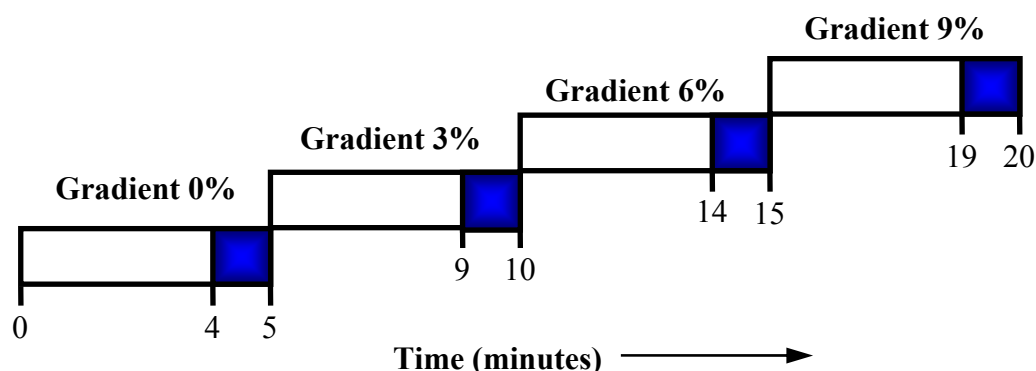
### **2.5 Bruce Protocol**

All subjects recruited for the exercise study described in [Chapter 5](#) underwent a modified (ACSM, 2006) Bruce protocol (Bruce *et al.*, 1973) under the supervision of a medical doctor: Dr Nicholas Barwell or Dr Lesley Hall, at the Institute of Diet, Exercise and Lifestyle (IDEAL) at the University of Glasgow. A modified twelve lead electrocardiographic (ECG) monitoring was conducted throughout the test and arterial blood pressure was measured at rest and immediately after a Bruce protocol during recovery.

### **2.6 Submaximal Exercise Test**

In the exercise study described in [Chapter 5](#), subjects undertook a submaximal exercise test to estimate the gradient necessary to elicit an intensity corresponding to 50%  $\dot{V}O_{2max}$ . The test consisted of a four-stage treadmill (Woodway GmbH, Weil am

Rhein, Germany) walk (**Figure 2.1**) to determine the relationship between increasing gradient and the subjects'  $\dot{V}O_2$  at their self-selected walking speed.  $\dot{V}O_{2\max}$  was estimated by extrapolation of the  $O_2$  uptake/heart rate relationship up to the subject's predicted maximum heart rate:  $[220 - \text{age}] \text{ beat} \cdot \text{min}^{-1}$  (ACSM, 2006). Each stage was five minutes long with expired air samples being taken during the last minute of each stage. Heart rates were recorded during the expired air collections as were perceived rates of exertion using the Borg scale (Borg, 1973). The first stage of the test was on a level treadmill with gradient increasing by 2.0 to 3.0% in each subsequent stage. The increase in gradient for each stage was established on an individual basis based on the subject's heart rate response. The relationship between  $\dot{V}O_2$  and treadmill gradient was determined for each subject and, together with the estimated  $\dot{V}O_{2\max}$ , the gradient necessary to elicit an intensity corresponding to 50%  $\dot{V}O_{2\max}$  was calculated.



**Figure 2.1:** A schematic diagram of a 4-stage submaximal exercise test performed for subjects in Chapter 5. The blue block represents an expired air sample and heart rate measurements.

## 2.7 Monitoring of Heart Rate

The subject's heart rate ([Chapter 5](#)) was monitored continuously during exercise and the recovery period by short range telemetry (Polar S610i, Polar Electro, Finland).

## 2.8 Measurement of Oxygen Uptake and Carbon Dioxide Production

Oxygen uptake and  $CO_2$  production in [Chapter 5](#) were determined at rest before and during exercise. Samples of expired air were collected into 100 or 150 L Douglas

bags. While wearing a nose clip, subjects breathed through a mouthpiece fitted to a lightweight large 2-way respiratory valve (2700 series, Hans Rudolph Inc. USA), which in turn was connected to a lightweight tube. The tubing was terminated at a two-way valve which opened and closed the Douglas bag. All equipment was supplied by Cranlea & Co. Birmingham, England.

An aliquot of expired air (measured using a flow meter) was removed from each Douglas bag to determine the fraction of O<sub>2</sub> and CO<sub>2</sub> using a gas analyser (Servomex 4100, Servomex Group Ltd., East Sussex, England). The analyser was calibrated before each use with certified reference gases (BOC Ltd, Surry, UK) and the reference gases were calibrated against a 'gold standard' reference gas to ensure consistency of results.

The remaining volume of expired air in each Douglas bag was measured by evacuation through a dry gas meter (Harvard apparatus, supplied by Cranlea & Co. Birmingham, England). The temperature of air in the Douglas bag was measured during evacuation using the same dry gas meter.

Barometric pressure was measured using a barometer and the measured expired gas volumes were corrected to standard temperature and pressure (STPD) for a dry gas using the universal gas equation. Inspired gas volumes were derived using the Haldane transformation (Consolazio *et al.*, 1963) and O<sub>2</sub> uptake, CO<sub>2</sub> production, minute ventilation, respiratory exchange ratio and the ventilatory equivalent for oxygen were calculated. Rates of substrate utilization expenditures were calculated via indirect calorimetry using the equations described by Frayn (Frayn, 1983) and energy expenditure was determined by multiplying the mass of substrates used by their respective energy densities.

## **2.9 Apolipoprotein B Measurements in VLDL<sub>1</sub> and VLDL<sub>2</sub> Fractions – Comparison of a Manual and an Automated Method**

### **2.9.1 Introduction**

In the Vascular Biochemistry laboratory, for kinetic studies, apoB is usually measured in lipoprotein fractions using a standardised manual method, which is applicable to VLDL, IDL and LDL fractions. ApoB is precipitated after the addition of an equal volume of absolute isopropanol (Egusa *et al.*, 1983). The supernatant is aspirated and its protein content is measured using a modified version of Lowry method (Lowry *et al.*, 1951). There is no apoB solubilisation in isopropanol and the precipitated apoB is virtually free of soluble apolipoproteins. ApoB protein concentrations are measured by the difference between total protein content in lipoprotein and the protein measured in supernatant. This method is standardised and widely used for the measurement of the relatively low concentrations of apoB in VLDL<sub>1</sub> and VLDL<sub>2</sub> fractions. However, it is laborious, time consuming and, being a manual technique, is subject to personal error.

Another method for measuring apoB concentrations is direct measurement using an automated *in vitro* immunoturbidimetric assay with the use of commercially available kits (see below). Being an automated method, the immunoturbidimetric assay saves time, effort and potentially improves reproducibility. However, these have been designed for apoB measurements in serum or plasma, where apoB concentrations are high and mainly represent apoB in the LDL fraction [85-90% of total apoB (Alaupovic, 1991)]. It was therefore unclear whether this method is suitable for the measurement of the low apoB concentrations in VLDL<sub>1</sub> and VLDL<sub>2</sub> fractions.

Therefore, the purpose of the present study was to compare the automated turbidimetry immunoassay method with the conventional precipitation one as an alternative method for the measurement of apoB protein in VLDL<sub>1</sub> and VLDL<sub>2</sub>.

## **2.9.2 Materials and Methods**

### **2.9.2.1 Specimens and Separation of VLDL<sub>1</sub> and VLDL<sub>2</sub> fractions**

A total of 260 EDTA plasma samples were used to separate VLDL<sub>1</sub> (S<sub>f</sub> 60-400) and VLDL<sub>2</sub> (S<sub>f</sub> 20-60) fractions by a modification of the cumulative ultracentrifugation density gradient technique as previously described ([section 2.4.1](#))

### **2.9.2.2 Isopropanol Precipitation Method (Egusa *et al.*, 1983)**

All VLDL<sub>1</sub> and VLDL<sub>2</sub> samples were brought to room temperature and 0.5 ml of absolute isopropanol was added to 0.5 ml of each sample in glass tubes (13/100 glass tubes, Labco Ltd., Buckinghamshire, UK). After immediate vigorous mixing, samples were then incubated for at least 24 h at 4° C, after which they were spun at 3000 rpm and 4° C for at least 30 min. Immediately, the supernatant was carefully aspirated using a drawn-out Pasteur pipette, without drawing out any of the precipitated apoB.

Total protein content in the VLDL<sub>1</sub> and VLDL<sub>2</sub> fractions and the non-apoB protein content of the supernatant were measured using the modified Lowry method as described earlier ([section 2.4.5](#)).

### **2.9.2.3 Immunoturbidimetric Method**

The apoB content of the same samples was measured directly using commercially available kits (WAKO Apolipoprotein B-HA, Wako Chemicals GmbH, Denmark; distributed by Alpha Laboratories). This method involved the specific combination of apoB in the sample with anti-human apoB antibodies in the reagent to yield an insoluble aggregate that causes increased turbidity. The degree of turbidity is measured optically using an autoanalyser (ILab<sup>TM</sup> 600, Clinical Chemistry System, Instrumentation Laboratory, USA). The method in Vascular Biochemistry has been optimised for apoB measurements in lipoprotein fractions by increasing the volume of sample used in the assay from 3 µl to 12 µl (results were then multiplied by a factor of 0.25). The supplied quality control was also diluted to account for the lower concentrations of apoB (WAKO 3 was diluted 1:5 to give a mean (range) concentration of 15.3 (12.3-18.4) mg.dl<sup>-1</sup>, which is within the range of expected results).

#### 2.9.2.4 Calculating ApoB %CV for Each Method

VLDL<sub>1</sub> and VLDL<sub>2</sub> were separated from two subjects and each fraction was pooled together. ApoB was measured in ten samples using the immunoturbidimetric method and the isopropanol-Lowry method. Means, SDs and CVs were calculated for each method.

#### 2.9.3 Results

**Figure 2.2** shows VLDL<sub>1</sub>- and VLDL<sub>2</sub>-apoB concentrations (mg.dl<sup>-1</sup>) measured by the immunoturbidimetry method in 260 samples plotted against that measured by the isopropanol precipitation method with the line of equality; the line all points would lie on if the two methods always gave exactly the same measurement. The two methods correlate well for both fractions with R<sup>2</sup> (goodness-of-fit) of 0.88 for VLDL<sub>1</sub> and 0.93 for VLDL<sub>2</sub>. However, the isopropanol method measured values as ~1.3 times higher than the immunoturbidimetric method for both VLDL<sub>1</sub> and VLDL<sub>2</sub> fractions. This was particularly apparent as the apoB concentrations increased.

**Figure 2.3** shows the percent difference between measurements by the two methods (isopropanol minus immunoturbidimetry) for each sample against their mean in both the VLDL<sub>1</sub> and VLDL<sub>2</sub> fractions. This is done to explicitly show extreme or outlying observations and any lack of agreement between the two methods (Bland & Altman, 1999). In addition, **Figure 2.3** shows the 95% limits of agreement estimated as mean percentage difference  $\pm$  2 SD for each fraction, which define the range within which most differences between measurements by the two methods will lie (Bland & Altman, 1999). This was:  $0.8 \pm 110.3\%$  for VLDL<sub>1</sub> and  $9.1 \pm 66.4\%$  for VLDL<sub>2</sub>, giving rise to 95% limits of agreement between (-109.4% to +111.1%) for VLDL<sub>1</sub> and between (-57.3% to +75.5%) for VLDL<sub>2</sub>. From **Figure 2.3**, a cut-off value of ~3.0 mg.dl<sup>-1</sup> could be made, above which measurements between the two methods appeared to show better agreement for both VLDL<sub>1</sub> and VLDL<sub>2</sub> fractions compared to values less than 3 mg.dl<sup>-1</sup>, where deviation around the mean was more scattered. This is better illustrated in **Figure 2.4**, which shows the %difference (isopropanol minus immunoturbidimetry) in apoB concentrations plotted against the mean of two methods for values  $\geq 3.0$  mg.dl<sup>-1</sup>. In this case, the deviation around the mean is evenly scattered, especially in the VLDL<sub>1</sub> fraction, and the 95% limits of



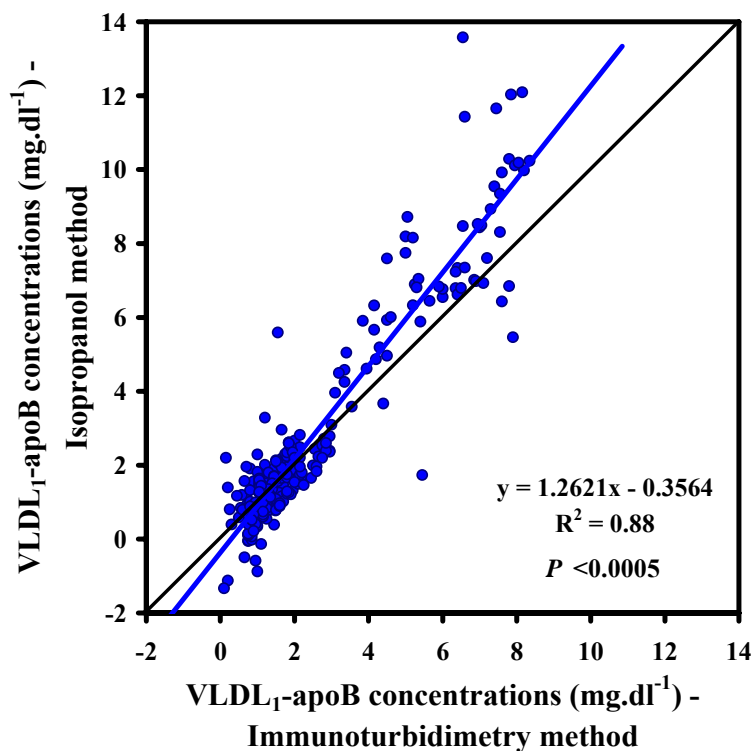
agreement (mean  $\pm$  2SD) became smaller: between -33.3% and 73.7% ( $20.2 \pm 53.5\%$ ,  $n = 65$ ) for VLDL<sub>1</sub> and between -17.5% and 53.0% ( $17.7 \pm 35.3\%$ ,  $n = 84$ ) for VLDL<sub>2</sub>. In addition, it is clear from **Figure 2.4A** that there are two outliers in the VLDL<sub>1</sub> fraction. Although excluding these two measurements did not influence the mean percentage difference, it reduced the SD value ( $20.7 \pm 37.5\%$ ) and consequently, the 95% limits of agreement (-16.8 to 58.2%).

Finally, **Table 2.2** shows the mean VLDL<sub>1</sub>- and VLDL<sub>2</sub>-apoB concentrations, SDs and CVs measured 10 times for each method in VLDL<sub>1</sub> and VLDL<sub>2</sub> fractions using pooled samples. The apoB concentrations were lower in VLDL<sub>1</sub> compared to VLDL<sub>2</sub> and it is clear that the immunoturbidimetric method was more reproducible than the isopropanol method at such low concentrations (CV 3.4% vs. 9.9% for VLDL<sub>1</sub> and 1.4% vs. 3.2% for VLDL<sub>2</sub>, respectively).

**Table 2.2:** Calculated mean, SD and CV for apoB concentrations measured 10 times for the same sample using the two methods

	VLDL <sub>1</sub> -apoB (mg.dl <sup>-1</sup> )		VLDL <sub>2</sub> -apoB (mg.dl <sup>-1</sup> )	
	Immunoturbidimetric method	Isopropanol method	Immunoturbidimetric method	Isopropanol method
<b>Mean</b>	3.74	2.07	6.96	5.65
<b>SD</b>	0.13	0.20	0.10	0.18
<b>CV (%)</b>	<b>3.38</b>	<b>9.88</b>	<b>1.39</b>	<b>3.19</b>

[A]



[B]

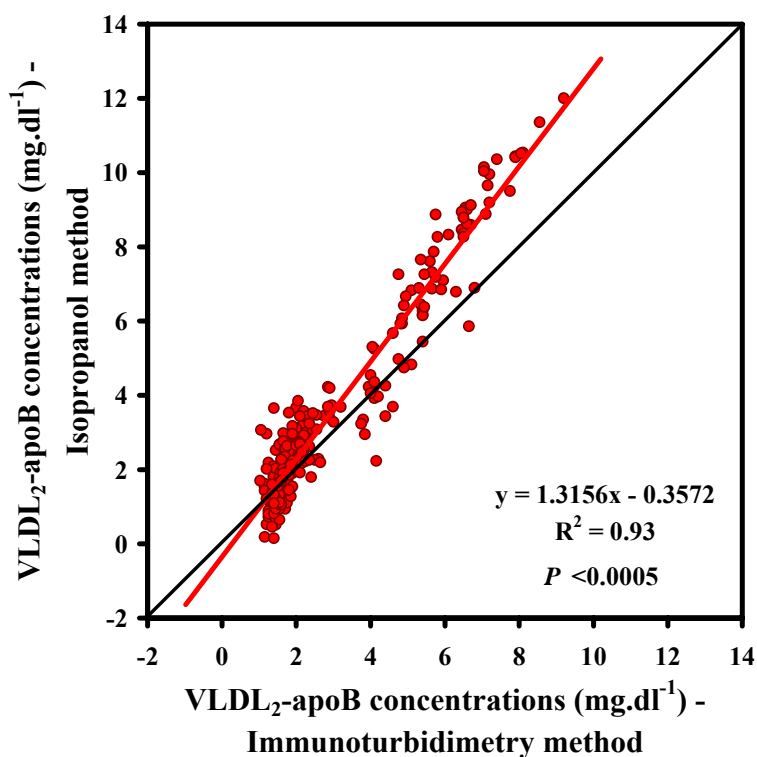


Figure 2.2: ApoB concentrations (mg.dl<sup>-1</sup>, with linear-regression lines of 'best-fit' and equation of the line) in [A] VLDL<sub>1</sub> and [B] VLDL<sub>2</sub> fractions measured using an automated immunoturbidimetric method and by a manual isopropanol method with the line of equality. (n = 260)

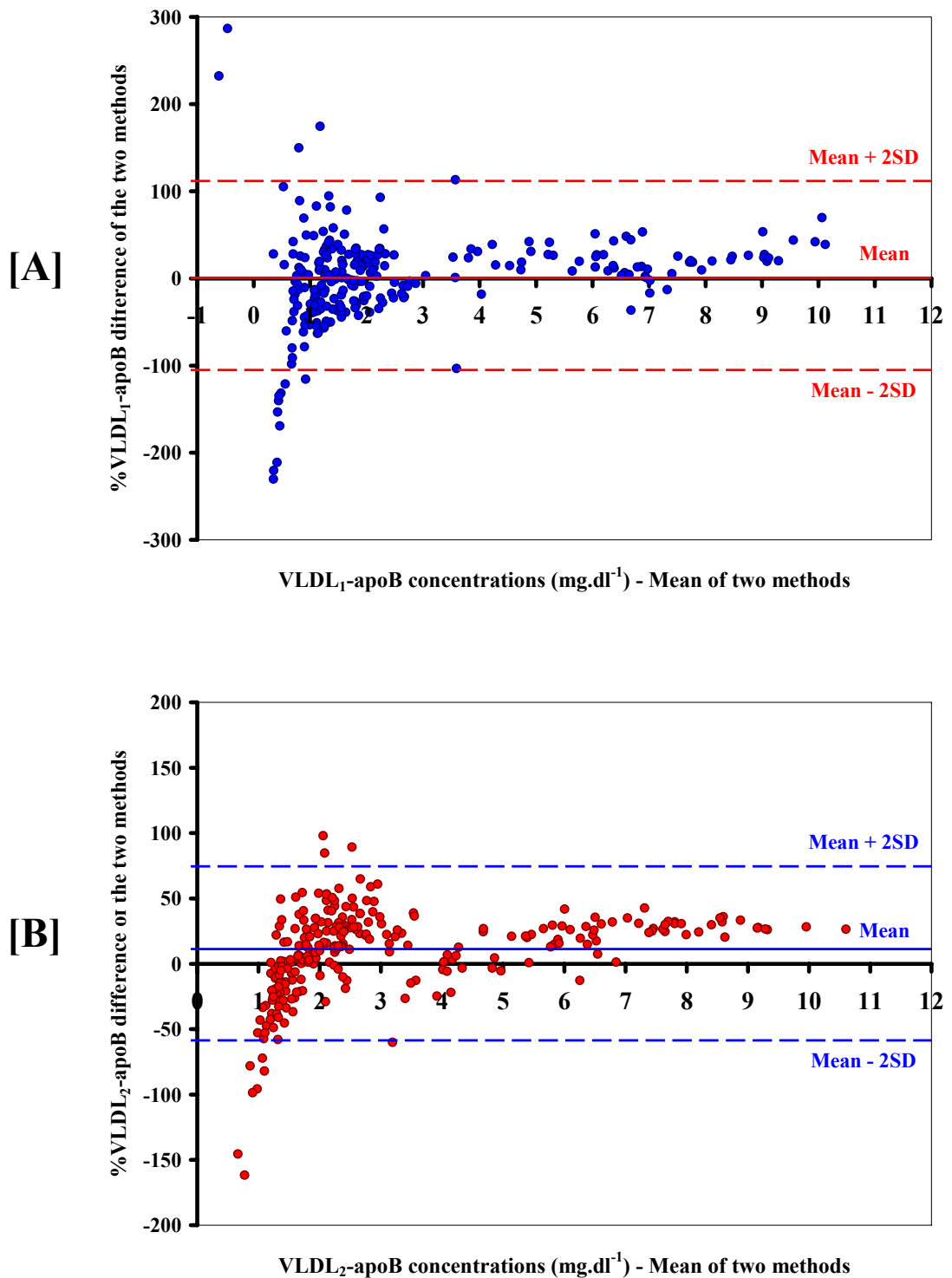


Figure 2.3: A scatterplot of the percent difference in apoB concentrations (mg.dl<sup>-1</sup>) between the two methods (isopropanol minus immunoturbidimetry) against their mean in [A] VLDL<sub>1</sub> and [B] VLDL<sub>2</sub>, with 95% limits of agreement (mean  $\pm$  2SD): VLDL<sub>1</sub> (0.8  $\pm$  110.2%) and VLDL<sub>2</sub> (9.1  $\pm$  66.4%), n = 260.

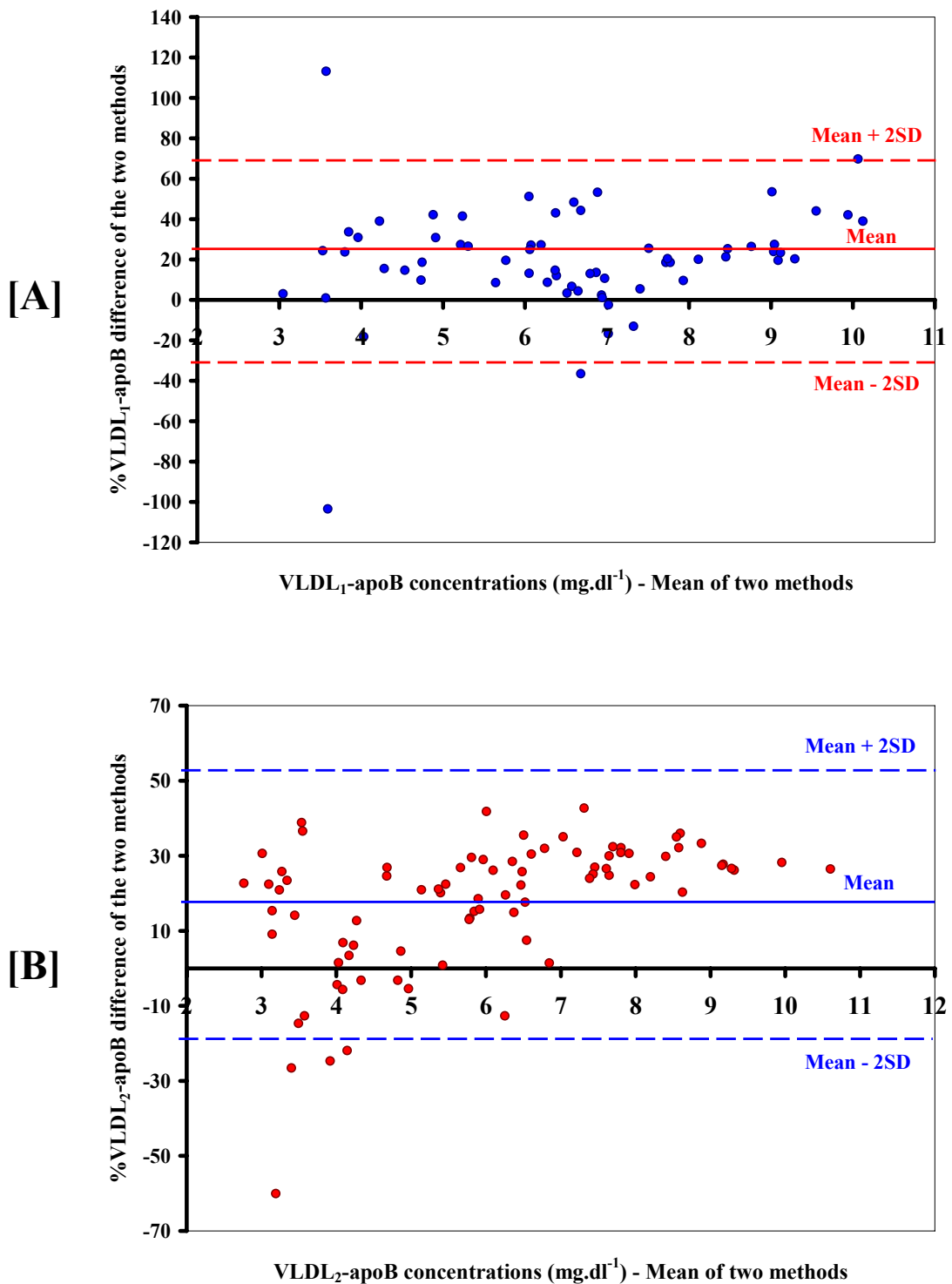


Figure 2.4: A scatterplot of the percent difference (isopropanol minus immunoturbidimetry) for apoB concentrations  $\geq 3.0$  mg.dl<sup>-1</sup> against the mean of the two methods in [A] VLDL<sub>1</sub> and [B] VLDL<sub>2</sub>, with 95% limits of agreement (mean  $\pm$  2SD): VLDL<sub>1</sub> (20.2  $\pm$  53.5%, n = 65) and VLDL<sub>2</sub> (17.7  $\pm$  35.3%, n = 85).

### 2.9.4 Discussion

Due to the low concentrations of apoB in VLDL fractions, the manual precipitation of apoB using isopropanol (Egusa *et al.*, 1983) has historically been the preferred choice, followed by a modification of another manual method (Lowry *et al.*, 1951) to determine the apoB protein concentrations in the fractions. Being a chemical method, this technique is not affected by the size of the lipoprotein particle, the possible masking of the apoB epitopes or changes in time over standardisation of commercial apoB assays. However, this method is time-consuming and requires a high degree of precision and skill in separating the non-apoB supernatant from the precipitated apoB protein, which may sometimes be difficult in cases of very high or very low apoB concentrations.

Conversely, apoB can be measured directly using an automated immunoturbidimetric assay, with the use of commercially available kits. The fact that this method is automated, makes it faster, allows for a minimal involvement of human error, and, consequently, better precision. One possible shortcoming of this immunological method is that it relies on detecting certain epitopes on the apoB particle and, therefore, has the disadvantage of being dependent on the manufacturer's choice of such epitopes, which could be subject to change at any time.

Because of the large number of samples per trial used in this thesis and the relative difficulty of using the manual isopropanol method for apoB measurements, the two methods were compared using 260 samples from the VLDL<sub>1</sub> and VLDL<sub>2</sub> fractions. This was done to assess the possibility of replacing the isopropanol precipitation method with the automated immunoturbidimetric method.

The present results show that the two methods agree well for both fractions with  $R^2$  values of  $\sim 0.9$ . This is especially true for apoB concentrations  $\geq 3.0$  mg.dl<sup>-1</sup>. Although the immunoturbidimetric method seems to give values  $\sim 20\%$  lower than the isopropanol method, this is consistent and the deviation from the mean of the two methods still lies within the 95% limits of agreement. However, the immunoturbidimetric method allows better precision and reproducibility than the manual isopropanol method, with CVs of 3.4% and 1.4% (compared to 9.9% and 3.2%) in VLDL<sub>1</sub> and VLDL<sub>2</sub> fractions, respectively. In addition, being an automated

method, the immunoturbidimetry allows the feasibility and the ease of analysing large number of samples in a relatively short period of time, compared to the isopropanol method.

In conclusion, it is inevitable to find some lack of agreement between different methods, but what matters is the amount by which methods disagree (Bland & Altman, 1999). The automated immunoturbidimetry method correlated consistently and reasonably well with the manual isopropanol precipitation method, with better precision, reproducibility and ease of use for measurements of multiple samples. Therefore, the immunoturbidimetric method was chosen to measure apoB concentrations in this thesis.

## **2.10 Glucose Pilot Study**

The purpose of this pilot study was to determine whether it was possible to maintain a ‘pseudo’ steady state of high circulating glucose and insulin concentrations by ingesting small amounts of a glucose solution at frequent intervals. If this approach was feasible, it would represent a more straightforward and less invasive method of inducing hyperinsulinaemia and hyperglycaemia than invasively infusing glucose and/or insulin. A number of different patterns of glucose ingestion were investigated to determine an appropriate protocol, which will be used in the study described in [Chapter 4](#).

### **2.10.1 Subjects and Methods**

Four subjects (2 women and 2 men – age (range) 20-33 years; body mass 55-95 kg; BMI 22.6-28.1 kg.m<sup>-2</sup>) were included in this study after giving informed consent. Subjects reported in the morning to the metabolic suite at the Institute of Diet, Exercise and Lifestyle, at the University of Glasgow, after a 12-h fast. Subjects were cannulated in an antecubital vein and after a 10-min interval a baseline EDTA fasting blood sample was drawn. A second baseline sample was taken 10 min later. All blood samples were placed immediately in ice. EDTA plasma was obtained after centrifugation for 10 min at 3000 rpm and divided into 5 aliquots in 1.5 ml Eppendroff tubes. Aliquots were stored immediately at -20° C for subsequent analysis of glucose, NEFA and insulin concentrations as previously described in [sections 2.4.2](#) and [2.4.3](#).

The glucose drinks were prepared by dissolving the appropriate weight of glucose powder (Thornton and Ross Ltd., Huddersfield, England) in hot water and making the required volume up with cold water. Approximately 5-6 ml of lemon juice per 100 ml of water were added for taste.

### 2.10.2 Protocol I

The first protocol was designed to introduce a glucose concentration of 30% (w/v) (15 g in 50 ml) every 15 min for 2 h. Two female subjects participated in this protocol (**Table 2.3**). Blood samples were drawn every 15 min.

**Table 2.3: Characteristics of the two female subjects who participated in Protocol I**

Subject	Age (y)	Body mass (kg)	BMI (kg.m <sup>-2</sup> )	HOMA
1	20	66.4	23.81	1.4
2	33	55	22.6	1.6

**Figure 2.5** shows glucose, insulin and NEFA responses for the two subjects. Glucose concentrations increased slowly in subject 1 and immediately in subject 2 before declining close to baseline at the end of 2 h. Consequently, insulin concentrations increased in response to the glucose drink, but differently in the two subjects: while subject 1 had a delayed insulin response corresponding to glucose concentrations, subject 2 had a dramatic increase in insulin concentrations for the duration of the trial. However, none of the two subjects reached any steady state during 2 h. Similarly, NEFA concentrations declined slowly in subject 1 but dramatically in subject 2 over the course of trial in response to insulin. NEFA concentrations were only suppressed at the end of the 2 h.

Despite different responses in glucose and insulin concentrations, neither subjects reached a ‘pseudo’ steady state. Because one subject had a delayed response, it was clear that an initial bolus glucose drink is needed to induce a more rapid response. Also, because the other subject had an exaggerated insulin response to the glucose drink, it might be beneficial to decrease its dose, and possibly increase its frequency to account for the reduced concentration.

### **2.10.3 Protocol II**

The second protocol started with a bolus drink of 20% glucose (30 g in 150 ml) followed by 20% glucose drink (5 g in 25 ml) every 10 min (instead of 15 min) for a period of 2 h. Blood samples were taken at 15-min intervals. One male subject participated in this protocol (aged 29 years, body mass 95 kg, BMI 28.1 kg.m<sup>-2</sup> and HOMA 1.4).

**Figure 2.6** shows glucose, insulin and NEFA responses during this protocol. The bolus glucose drink successfully increased glucose and insulin concentrations shortly after ingestion. However, the glucose drink, thereafter, was not sufficient to maintain a steady state of glucose and insulin as they started to decline after 90 min. In addition, the drink did not induce a desirable suppression in NEFA concentrations before the 2 h.

Therefore, a bolus glucose drink seemed essential to induce a faster and immediate response in glucose and insulin concentrations. However, a dose of 5 g glucose every 10 minutes did not seem sufficient to maintain elevated glucose and insulin concentrations. Furthermore, it would be helpful to increase the observation time of the study to investigate whether a steady state could be reached and/or maintained for a longer period of time.

### **2.10.4 Protocol III**

In this protocol, glucose drink concentration and the observation time were increased. It started with 30% bolus glucose drink (30 g in 100 ml) followed by 60% glucose drink (15 g in 25 ml) every 15 min for 4 h. These doses were chosen as it became obvious from the previous two protocols that a stronger glucose drink might be needed to achieve the desired ‘pseudo’ steady state and suppress NEFA at a time shorter than 2 h. One subject participated in this protocol (aged 22 years, body mass 80.5 kg, BMI 27.8 kg.m<sup>-2</sup> and HOMA 4.0).

**Figure 2.7** shows glucose, insulin and NEFA responses during the 4-h observation period for this protocol. Glucose and insulin concentrations increased quickly in response to the bolus glucose drink, which caused a desirable, fast reduction in NEFA concentrations. However, the glucose drink caused abrupt responses in both



glucose and insulin concentrations. In addition, there was tendency for insulin concentrations to continually increase during the 4 h. It is uncertain, however, whether this increase is related to the subject's HOMA of 4.0, which may indicate insulin resistance, or whether it was caused by the increased glucose drink concentration.

#### **2.10.5 Conclusion**

From these protocols, it was obvious that (1) a bolus glucose drink was needed to trigger an immediate insulin response, (2) providing 5 g of glucose every 10 minutes was not sufficient to maintain elevated levels of glucose and insulin, while providing 15 g glucose every 15 minutes produced abrupt responses and (3) although NEFA concentrations started to decline immediately in response to insulin, it took about 60 min to be suppressed almost completely in the last protocol.

In conclusion, taking these findings into account, it was decided that the final protocol for the glucose ingestion would be; a bolus glucose drink of 25% (w/v) (30 g of glucose in 120 ml), followed by 25% glucose drink (10 g in 40 ml) every 15 min for the duration of the trial. The glucose ingestion protocol would start an hour prior to the Intralipid infusion described in [Chapter 4](#) to ensure adequate suppression of NEFA concentrations.

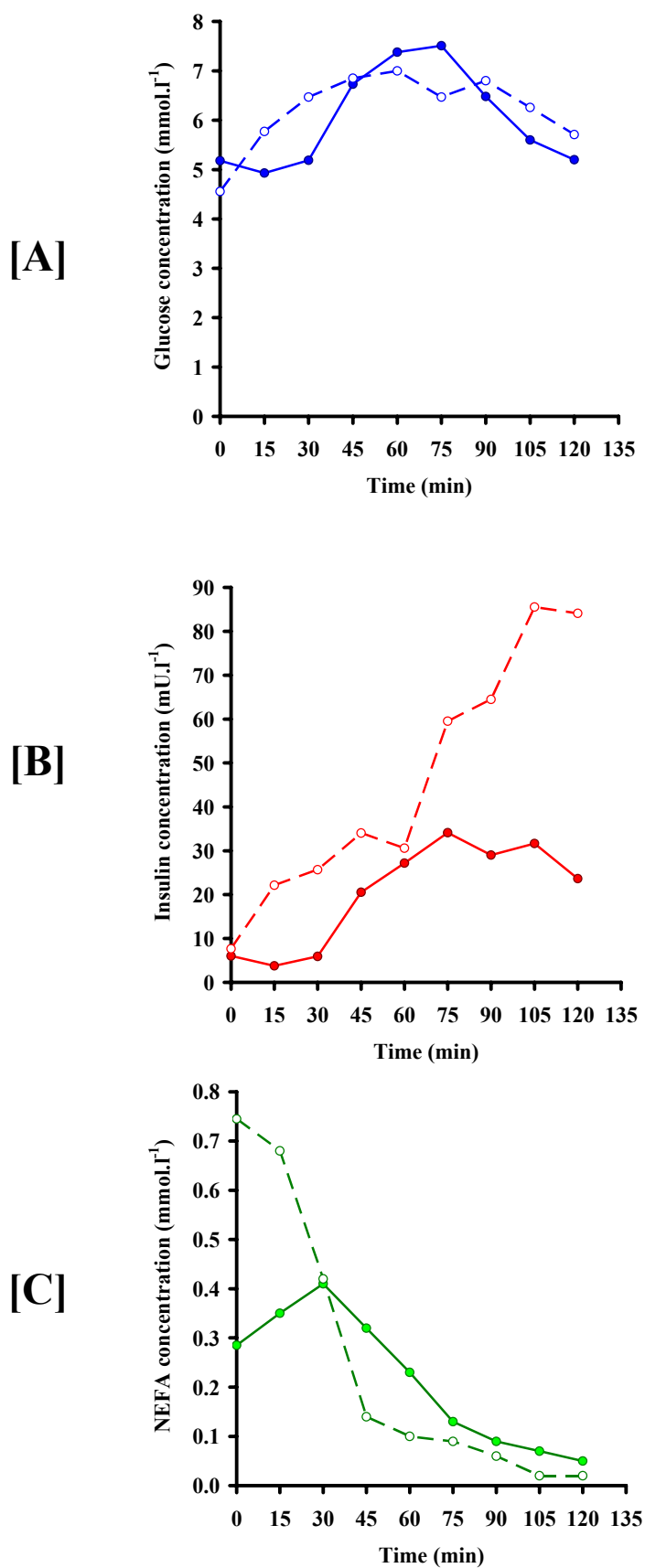
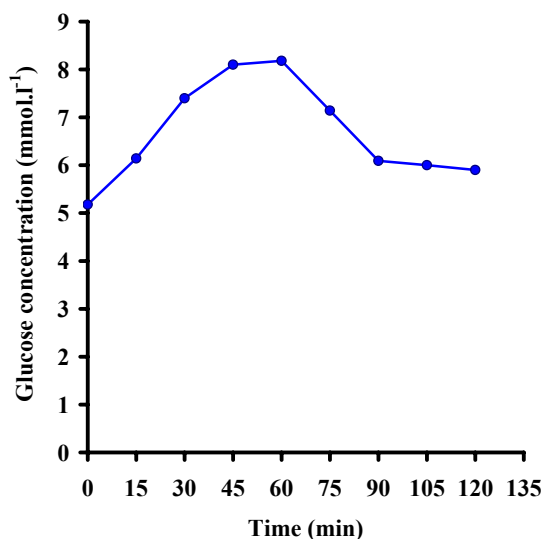
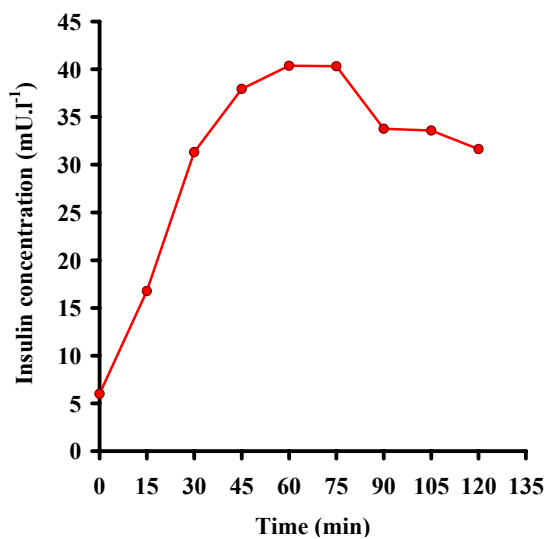


Figure 2.5: [A] glucose, [B] insulin and [C] NEFA response ( $n = 2$ ) for the glucose pilot - Protocol I: 30% (w/v) glucose (15 g in 50 ml) every 15 min for 2 h. The solid lines represent subject 1, while the dotted lines represent subject 2.

[A]



[B]



[C]

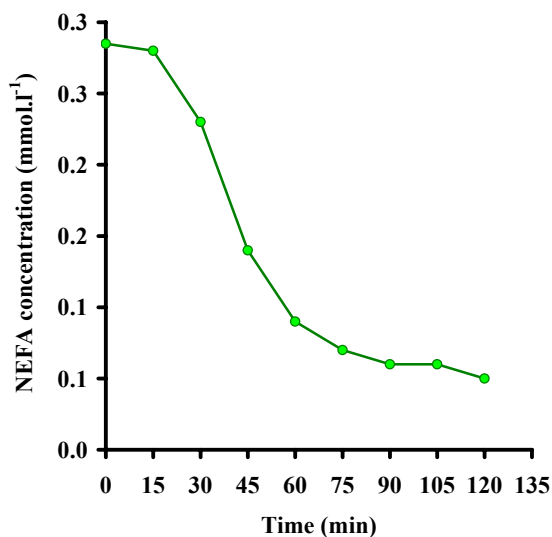
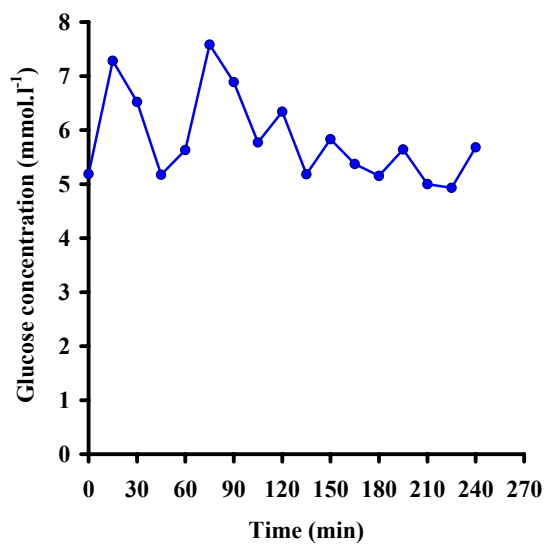
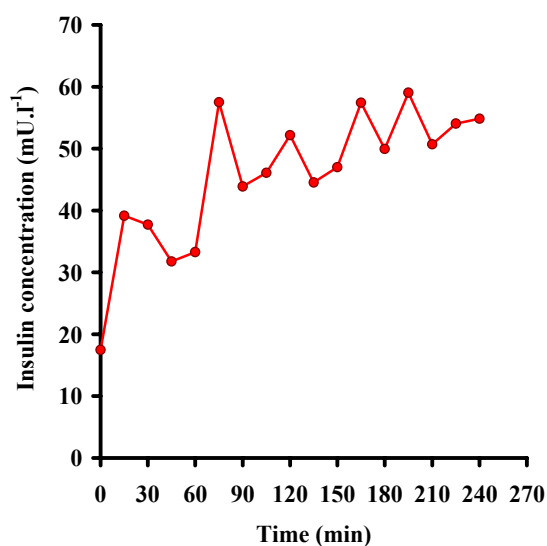


Figure 2.6: [A] glucose, [B] insulin and [C] NEFA response ( $n = 1$ ) for the glucose pilot - Protocol II: a bolus drink of 20% (w/v) of glucose (30 g in 150 ml) followed by a 20% glucose drink (5 g in 25 ml) every 10 min for 2 h.

[A]



[B]



[C]

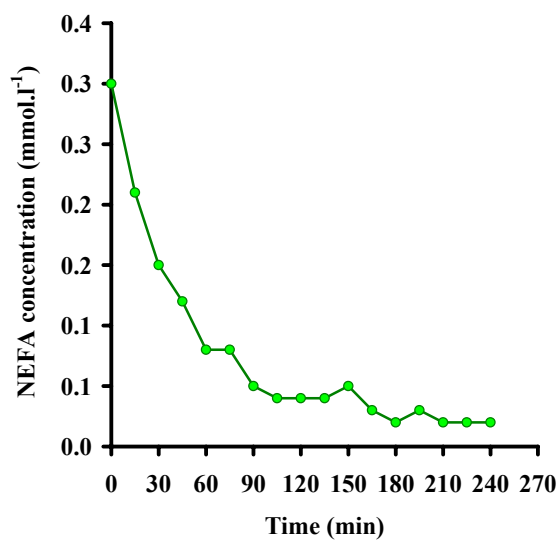


Figure 2.7: [A] glucose, [B] insulin and [C] NEFA response ( $n = 1$ ) for the glucose pilot - Protocol III: a bolus glucose drink of 30 g (in 100 ml) followed by 15 g of glucose (in 25 ml) every 15 min for 4 hours.

### 3. Development of a Novel Method to Determine VLDL<sub>1</sub> Kinetics

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#### 3.1 Introduction

A large body of evidence suggests that increased circulating concentrations of TRLs increase the risk of atherosclerosis (Malloy & Kane, 2001; Tanaka *et al.*, 2001). This is particularly evident in the postprandial state (Karpe, 1999; Schneeman *et al.*, 1993). However, the measurement of a high TRL concentration provides no information regarding the mechanisms responsible for this elevation; i.e. increased rate of synthesis and/or reduced rate of catabolism. As it is important to understand the mechanisms responsible for increased TRL concentrations in different metabolic states, both to advance basic scientific understanding and to help guide therapeutic treatments, studies investigating kinetics of TRL can yield useful data. Such an approach has, for example, revealed that the dyslipidaemia associated with insulin resistance and type 2 diabetes is largely due to an overproduction of hepatically derived large VLDL<sub>1</sub> (S<sub>f</sub> 60-400) (Watson *et al.*, 2003; Adiels *et al.*, 2005a). These studies typically use precursors labelled with stable or radioactive isotope tracers to measure the synthesis of lipids and apolipoproteins directly (Adiels *et al.*, 2005a; Gill *et al.*, 2004a; Packard *et al.*, 2000; Packard, 1995). Although these techniques yield detailed kinetic data, they are costly, time consuming, labour-intensive and require the use of specialised equipment and techniques in research laboratories.

The aim of the present study was, therefore, to develop a relatively straightforward method of obtaining TRL kinetic data. The method relies on the fact that chylomicrons compete with VLDL<sub>1</sub> particles for the same catalytic pathway; i.e. hydrolysis of their TG content by the action of LPL. Previous studies (Karpe & Hultin, 1995; Björkegren *et al.*, 1996) have shown that VLDL<sub>1</sub> accumulates in plasma after fat ingestion or intravenous infusion of a lipid emulsion (e.g. Intralipid) due to the presence of the newly secreted chylomicrons or chylomicron-like particles, which are the preferred substrate for LPL because of their larger size and TG content (Fisher *et al.*, 1995). Indeed, using stable isotope methods, Björkegren *et*

*al.* demonstrated that infusion of Intralipid prevents over 90% of VLDL<sub>1</sub> catabolism (Bjorkegren *et al.*, 1996). It was therefore hypothesized in the present study that it would be possible to calculate the production rates of VLDL<sub>1</sub>-TG and -apoB from the rate of their accumulation during an infusion of Intralipid. The former would be a measure of lipid production while the latter would represent the rate of VLDL<sub>1</sub> particle production, as there is one apoB molecule per VLDL<sub>1</sub> particle (Elovson *et al.*, 1988). Furthermore, using this approach, it is possible to calculate the rate of Intralipid-TG clearance (a surrogate measure of TRL-TG clearance) from either the steady-state Intralipid-TG concentration during infusion (Rang *et al.*, 2003) or from the exponential decay in Intralipid-TG concentration post-infusion (Rossner, 1974). Here the author reports the development and validation of this 'Intralipid method' to determine TRL kinetics.

## 3.2 Material and Methods

### 3.2.1 Subjects

Ten non-smoking healthy subjects (7 males and 3 females) were included in this study after giving written informed consent. All subjects had normal thyroid, liver and renal function and none had acute illness, a history of known cardiovascular disease and hypertension, nor were under medication known to influence carbohydrate or lipid metabolism. The subject information sheet, consent form and health screen for the present study are shown in Appendices A1 and B1. Two subjects had the E3/2 apoE phenotype, seven had the E3/3 phenotype and one had the E4/3 phenotype. The subjects' characteristics are shown in **Table 3.1**. Subjects were requested not to exercise for three days before their study days as this is known to affect TRL metabolism (Gill & Hardman, 2003). In addition, they were asked to weigh and record their dietary intake for two days prior to the Intralipid test and this diet was replicated in those subjects who underwent a second Intralipid test. The study protocol was approved by the Research Ethics Committee of the North Glasgow University Hospitals NHS Trust.

**Table 3.1: Subjects' physical and metabolic characteristics (n=10).**

	Mean	Range
Age (years)	33.5	(20.0 – 55.0)
Body mass index (kg.m <sup>-2</sup> )	25.9	(20.8 – 34.7)
Waist circumference (cm)	85.1	(65.0 – 113.5)
Waist/hip ratio	0.84	(0.71 – 1.04)
Triglycerides (mmol.l <sup>-1</sup> )	1.36	(0.40 – 4.43)
Total cholesterol (mmol.l <sup>-1</sup> )	4.15	(2.85 – 5.90)
HDL cholesterol (mmol.l <sup>-1</sup> )	1.30	(0.75 – 1.85)
LDL cholesterol (mmol.l <sup>-1</sup> )	2.20	(1.22 – 4.00)
Glucose (mmol.l <sup>-1</sup> )	5.5	(4.3 – 8.0)
Insulin (mU.l <sup>-1</sup> )	8.92	(2.78 – 24.81)
HOMA <sub>IR</sub>	2.41	(0.56 - 8.82)
NEFA (mmol.l <sup>-1</sup> )	0.51	(0.34 – 0.70)
ALT (U.l <sup>-1</sup> )	23	(16-38)

### 3.2.2 Intravenous Intralipid Test

Each subject reported to the Clinical Investigation Suite in the Department of Vascular Biochemistry in Glasgow Royal Infirmary after an overnight fast of 12 h. Transportation to the hospital was provided for the subjects, when needed, to ensure that they arrived in a rested state. A cannula was introduced into an antecubital vein in both arms; one for administration of Intralipid (purified soybean oil emulsion, Fresenius Kabi Ltd., Warrington, UK) and the other for blood sampling. The cannulae were kept patent by flushing with non-heparinised saline solution (0.9% NaCl). Ten min after cannulation a first baseline blood sample was obtained. A second baseline blood sample was obtained 10 min later.

The intravenous Intralipid test used was a modification of that described by Björkegren and colleagues (Björkegren *et al.*, 1996). A bolus dose of 20% Intralipid (0.1 g.kg<sup>-1</sup> body mass) was injected within one min. This was followed immediately by a constant continuous infusion of 10% Intralipid (0.1 g.kg<sup>-1</sup>.h<sup>-1</sup>). This dose was chosen as Björkegren *et al.* reported that the rate of rise of VLDL<sub>1</sub>-apoB during Intralipid infusion was no greater for a 0.2 g.kg<sup>-1</sup>.h<sup>-1</sup> infusion dose compared with 0.1

$\text{g.kg}^{-1}.\text{h}^{-1}$ , suggesting that the lower dose was sufficient to saturate LPL and prevent measurable VLDL<sub>1</sub> catabolism (Bjorkegren *et al.*, 1996). However, experiments with the  $0.2 \text{ g.kg}^{-1}.\text{h}^{-1}$  dose were also performed in the present study to confirm that this was the case in the author's hands (see below). Initially, the infusion period was 120 min, however, during development of the technique this was subsequently shortened to 75 min after it became clear that a 75-min infusion was long enough to induce a sufficient measurable rise in VLDL<sub>1</sub>-TG and -apoB. **Figure 3.1** shows an example of a subject during an Intralipid trial.

Blood samples were obtained at 15-min intervals during the infusion. Further blood samples were drawn 2.5, 5, 10, 15, 20, 30, 45, 60 and 75 min post-infusion. Initially, the post-infusion period was 3.25 h. However, this was subsequently shortened to 75 min when it became clear that this was sufficient to calculate the Intralipid-TG clearance rate using the exponential decay. All samples were obtained directly into 10-ml potassium EDTA tubes (BD Vacutainer Systems, Plymouth, UK) and placed immediately in ice before centrifuging for 15 min at 3000 rpm and 4°C.

Aliquots of plasma were frozen immediately at -70°C, as described in [section 2.3](#), for subsequent analysis of insulin, NEFA, glucose, TG, and total and HDL-cholesterol. The remaining plasma was stored overnight at 4°C prior to separation of Intralipid and lipoproteins.

### 3.2.3 Increasing the Intralipid Infusion Rate

Five subjects (2 females and 3 males, age 23-47 y [range], BMI 20.8-28.7  $\text{kg.m}^{-2}$ , fasting TG 0.47-2.45  $\text{mmol.l}^{-1}$ ) underwent a second test using a higher infusion dose ( $0.2 \text{ g.kg}^{-1}.\text{h}^{-1}$ ) of 10% Intralipid with the same  $0.1 \text{ g.kg}^{-1}$  bolus dose. This was carried out to determine whether an Intralipid infusion dose of  $0.1 \text{ g.kg}^{-1}.\text{h}^{-1}$  was sufficient to completely prevent measurable lipolysis of VLDL<sub>1</sub> by LPL, and therefore enable determination of VLDL<sub>1</sub>-TG and -apoB production rates from their rises in concentration. If the infusion rate of  $0.1 \text{ g.kg}^{-1}.\text{h}^{-1}$  was sufficient to saturate LPL and block lipolysis of VLDL<sub>1</sub>, the  $0.2 \text{ g.kg}^{-1}.\text{h}^{-1}$  dose would not result in higher calculated production rates of VLDL<sub>1</sub>-TG or -apoB compared to the  $0.1 \text{ g.kg}^{-1}.\text{h}^{-1}$  dose. The order of testing was randomized. Other than the higher infusion dose, all conditions of the tests were the same.





**Figure 3.1: A picture of a subject during an Intralipid trial.** A cannula is placed in an antecubital vein in each arm; one for the Intralipid infusion of  $0.1 \text{ g.kg}^{-1}.\text{h}^{-1}$  (right arm) using an IVAC Signature Edition Gold pump (ALARIS Medical Systems, Inc.) and one for blood sampling (left arm). An initial bolus Intralipid dose ( $0.1 \text{ g.kg}^{-1}$  body mass) was given in the right arm via a three-way stopcock immediately before starting the infusion. (Published with permission from the subject).

### 3.2.4 Intralipid ( $S_f > 400$ ) Separation from Whole Plasma

Two ml of plasma were overlaid with 4 ml of 1.006 g.ml<sup>-1</sup> density solution in ultra-clear centrifuge tubes and spun at 10,000 rpm and 4° C for 30 min (Lindgren *et al.*, 1972) using Beckman L8-60M Ultracentrifuge and Beckman 50.4 rotor (Beckman Instruments Inc., UK). Intralipid ( $d < 1.006$  g.ml<sup>-1</sup>) was removed in the top 2 ml (IL-1) for subsequent measurements of TG using commercially available kits as described in [section 2.4.2](#). TG concentration was also measured in the middle 1.5 ml fraction (IL-2) to verify complete separation of Intralipid. The final Intralipid-TG concentration was calculated as the addition of these two fractions [IL-1 + (IL-2 × 1.5/2)]. In addition, glycerol was measured in these IL-1 and IL-2 fractions ([section 2.4.2](#)) to determine the amount of free glycerol. The final 0.5 ml of the density solution was discarded and the remaining 2 ml Intralipid-free plasma was used for separation of VLDL<sub>1</sub> and VLDL<sub>2</sub>. The CV for the Intralipid-TG separation was 6.9%.

### 3.2.5 VLDL<sub>1</sub> and VLDL<sub>2</sub> Separation

VLDL<sub>1</sub> ( $S_f$  60-400) and VLDL<sub>2</sub> ( $S_f$  20-60) were isolated from plasma using a modification of the cumulative ultracentrifugation density gradient technique as previously described in [section 2.4.1](#). TG concentrations were then measured in the VLDL<sub>1</sub> and VLDL<sub>2</sub> fractions in all time points and apoB concentrations were also measured directly by immunoturbidimetry as described in [section 2.9.2.3](#). The CVs for separation of VLDL<sub>1</sub>-TG and -apoB were 5.0% and 3.4%, respectively and for VLDL<sub>2</sub>-TG and -apoB were 5.8% and 1.4%, respectively.

### 3.2.6 Fasting Plasma Analysis

Plasma glucose, insulin, total and HDL cholesterol concentrations were analysed in the fasted state and TG and NEFA concentrations were analysed at all time points. LDL cholesterol was calculated in the fasted state using the Friedewald equation (Friedewald *et al.*, 1972). Serum ALT concentrations were measured at screening. All analyses were done as previously described in [section 2.4.2](#).

### 3.2.7 Correction for Glycerol

Enzymatic kits for TG analysis measure the glycerol that is hydrolyzed from TG by LPL. As Intralipid contains free glycerol as an excipient, it has been reported that it overestimates the true TG concentrations of Intralipid (Howdieshell *et al.*, 1995). Therefore, all Intralipid-TG measurements were corrected for free glycerol and are reported as ‘true’ TG concentrations [true TG concentration (mmol.l<sup>-1</sup>) = measured TG (mmol.l<sup>-1</sup>) - glycerol (mmol.l<sup>-1</sup>)]. Glycerol concentrations were also measured in 5 subjects in the VLDL<sub>1</sub> fraction during infusion and were found to be negligible (influencing VLDL<sub>1</sub>-TG concentrations by less than 1%).

### 3.2.8 Kinetic Data Calculations

The clearance rates of Intralipid-TG and production rates of VLDL<sub>1</sub>-TG and -apoB were calculated as described below using examples from individual subjects.

#### 3.2.8.1 Calculating VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-ApoB Production Rates

The production rates (mg.h<sup>-1</sup>) of VLDL<sub>1</sub>-TG and -apoB were calculated from the gradient of the linear rise in their pools (total mg in plasma) over time (min) multiplied by 60 min. Pool sizes (mg) were calculated as concentrations in mg.dl<sup>-1</sup> multiplied by plasma volume [4% of body mass (Packard *et al.*, 1984)] in decilitres.

**Figure 3.2A** represents the linear increase in TG pool (mg) in the VLDL<sub>1</sub> fraction of subject No.3 (female, 55 y, 84.5 kg) with R<sup>2</sup> (goodness-of-fit) value of 0.97 and a gradient of 35.8. Thus, the VLDL<sub>1</sub>-TG production rate of this subject would be (35.8 × 60) 2147.4 mg.h<sup>-1</sup> (609.9 mg.kg<sup>-1</sup>.d<sup>-1</sup>). Similarly, from **Figure 3.2B**, the VLDL<sub>1</sub>-apoB production rate of the same subject was (0.85 × 60) 50.9 mg.h<sup>-1</sup> (14.5 mg.kg<sup>-1</sup>.d<sup>-1</sup>).

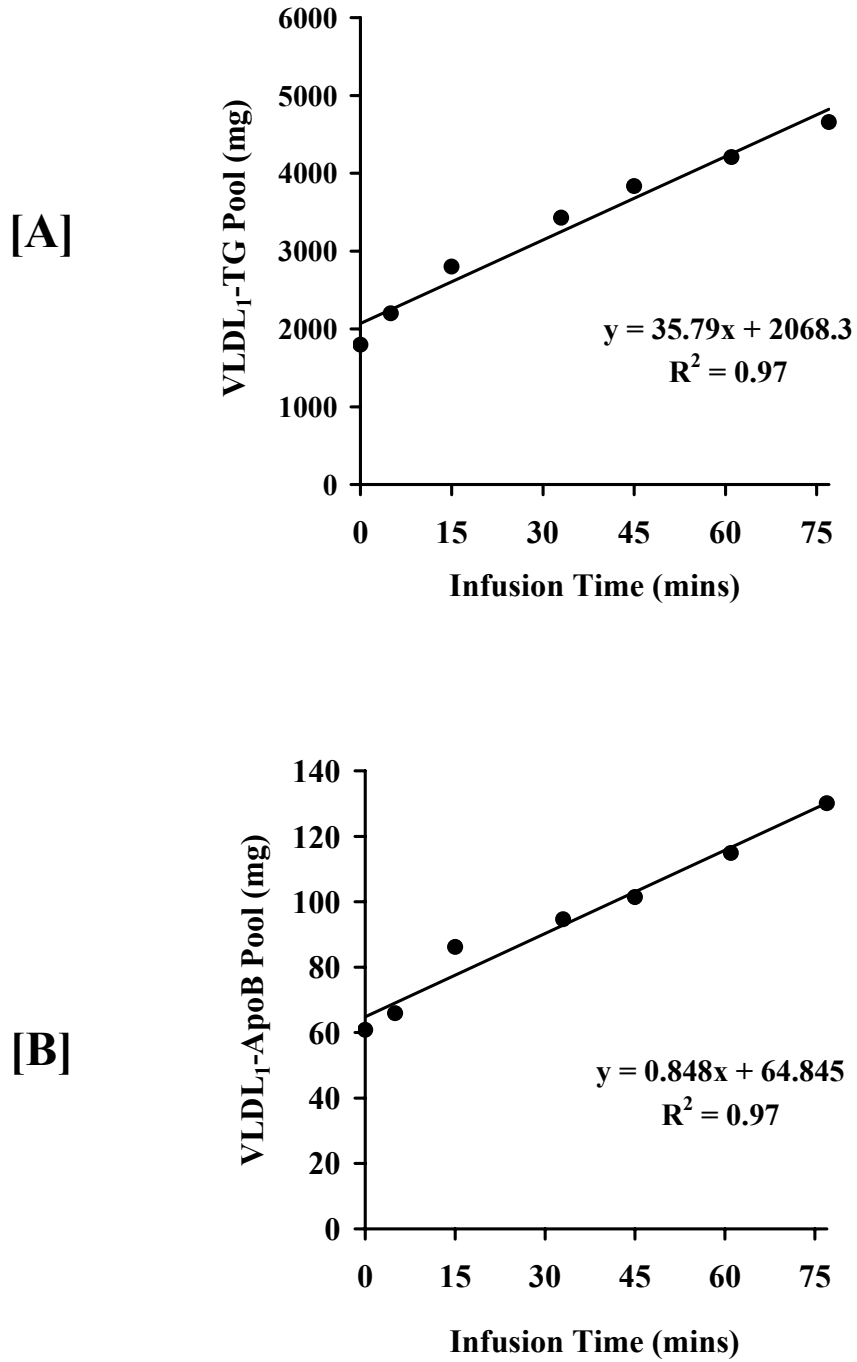


Figure 3.2: Changes in [A] VLDL<sub>1</sub>-TG and [B] VLDL<sub>1</sub>-apoB pools (mg) over time in a female subject No. 3 (55 yr, 84.5 kg) during infusion of 10% Intralipid (0.1 g.kg<sup>-1</sup>.h<sup>-1</sup>). An initial bolus dose of 20% Intralipid (0.1 g.kg<sup>-1</sup> body mass) was given at 0 min.

### 3.2.8.2 Calculating VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-ApoB Fractional Synthetic and Catabolic Rates

The fractional synthetic rate (FSR) is defined as the rate of incorporation of a precursor into a product per unit of product mass (Foster *et al.*, 1993), which can be calculated as:

$$\text{FSR} = \frac{\text{initial rate of change in product}}{\text{initial precursor concentration}} \quad (\text{Eq. 1})$$

In this method, the FSR of VLDL<sub>1</sub>-TG and -apoB (pools.d<sup>-1</sup>) can be calculated from the gradient of the linear rise in their pool (mg) over time (min) divided by fasting pool size (mg) then multiplied by 60 and 24 (to account for number of minutes in 1 day). However, since the system is in a steady state in fasting conditions, where synthesis of VLDL<sub>1</sub> equals its clearance, FSR will also be equivalent to the fractional clearance rate (FCR) in the fasted state (Foster *et al.*, 1993).

Subject No. 3 had a VLDL<sub>1</sub>-TG and -apoB fasting pool sizes of 1669.5 and 61.7 mg, respectively. According to the equation of the line in **Figure 3.2A**, the subject's VLDL<sub>1</sub>-TG FSR and FCR were  $(35.8 \div 1669.5 \times 60 \times 24)$  30.9 pools.d<sup>-1</sup>. Similarly, from **Figure 3.2B**, VLDL<sub>1</sub>-apoB FSR and FCR were 19.8 pools.d<sup>-1</sup>.

### 3.2.8.3 Calculating Intralipid-TG Clearance Rate

Assuming that all TG clearance is Intralipid-TG clearance, it is possible to determine Intralipid-TG clearance rate in two ways:

#### 3.2.8.3.1 The Steady-State (SS) Method

The clearance rate of Intralipid-TG can be calculated from the steady-state concentration during infusion using the following equation (Rang *et al.*, 2003):

$$\text{Clearance rate (ml.min}^{-1}\text{)} = \frac{\text{infusion rate (mg.min}^{-1}\text{)}}{\text{steady state concentration (mg.ml}^{-1}\text{)}} \quad (\text{Eq. 2})$$

In this method, steady state is defined as being achieved when the final 3 values of the Intralipid-TG concentrations differed by less than 13.8% (i.e. two times the CV for the separation of the Intralipid fraction and measurement of the TG. This

represents the 95% confidence interval for the measured value). **Figure 3.3A** shows the concentration of Intralipid-TG  $\pm 13.8\%$  during a 75-min infusion for subject No.8 (female, 23 y, 52.0 kg, BMI 20.8 kg.m<sup>-2</sup>). The steady state concentration, i.e. the mean of the last 3 points of infusion, is 0.86 mmol.l<sup>-1</sup> (0.76 mg.ml<sup>-1</sup>). Since this subject's body mass is 52.0 kg, the infusion rate of 0.1 g of 10% Intralipid per kg of body mass per h equals:

$$\frac{0.1 \text{ g.h}^{-1}}{\text{kg}} \times 52.0 \text{ kg} = 5.2 \text{ g.h}^{-1} = 5200 \text{ mg.h}^{-1} = 86.67 \text{ mg.min}^{-1}$$

From equation (2):

$$\text{Clearance rate} = \frac{86.67 \text{ mg.min}^{-1}}{0.76 \text{ mg.ml}^{-1}} = 113.7 \text{ ml.min}^{-1}$$

Assuming plasma volume [4% of body mass, (Packard *et al.*, 1984)] for this subject was 2080 ml, the clearance rate can also be expressed in pools.d<sup>-1</sup> as follows:

$$\text{Intralipid-TG clearance rate} = \frac{113.7 \text{ ml.min}^{-1}}{2080 \text{ ml}} \times 60 \text{ min} \times 24 \text{ h} = 78.9 \text{ pools.d}^{-1}$$

### 3.2.8.3.2 The Exponential (Exp) Method

After stopping the intravenous infusion, Intralipid-TG declines exponentially following first-order kinetics as described by Rössner (Rossner, 1974). **Figure 3.3B** shows the Intralipid-TG concentrations (mmol.l<sup>-1</sup>) post-infusion over time (min) curve for the same female subject (No.8) on a semi-log scale. (In this case, data for the first 45 min post-infusion were used for calculation, as beyond this Intralipid-TG concentrations approached zero increasing the error associated with an exponential curve fit). The equation of the fitted line is:

$$y = k e^{-bt} \quad (\text{Eq. 3})$$

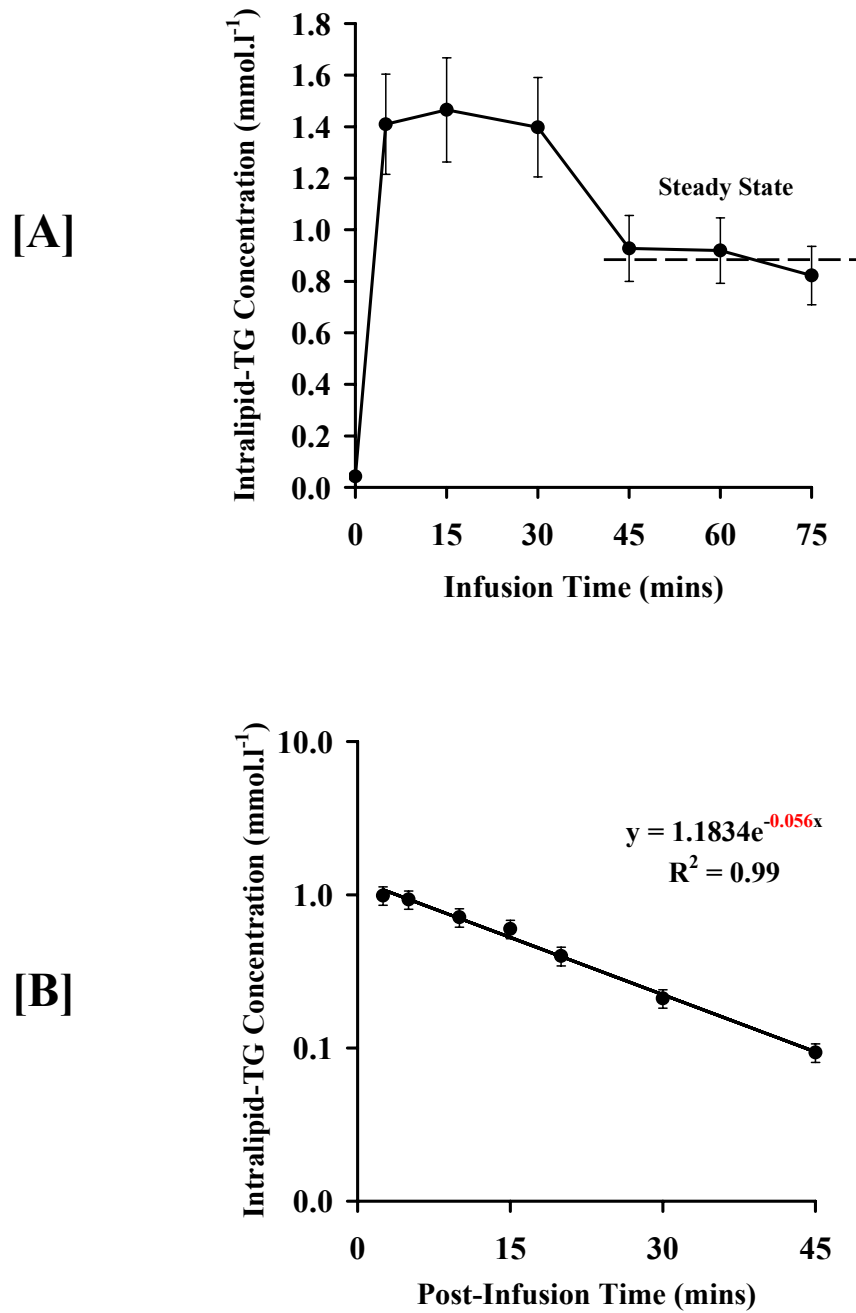


Figure 3.3: Intralipid-TG concentrations (mmol.l<sup>-1</sup>) [A] during infusion and [B] post-infusion of 0.1 g.kg<sup>-1</sup>.h<sup>-1</sup> of 10% Intralipid for a female subject (23 y, 52.0 kg). An initial bolus dose of 20% Intralipid (0.1 g.kg<sup>-1</sup> body mass) was given at 0 min. The Y-axis error bars are  $\pm 13.8\%$  (i.e. the 95% confidence interval for the measured Intralipid-TG concentration).

where  $k$  is the proportionality constant,  $t$  is the time, and  $b$  is the exponential decay constant, which in turn is defined as:

$$b = \frac{\text{clearance rate (ml.min}^{-1}\text{)}}{\text{plasma distribution volume (ml)}} \quad (\text{Eq. 4})$$

Hence,

$$\text{Clearance rate (ml.min}^{-1}\text{)} = b \times \text{plasma volume (ml)} \quad (\text{Eq. 5})$$

$$\text{or} \quad \text{in pools.d}^{-1} = b \times 60 \text{ min} \times 24 \text{ h}$$

Since the  $b$  value from **Figure 3.3B** is 0.056 and the subject's plasma volume was 2080 ml, then from equation (5):

$$\begin{aligned} \text{Intralipid-TG clearance rate} &= 0.056 \times 2080 \text{ ml} = 116.5 \text{ ml.min}^{-1} \\ \text{or} \quad &= b \times 60 \text{ min} \times 24 \text{ h} = 80.6 \text{ pools.d}^{-1} \end{aligned}$$

**Figure 3.4** shows a schematic diagram of the Intralipid method protocol. (For comparison purposes with the stable-isotope method see **Figure 1.7**).

### 3.2.9 Intralipid Recovery

In order to assess the recovery of plasma-Intralipid in the Intralipid fraction ( $S_f > 400$ ), EDTA plasma was spiked with Intralipid to produce an Intralipid-TG concentration in plasma of  $\sim 1.5 \text{ mmol.l}^{-1}$  and  $\sim 4 \text{ mmol.l}^{-1}$ . These reflect approximate Intralipid-TG concentrations at the  $0.1 \text{ g.kg}^{-1}.\text{h}^{-1}$  and  $0.2 \text{ g.kg}^{-1}.\text{h}^{-1}$  infusion doses. For each Intralipid concentration, samples of spiked plasma were divided into 10 aliquots and the Intralipid fractions were separated as described in [section 3.2.4](#). TG and glycerol concentrations were measured in plasma before and after addition of the Intralipid (to calculate the actual Intralipid-TG concentration), as well as in the separated Intralipid fractions. The Intralipid recovery was calculated as follows:

$$\% \text{ recovery} = \frac{\text{separated Intralipid-TG}}{\text{actual Intralipid-TG}} \times 100 \quad (\text{Eq. 6})$$

where actual Intralipid-TG = total TG (plasma with Intralipid) – TG (Intralipid-free plasma).



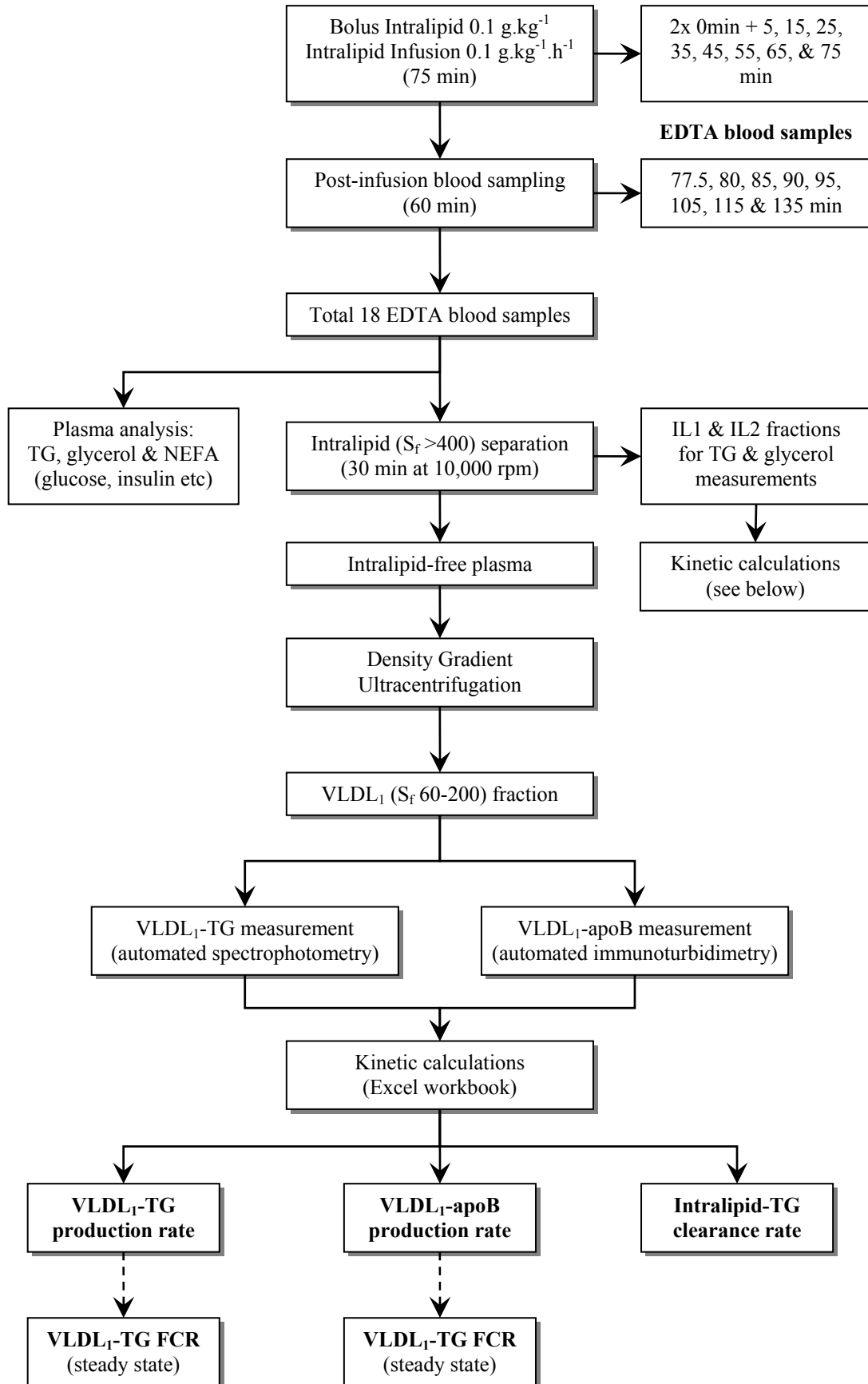


Figure 3.4: Schematic diagram of the 'Intralipid Method' protocol

### 3.2.10 Statistical Analyses

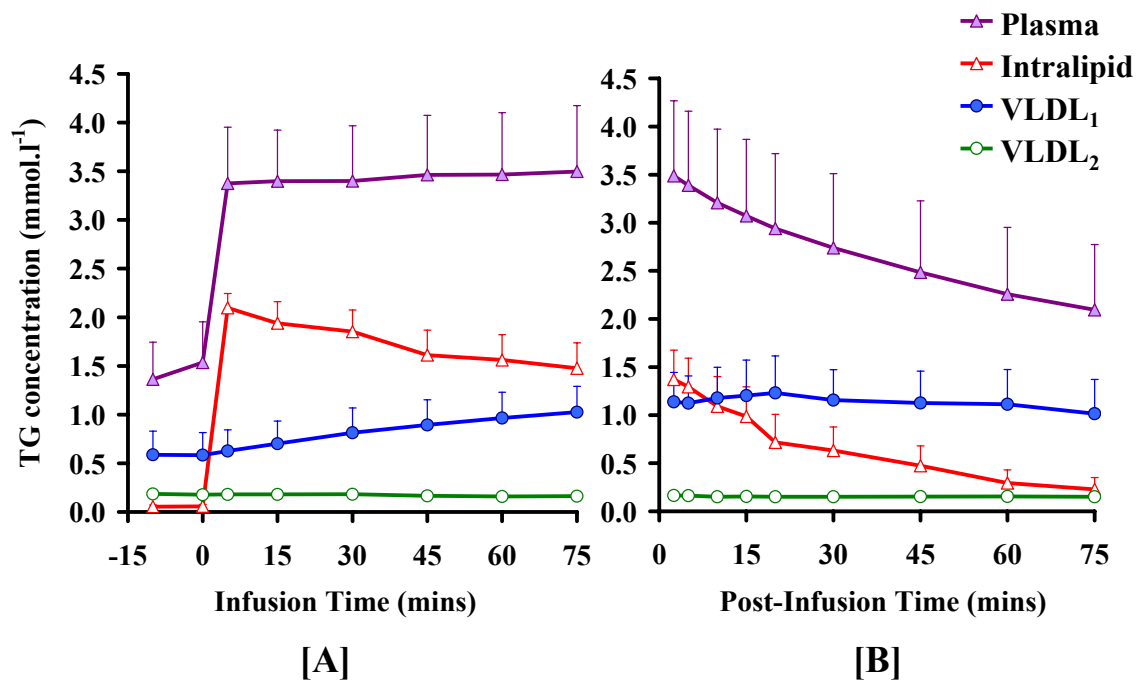
Statistical analyses were performed using MINITAB for Windows (Version 13.1, MINITAB Inc., State College, PA) and STATISTICA (Release 6.0, StatSoft, Inc, USA). Normality was checked for all the data using the Anderson-Darling test. When data did not approximate a normal distribution, these were log-transformed, specifically TG, glucose, insulin, HOMA<sub>IR</sub>, production rates of VLDL<sub>1</sub>-TG (expressed in both mg.h<sup>-1</sup> and mg.kg<sup>-1</sup>.d<sup>-1</sup>) and VLDL<sub>1</sub>-apoB (expressed in mg.h<sup>-1</sup>), Intralipid-TG clearance rate and VLDL<sub>1</sub>-apoB FSR required transformation. Time trends were tested using one-way ANOVA with repeated measures. Paired *t*-tests were used to compare between the Intralipid-TG clearance rates calculated from the steady state and the exponential decay and between the kinetic data obtained from the low and high Intralipid doses. HOMA [calculated as insulin (mU.l<sup>-1</sup>) × glucose (mmol.l<sup>-1</sup>)/22.5] was used as a validated surrogate measure of insulin resistance (Matthews *et al.*, 1985). Relationships between HOMA-estimated insulin resistance (HOMA<sub>IR</sub>), NEFA, BMI, waist circumference and kinetic parameters were assessed using Pearson product-moment correlations. Significance was accepted at the *P* < 0.05 level. Data are presented as mean ± SEM unless otherwise stated. It was not possible to perform a power calculation for this study as the method was still in the developmental stage and thus no data on the reliability of the method were available at this stage.

## 3.3 Results

### 3.3.1 Plasma, Intralipid, VLDL<sub>1</sub>- and VLDL<sub>2</sub>-TG Concentrations During and Post-Infusion

**Figure 3.5A** shows the mean plasma-, Intralipid-, VLDL<sub>1</sub>- and VLDL<sub>2</sub>-TG responses in 10 subjects during 75-min Intralipid infusion (0.1 g.h<sup>-1</sup>.kg<sup>-1</sup> body mass). Plasma-TG concentrations were increased to ~2-3 times the fasting value during the infusion. Similarly, mean Intralipid-TG concentrations increased in response to the bolus dose (*P* < 0.001). VLDL<sub>1</sub>-TG concentrations rose linearly during the infusion (*P* < 0.001), but VLDL<sub>2</sub>-TG did not change significantly during the course of the infusion (*P* = 0.14).

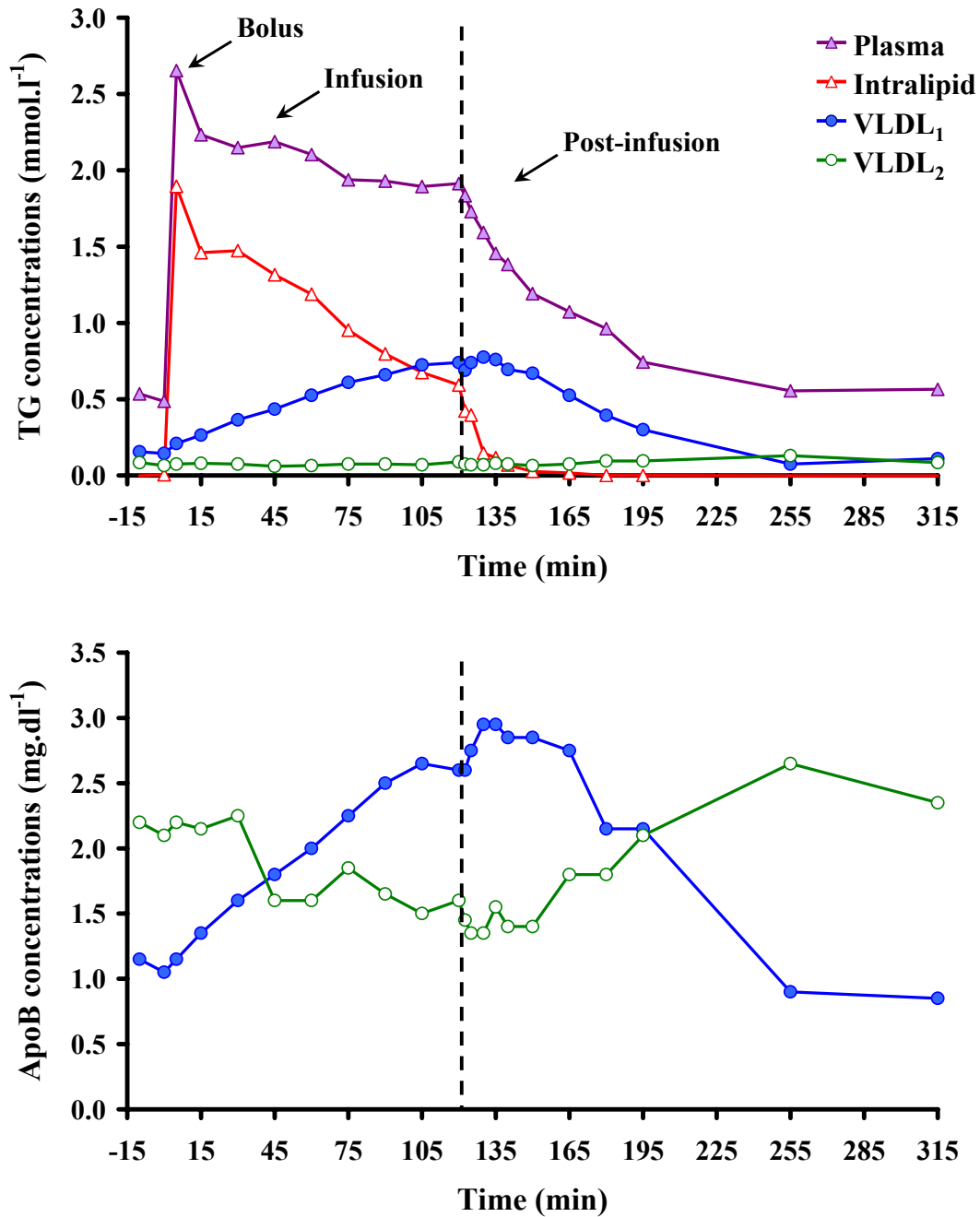
After stopping the infusion, the plasma- and Intralipid-TG concentrations decreased exponentially (both  $P < 0.001$ , **Figure 3.5B**). VLDL<sub>1</sub>-TG continued to rise for ~20 min ( $P < 0.001$ ) before plateauing and subsequently decreasing. In subjects where the post-infusion period was extended, VLDL<sub>1</sub>-TG returned to baseline concentrations within 105-135 min (an example of an earlier Intralipid trial is shown in **Figure 3.6**). The mean VLDL<sub>2</sub>-TG concentrations remained unchanged for the 75 min post-infusion observation period.



**Figure 3.5:** Plasma-, Intralipid-, VLDL<sub>1</sub>- and VLDL<sub>2</sub>-TG concentrations (mmol.l<sup>-1</sup>) [A] during infusion of 0.1 g.kg<sup>-1</sup>.h<sup>-1</sup> of 10% Intralipid and [B] for 75 min post-infusion. An initial bolus dose of 20% Intralipid (0.1 g.kg<sup>-1</sup> body mass) was given at 0 min. N = 10, values are mean  $\pm$  SEM.

### 3.3.2 VLDL<sub>1</sub>- and VLDL<sub>2</sub>-ApoB Concentrations During Infusion

The mean apoB concentration in the VLDL<sub>1</sub> (S<sub>f</sub> 60-400) fraction increased steadily from fasting levels throughout the infusion and was significantly higher than baseline within 15 min ( $P < 0.001$ ). On the other hand, mean VLDL<sub>2</sub>-apoB concentrations declined significantly ( $P < 0.001$ ) during 75 min of infusion. Total (VLDL<sub>1</sub> + VLDL<sub>2</sub>) VLDL-apoB concentrations rose slightly but significantly ( $P < 0.05$ ) during infusion (**Figure 3.7**). There was no significant change in the VLDL<sub>1</sub> or VLDL<sub>2</sub> TG/apoB ratio (expressed in mol:mol) over the 75 min of infusion ( $P = 0.21$  and  $P = 0.16$ , respectively) (**Figure 3.8**).



**Figure 3.6: An example of a full 135min-Intralipid trial during development of the method.** [Top] Plasma-, Intralipid-, VLDL<sub>1</sub>- and VLDL<sub>2</sub>-TG and [Bottom] VLDL<sub>1</sub>- and VLDL<sub>2</sub>-apoB concentrations during and post-infusion for a female subject (34 y, 55.2 kg). A bolus dose of 20% Intralipid (0.1 g.kg<sup>-1</sup> body mass) was injected at zero min, followed immediately by a constant infusion of 10% Intralipid (0.1 g.kg<sup>-1</sup>.h<sup>-1</sup>) for 120 min. The dotted line represents the end of infusion. VLDL<sub>1</sub> kinetics for this subject were calculated from the first 75 min of infusion: VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB production rates were 711.2 and 20.7 mg.h<sup>-1</sup> and fractional catabolic rates were 56.3 and 20.0 pools.d<sup>-1</sup>, respectively. Intralipid-TG clearance rate was calculated only from its exponential decay post-infusion as 146.9 pools.d<sup>-1</sup>, as no steady state was achieved during infusion.

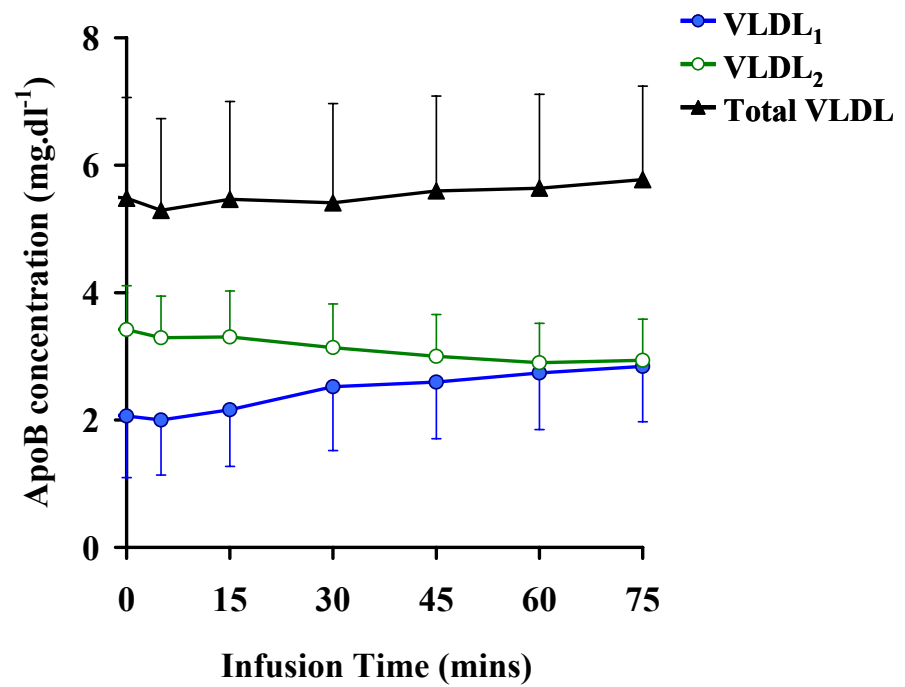


Figure 3.7: VLDL<sub>1</sub>-, VLDL<sub>2</sub>- and total VLDL-apoB concentrations (mg.dl<sup>-1</sup>) during infusion of 10% Intralipid (0.1 g.kg<sup>-1</sup>.h<sup>-1</sup>). An initial bolus dose of 20% Intralipid (0.1 g.kg<sup>-1</sup> body mass) was given at 0 min. N = 10, values are mean ± SEM.

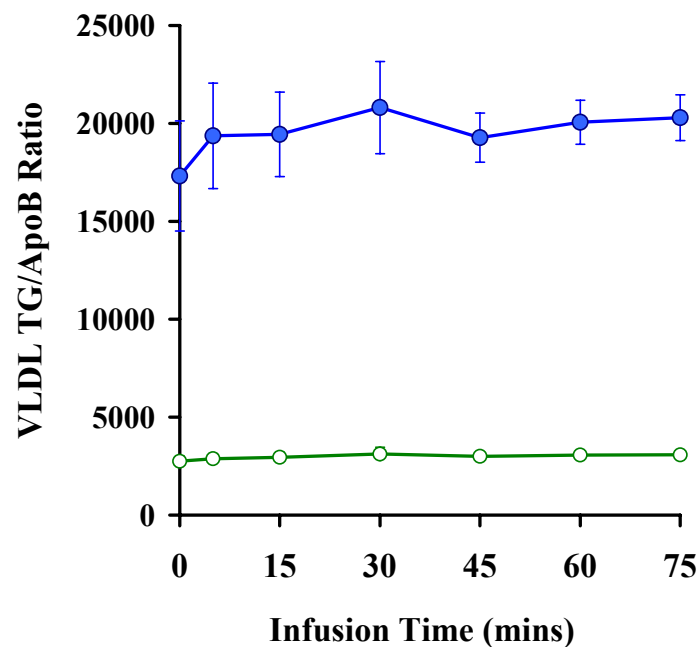


Figure 3.8: VLDL<sub>1</sub>- and VLDL<sub>2</sub>-TG/apoB ratios during infusion of 10% Intralipid (0.1 g.kg<sup>-1</sup>.h<sup>-1</sup>). An initial bolus dose of 20% Intralipid (0.1 g.kg<sup>-1</sup> body mass) was given at 0 min. N = 10, values are mean ± SEM.

### 3.3.3 Kinetic Data

**Table 3.2** shows the production rates and FSR of VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB as well as the clearance rates of Intralipid-TG calculated for each subject (n = 10) as previously described. Fasting VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>- and VLDL<sub>2</sub>-apoB concentrations are also presented.

#### 3.3.3.1 VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-ApoB Production Rates

The mean  $\pm$  SEM (range) production rates for VLDL<sub>1</sub>-TG and -apoB were  $1076.7 \pm 224.7$  (446.0 – 2563.2) mg.h<sup>-1</sup> and  $25.4 \pm 3.9$  (12.0 – 50.9) mg.h<sup>-1</sup>, respectively. These corresponded to  $333.6 \pm 49.1$  (198.5 – 609.9) mg.kg<sup>-1</sup>.d<sup>-1</sup> and  $8.1 \pm 0.9$  (4.4 – 14.5) mg.kg<sup>-1</sup>.d<sup>-1</sup>, respectively.

#### 3.3.3.2 VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-ApoB Fractional Synthetic and Catabolic Rates

The mean  $\pm$  SEM VLDL<sub>1</sub>-TG and -apoB FSRs, which are equal to the VLDL<sub>1</sub>-TG and -apoB FCRs in the fasted state (Foster *et al.*, 1993), were  $30.2 \pm 5.7$  (6.5 – 57.8) pools.d<sup>-1</sup> and  $21.1 \pm 5.1$  (2.2 – 56.2) pools.d<sup>-1</sup>, respectively.

#### 3.3.3.3 Intralipid-TG Clearance Rate

The Intralipid-TG clearance rates calculated for individual subjects by two methods as described above (i.e. SS and Exp) are shown in **Table 3.2**. Eight out of the 10 subjects reached the defined steady state during infusion. The mean  $\pm$  SEM Intralipid-TG clearance rates in these 8 subjects did not differ significantly between the two calculation methods ( $52.4 \pm 8.6$  pools.d<sup>-1</sup> for SS vs  $55.3 \pm 9.2$  pools.d<sup>-1</sup> for Exp,  $P = 0.45$ ) and the values obtained were strongly correlated ( $r = 0.96$ ,  $P < 0.001$ ). However, since not all subjects reached a steady state, the Intralipid-TG clearance rates mentioned hereafter will refer to that calculated using the exponential method.

**Table 3.2: Fasting concentrations, apoE phenotypes and individual lipoprotein kinetic parameters calculated using the ‘Intralipid method’ (0.1 g.kg<sup>-1</sup>.h<sup>-1</sup> infusion dose) (n=10).**

Subjects	Fasting Concentrations			Production Rates				Intralipid-TG Clearance Rate		FSR <sup>§</sup>	
(apoE phenotype)	VLDL <sub>1</sub> -TG	VLDL <sub>1</sub> -ApoB	VLDL <sub>2</sub> -ApoB	VLDL <sub>1</sub> -TG		VLDL <sub>1</sub> -ApoB		SS	Exp	VLDL <sub>1</sub> -TG	VLDL <sub>1</sub> -ApoB
	<i>mmol.l<sup>-1</sup></i>	<i>mg.dl<sup>-1</sup></i>	<i>mg.dl<sup>-1</sup></i>	<i>mg.h<sup>-1</sup></i>	<i>mg.kg<sup>-1</sup>.d<sup>-1</sup></i>	<i>mg.h<sup>-1</sup></i>	<i>mg.kg<sup>-1</sup>.d<sup>-1</sup></i>	<i>pools.d<sup>-1</sup></i>	<i>pools.d<sup>-1</sup></i>	<i>pools.d<sup>-1</sup></i>	<i>pools.d<sup>-1</sup></i>
<b>1</b> (3/3)	0.76	0.8	1.6	625.0	241.1	17.1	6.6	**	79.8	30.2	22.0
<b>2</b> (3/2)	0.83	1.1	2.9	719.9	214.6	14.8	4.4	82.6	66.1	25.8	10.3
<b>3</b> (3/2)	1.81	1.8	6.8	2147.4	609.9	50.9	14.5	56.9	56.4	30.9	19.8
<b>4</b> (4/3)	0.40	0.8	0.9	642.3	235.3	12.0	4.4	72.8	96.5	57.8	13.7
<b>5</b> (3/3)	0.54	1.1	2.2	711.2	309.2	20.7	9.0	**	146.9	56.3	20.0
<b>6</b> (3/3)	4.43	10.6	7.7	2563.2	593.2	40.2	9.3	17.3	18.1	6.5	2.2
<b>7</b> (3/3)	2.15	3.8	3.7	1228.1	406.5	24.9	8.2	39.8	40.5	9.3	5.4
<b>8</b> (3/3)	0.65	0.3	3.0	446.0	205.8	16.1	7.4	78.7	80.6	44.7	56.2
<b>9</b> (3/3)	1.04	1.0	3.0	885.5	322.0	24.1	8.8	43.4	54.3	22.6	21.9
<b>10</b> (3/3)	1.03	0.5	2.9	798.2	198.5	33.0	8.2	27.2	29.5	17.4	39.1

**Exp**, exponential decay; **FCR**, fractional catabolic rate; **FSR**, fractional synthetic rate; **SS**, steady state.

Subjects 3, 5 and 8 were female subjects.

\*\* Subject did not reach a steady state.

§ As FSR equals FCR in under steady state conditions, the VLDL<sub>1</sub>-TG and -apoB FSR values are equal to the FCR values in the fasted state.

### 3.3.4 Effect of Increasing the Intralipid Infusion Rate

For the 5 subjects who underwent Intralipid infusion at the Low ( $0.1 \text{ g.kg}^{-1}.\text{h}^{-1}$ ) and the High ( $0.2 \text{ g.kg}^{-1}.\text{h}^{-1}$ ) doses, there were no significant differences in the mean VLDL<sub>1</sub>-apoB production rates (Low dose:  $23.8 \pm 2.8$  vs High dose:  $22.0 \pm 1.9 \text{ mg.h}^{-1}$ ,  $P = 0.21$ ) or VLDL<sub>1</sub>-TG production rates (Low dose:  $813.8 \pm 127.0$  vs High dose:  $960.9 \pm 136.8 \text{ mg.h}^{-1}$ ,  $P = 0.10$ ) between both doses, although there was a tendency for the VLDL<sub>1</sub>-TG production rate to be higher at the High Intralipid dose. However, it was observed that separation of the large amount of Intralipid from plasma at the higher  $0.2 \text{ g.kg}^{-1}.\text{h}^{-1}$  dose was technically quite difficult and it was suspected that the VLDL<sub>1</sub> fraction in some samples may have become slightly contaminated with Intralipid at this dose (**Figure 3.9**). This suggestion is supported by the substantially lower recovery of Intralipid in the Intralipid fraction at high Intralipid concentrations (see below). Calculated FSRs for VLDL<sub>1</sub>-TG (Low dose:  $30.1 \pm 8.8$  vs High dose:  $30.4 \pm 8.1 \text{ pools.d}^{-1}$ ,  $P = 0.94$ ) and VLDL<sub>1</sub>-apoB (Low dose:  $28.5 \pm 8.7$  vs High dose:  $33.4 \pm 14.6 \text{ pools.d}^{-1}$ ,  $P = 0.64$ ) did not differ between the two doses. In addition, FSRs for VLDL<sub>1</sub>-TG ( $r = 0.88$ ,  $P = 0.05$ ) and VLDL<sub>1</sub>-apoB ( $r = 0.95$ ,  $P = 0.01$ ) between the two doses correlated highly with each other and, when plotted, followed the line-of-equality (**Figure 3.10**).

Furthermore, although a formal reproducibility test was not performed, the repeated two doses provided an opportunity to estimate the test/re-test reproducibility of VLDL<sub>1</sub>-TG and -apoB production rates. Based on data from these 5 subjects, the within-subject CV for VLDL<sub>1</sub>-TG production was 20.1% and for VLDL<sub>1</sub>-apoB production was 12.7%.

### 3.3.5 Intralipid Recovery

Recovery of the Intralipid-TG in the Intralipid ( $S_f > 400$ ) fraction was  $95 \pm 7\%$  (mean  $\pm$  SD) for the  $1.5 \text{ mmol.l}^{-1}$  Intralipid-TG concentrations and  $71 \pm 4\%$  for the  $\sim 4.0 \text{ mmol.l}^{-1}$  Intralipid-TG concentrations.



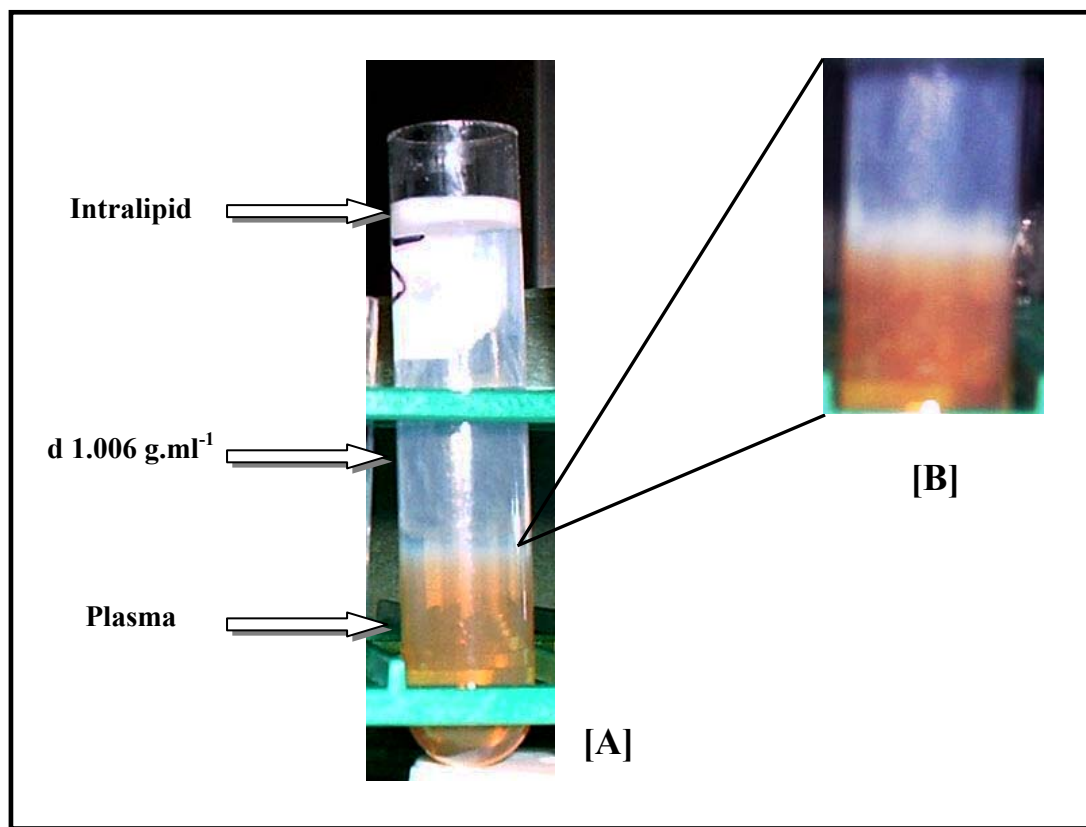


Figure 3.9: [A] An example of Intralipid separation in the High dose of Intralipid infusion (0.2 g.kg<sup>-1</sup>.h<sup>-1</sup>). Intralipid separation is done by overlaying 2 ml of plasma with 4 ml of density solution 1.006 g.ml<sup>-1</sup> and centrifuged at 10,000 rpm for 30 min. [B] A magnification of the bottom of the tube showing a thin layer of possible contamination of Intralipid at the top of plasma due to the high concentration of Intralipid.

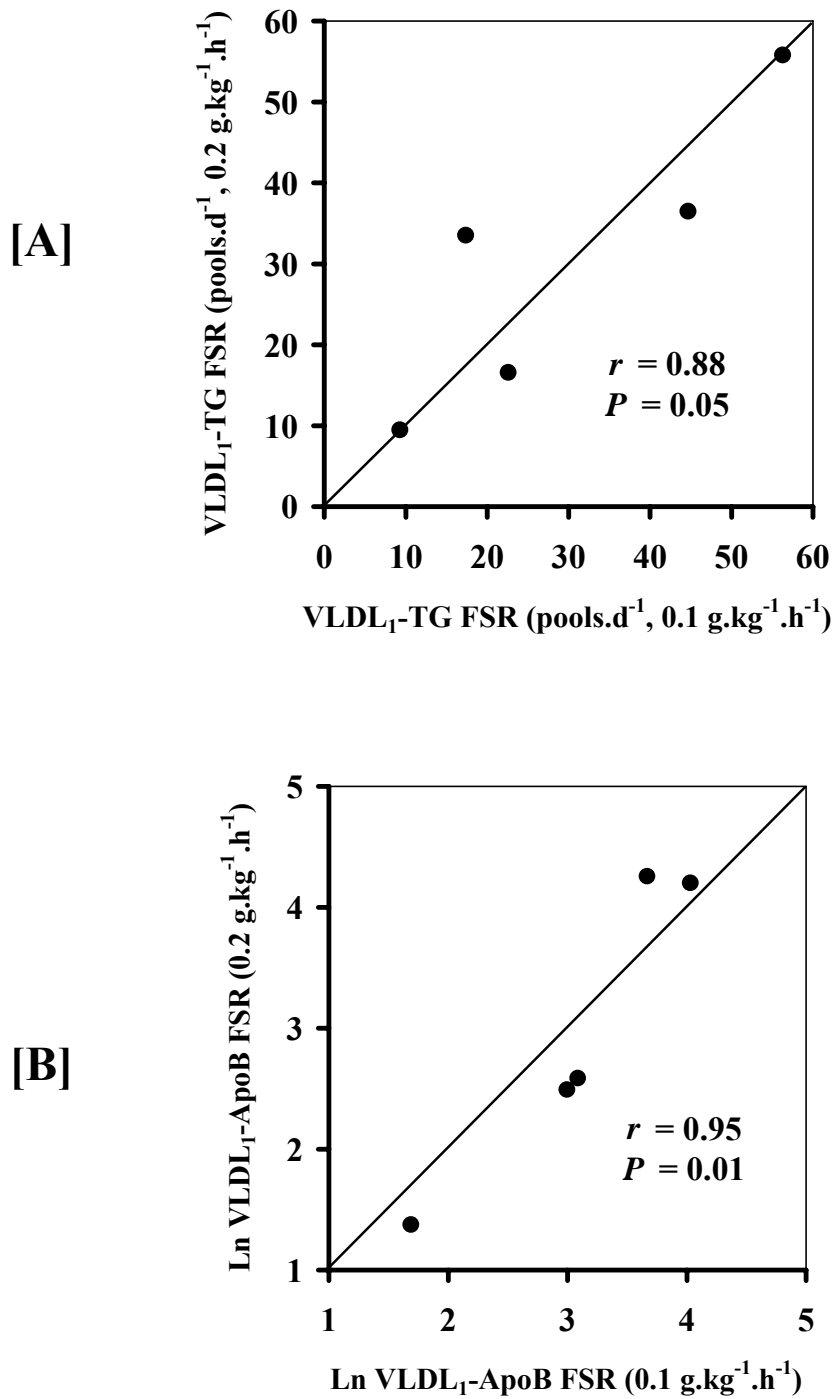


Figure 3.10: Scatterplots (with the line of equality) illustrating the agreement between [A] VLDL<sub>1</sub>-TG FSR (pools.d<sup>-1</sup>) and [B] Ln VLDL<sub>1</sub>-apoB FSR between the Low (0.1 g.kg<sup>-1</sup>.h<sup>-1</sup>) and High (0.2 g.kg<sup>-1</sup>.h<sup>-1</sup>) doses. FSR is equivalent to FCR in the fasted state. N = 5,  $r$  and  $P$  values are for Pearson product-moment correlations between variables. VLDL<sub>1</sub>-apoB FSR was expressed in pools.d<sup>-1</sup> prior to log-transformation.

### 3.3.6 Interrelationships between VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB Production and TG Clearance

Intralipid-TG clearance rate and VLDL<sub>1</sub>-TG and -apoB production rates (expressed in mg.h<sup>-1</sup>) were significantly inter-related (**Figure 3.11**) with the expected negative correlation between Intralipid-TG clearance and VLDL<sub>1</sub>-TG ( $r = -0.67$ ,  $P = 0.04$ ) and VLDL<sub>1</sub>-apoB ( $r = -0.69$ ,  $P = 0.03$ ) production rates and a positive correlation between VLDL<sub>1</sub>-TG and -apoB production rates ( $r = 0.85$ ,  $P = 0.002$ ). There was also a very strong relationship between VLDL<sub>1</sub>-TG FSR (which equals the VLDL<sub>1</sub>-TG FCR in the fasted state) and Intralipid-TG clearance rate ( $r = 0.90$ ,  $P < 0.005$ ). The positive correlation between VLDL<sub>1</sub>-TG and -apoB production rates remained significant between production rates when values were expressed in mg.kg<sup>-1</sup>.d<sup>-1</sup> ( $r = 0.73$ ,  $P = 0.02$ ). However, the relationships between VLDL<sub>1</sub>-TG production rate expressed in mg.kg<sup>-1</sup>.d<sup>-1</sup> and Intralipid-TG clearance ( $r = -0.46$ ,  $P = 0.18$ ) and between VLDL<sub>1</sub>-apoB production rate expressed in mg.kg<sup>-1</sup>.d<sup>-1</sup> and Intralipid-TG clearance ( $r = -0.28$ ,  $P = 0.44$ ) were not statistically significant.

### 3.3.7 Relationships between Kinetic Variables and Subjects Characteristics

**Figure 3.12** shows the relationships between the calculated kinetic variables and subjects characteristics, with VLDL<sub>1</sub>-TG and -apoB production rates expressed in mg.h<sup>-1</sup>. VLDL<sub>1</sub>-TG and -apoB production rates (mg.h<sup>-1</sup>) correlated strongly and significantly with waist circumference and fasting TG concentration. VLDL<sub>1</sub>-TG production rate also correlated significantly with HOMA<sub>IR</sub>. Similarly, Intralipid-TG clearance rate was significantly and inversely correlated with waist circumference, fasting TG concentrations and HOMA<sub>IR</sub>. In addition, BMI correlated significantly and positively with VLDL<sub>1</sub>-TG ( $r = 0.83$ ,  $P = 0.003$ ) and VLDL<sub>1</sub>-apoB ( $r = 0.81$ ,  $P = 0.004$ ) production rates and inversely with Intralipid-TG clearance rate ( $r = -0.60$ ,  $P = 0.07$ ). Fasting NEFA concentrations were not significantly correlated with any of the kinetic variables. The relationships between VLDL<sub>1</sub>-TG and -apoB production rates expressed in mg.kg<sup>-1</sup>.d<sup>-1</sup> units with BMI, waist circumference, fasting TG concentration and HOMA<sub>IR</sub> are shown in **Table 3.3**. The correlations between VLDL<sub>1</sub>-TG production and all of these variables remained strong and statistically significant, however the correlations between VLDL<sub>1</sub>-apoB production rate and

waist circumference and fasting TG were not statistically significant when the production rates were normalized for body mass.

### 3.3.8 Relationships between Kinetic Variables and ALT Concentrations

**Figure 3.13** shows the relationships between serum ALT concentrations and kinetic variables. Although all subjects had ALT concentrations within the normal range, one subject was on antibiotics shortly prior to participating in the study and was excluded from the following correlations. ALT concentrations correlated significantly with VLDL<sub>1</sub>-apoB production rate ( $r = 0.73$ ,  $P = 0.03$ ) and Intralipid-TG clearance rate ( $r = -0.79$ ,  $P = 0.01$ ). This indicates that, within normal range, ALT concentrations explained 53% ( $0.73^2$ ) and 62% ( $0.79^2$ ) of the variance in VLDL<sub>1</sub>-apoB production rate and Intralipid-TG clearance rate, respectively. There was a tendency toward a positive correlation with VLDL<sub>1</sub>-TG production rate, but it did not reach statistical significance ( $r = 0.59$ ,  $P = 0.09$ ).

**Table 3.3: Correlations between VLDL<sub>1</sub>-apoB and -TG production rates (expressed in mg.kg<sup>-1</sup>.d<sup>-1</sup>) and subjects' characteristics.**

	VLDL <sub>1</sub> -ApoB Production Rate	Ln VLDL <sub>1</sub> -TG Production Rate
<b>BMI</b>	$r = 0.63$ $P = 0.05$	$r = 0.65$ $P = 0.04$
<b>WC</b>	$r = 0.45$ $P = 0.19$	$r = 0.70$ $P = 0.02$
<b>Ln Fasting TG</b>	$r = 0.54$ $P = 0.11$	$r = 0.79$ $P = 0.01$
<b>Ln HOMA<sub>IR</sub></b>	$r = 0.40$ $P = 0.25$	$r = 0.79$ $P = 0.01$

**BMI**, body mass index; **HOMA<sub>IR</sub>**, HOMA-estimated insulin resistance; **WC**, waist circumference. N = 10,  $r$  and  $p$  values are for Pearson product-moment correlations between values.

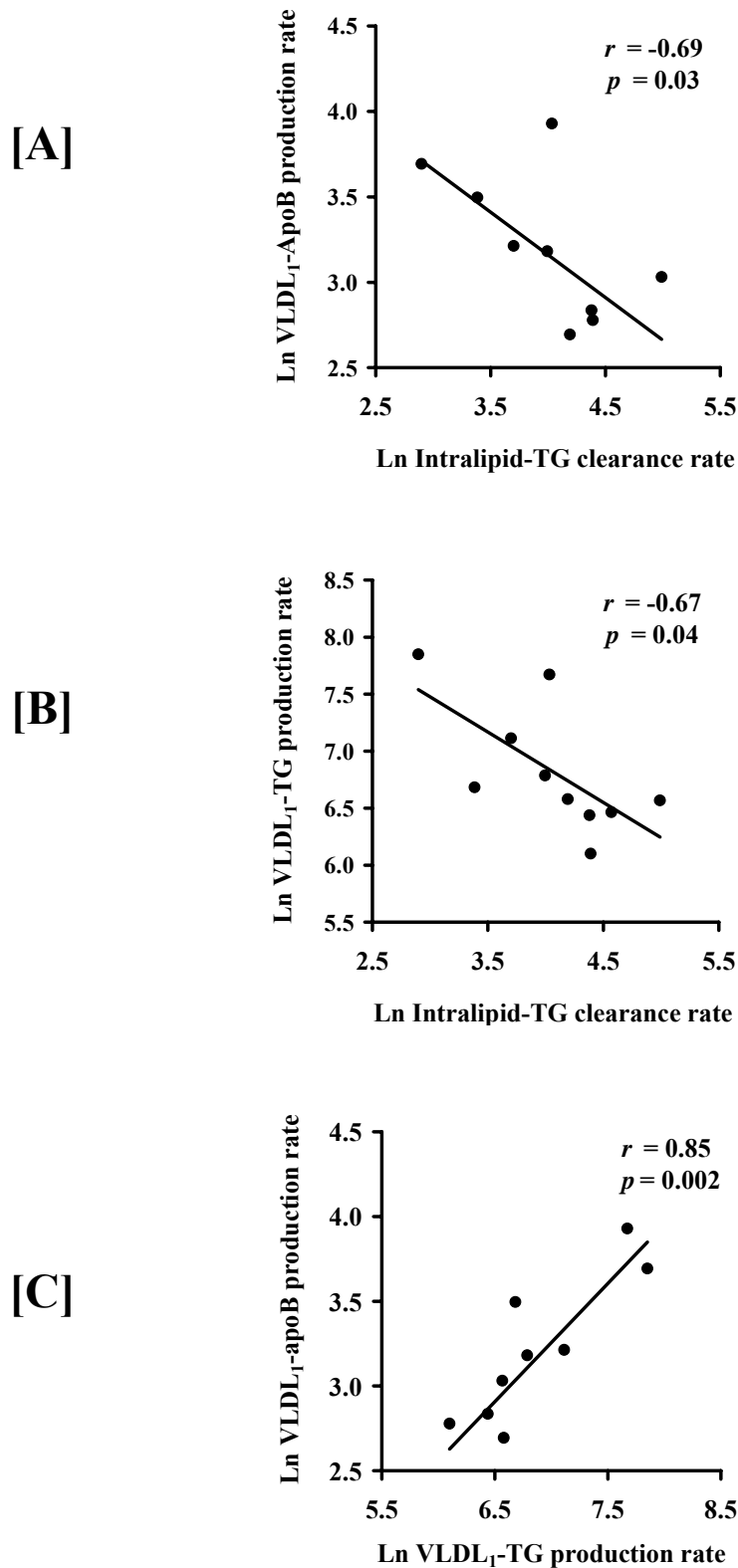
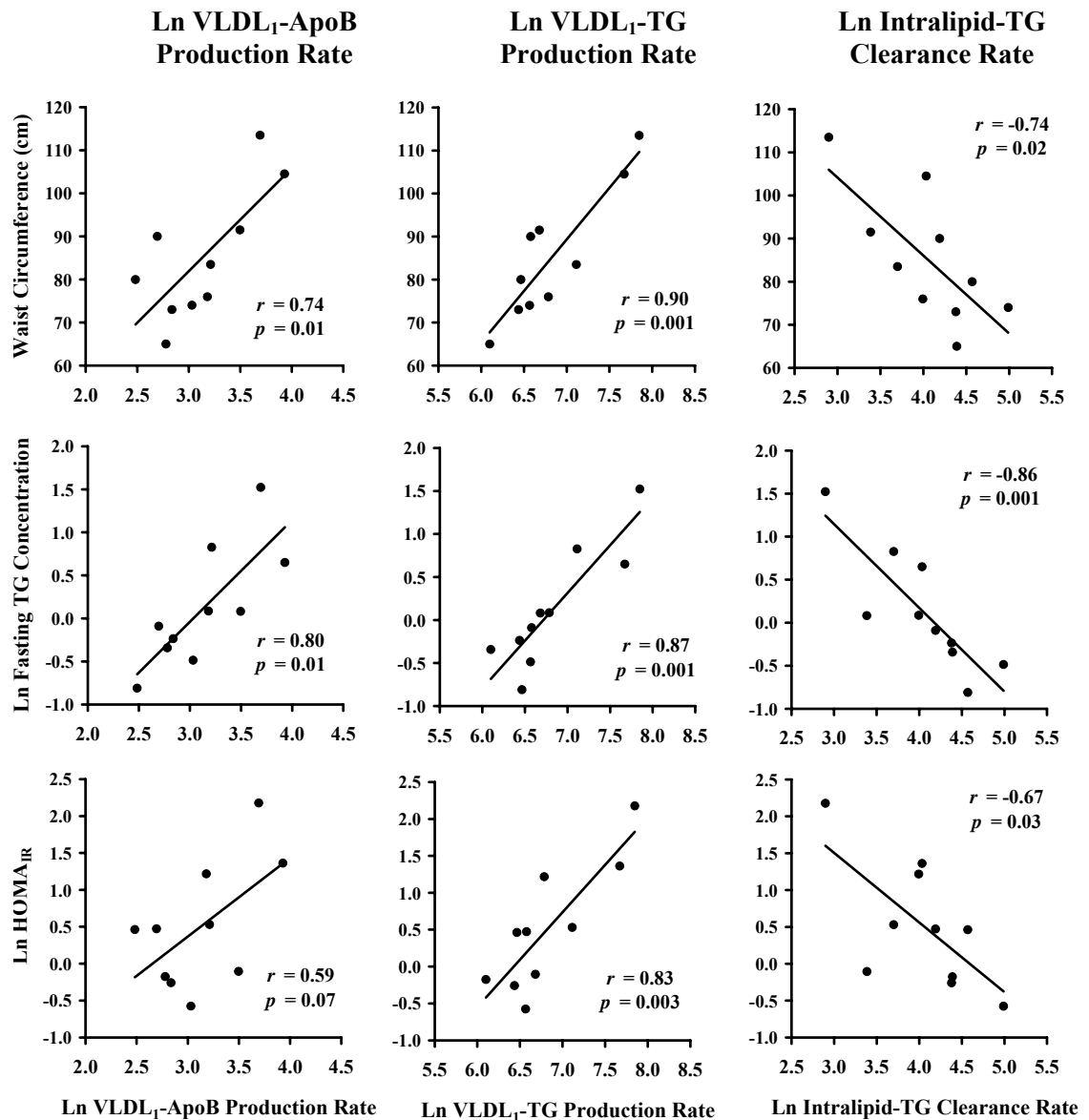
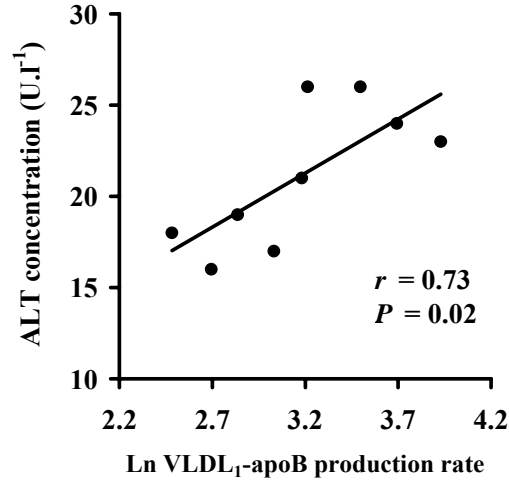


Figure 3.11: Scatterplots (with linear-regression lines of ‘best-fit’) illustrating the interrelationships between [A] Ln Intralipid-TG clearance rate and Ln VLDL<sub>1</sub>-apoB production rate, [B] Ln Intralipid-TG clearance rate and Ln VLDL<sub>1</sub>-TG production rate, and [C] Ln VLDL<sub>1</sub>-apoB and Ln VLDL<sub>1</sub>-TG production rates. N = 10,  $r$  and  $P$  values are for Pearson product-moment correlations between variables. Production rates and Intralipid-TG clearance rate were expressed in  $\text{mg.h}^{-1}$  and  $\text{pools.d}^{-1}$  prior to log-transformation, respectively.

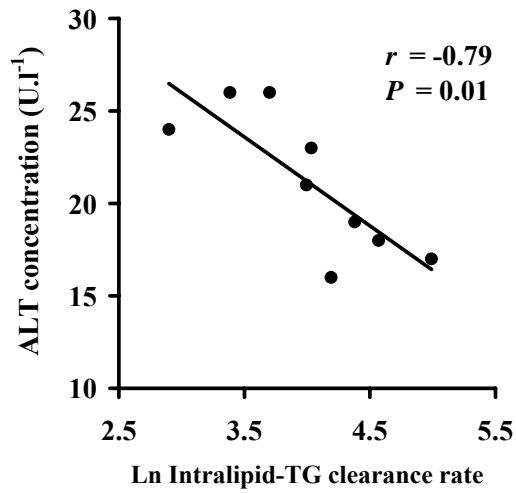


**Figure 3.12:** Scatterplots (with linear-regression lines of ‘best-fit’) illustrating the relationships between the kinetic variables [Ln VLDL<sub>1</sub>-TG production rate (left), Ln VLDL<sub>1</sub>-apoB production rate (middle) and Ln Intralipid-TG clearance rate (right)] and subjects’ characteristics: waist circumference (top), Ln fasting TG concentrations (middle) and Ln HOMA-estimated insulin resistance (HOMA<sub>IR</sub>) (bottom). N = 10,  $r$  and  $p$  values are for Pearson product-moment correlations between variables.

[A]



[B]



[C]

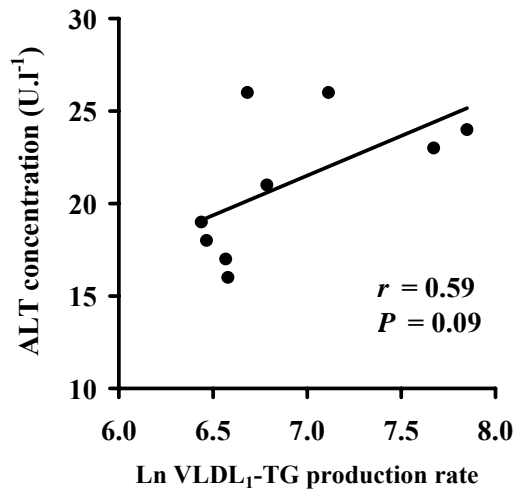


Figure 3.13: Scatterplots (with linear-regression lines of ‘best-fit’) illustrating the relationships between serum ALT concentration (mU.l<sup>-1</sup>) and [A] Ln VLDL<sub>1</sub>-apoB production rate, [B] Ln Intralipid-TG clearance rate and [C] Ln VLDL<sub>1</sub>-apoB production rate. N = 9,  $r$  and  $P$  values are for Pearson product-moment correlations between variables.

### 3.4 Discussion

In the present study, a relatively straightforward method of determining TRL kinetics was developed. The method relies on the fact that chylomicrons or chylomicron-like particles, such as Intralipid, compete with hepatically-derived large VLDL<sub>1</sub> particles for clearance by a common saturable pathway – i.e. hydrolysis of their TG content by LPL (Björkegren *et al.*, 1996; Karpe & Hultin, 1995) – and that chylomicrons or chylomicron-like particles are the preferred substrate for LPL (Fisher *et al.*, 1995). Thus, the presence of a sufficient concentration of chylomicrons or chylomicron-like particles in the circulation will almost entirely prevent clearance of VLDL<sub>1</sub> by LPL (Björkegren *et al.*, 1996) and the rates of VLDL<sub>1</sub>-TG and -apoB production can therefore be calculated from their rates of rise in concentration. The present work builds on the findings of Björkegren *et al.* who, in studies designed to evaluate the effects of Intralipid infusion on VLDL<sub>1</sub> (S<sub>f</sub> 60-400) and VLDL<sub>2</sub> (S<sub>f</sub> 20-60) kinetics, found that individual rates of VLDL<sub>1</sub>-apoB production calculated from the rate of rise of VLDL<sub>1</sub>-apoB during infusion were virtually identical to those calculated from the ‘gold-standard’ stable isotope method (see below) (Björkegren *et al.*, 1996). The ‘Intralipid method’ described in the present study enables the determination of the rate of VLDL<sub>1</sub>-TG (i.e. VLDL<sub>1</sub> lipid) and VLDL<sub>1</sub>-apoB (i.e. VLDL<sub>1</sub> particle) production as well as the clearance rate of chylomicron-like particles.

The Intralipid method specifically measures the production rate of large VLDL<sub>1</sub> (S<sub>f</sub> 60-400) rather than total VLDL (i.e. S<sub>f</sub> 20-400). VLDL is a metabolically heterogeneous class of lipoproteins, and it is the larger VLDL<sub>1</sub> subclass which competes with chylomicrons/chylomicron-like particles for LPL-mediated clearance and would have its clearance blocked by the presence of Intralipid (Björkegren *et al.*, 1996). In contrast, catabolism of the smaller VLDL<sub>2</sub> subclass would not be completely blocked by Intralipid as its clearance can occur via the action of hepatic lipase as well as LPL (Packard & Shepherd, 1997). Indeed, as one source of VLDL<sub>2</sub> is from catabolism of VLDL<sub>1</sub> (the other being direct hepatic production), and this process was blocked by Intralipid infusion, mean VLDL<sub>2</sub>-apoB concentrations decreased slightly during the infusion although individual responses were more heterogeneous than that observed in VLDL<sub>1</sub>-apoB; a finding also reported by Björkegren *et al.* (Björkegren *et al.*, 1996). This heterogeneity in individual VLDL<sub>2</sub>-



apoB responses meant that it was not possible to perform any kinetic analyses using the VLDL<sub>2</sub> data.

To validate the calculation of VLDL<sub>1</sub>-TG and -apoB production rates using the 'Intralipid method' it was necessary to consider a number of issues. The first was to determine whether infusing a higher Intralipid dose would influence the calculated VLDL<sub>1</sub>-TG and -apoB production rates. This was necessary to establish whether the proposed Intralipid infusion dose of 0.1 g.kg<sup>-1</sup>.h<sup>-1</sup>, was sufficient to saturate LPL and block clearance of VLDL<sub>1</sub>: if the 0.1 g.kg<sup>-1</sup>.h<sup>-1</sup> dose was sufficient, infusing a higher Intralipid dose should not affect calculated production rates. In agreement with the findings of Björkegren *et al.* (Björkegren *et al.*, 1996), the calculated VLDL<sub>1</sub>-apoB production rate in the present study was not changed when a higher, 0.2 g.kg<sup>-1</sup>.h<sup>-1</sup> Intralipid infusion dose was used. Similarly, FSRs, which correspond to the FCRs in the fasted state, for VLDL<sub>1</sub>-TG and -apoB did not differ between the two doses (**Figure 3.10**). Although there was tendency for the calculated VLDL<sub>1</sub>-TG production rate to be higher with the 0.2 g.kg<sup>-1</sup>.h<sup>-1</sup> dose, this was not statistically significant. However, the author feels that the slightly higher apparent VLDL<sub>1</sub>-TG production rate at the High dose was a methodological, rather than a physiological, issue caused by the difficulty in separating Intralipid at the High dose, leading to the potential contamination of the VLDL<sub>1</sub> fraction with Intralipid. This is supported by the fact that Intralipid recoveries at high Intralipid doses were relatively low (71% at an Intralipid concentration of 4 mmol.l<sup>-1</sup>). This contrasts with the near complete recovery of Intralipid at lower Intralipid doses (95% at an Intralipid concentration of 1.5 mmol.l<sup>-1</sup>). This, of course, would not influence the VLDL<sub>1</sub>-apoB production rate calculations, as Intralipid particles do not contain apoB.

A further issue to consider is whether, following lipolysis by LPL, Intralipid 'remnant' particles may have appeared in the VLDL<sub>1</sub> fraction, thereby increasing the measured VLDL<sub>1</sub>-TG concentration and the apparent VLDL<sub>1</sub>-TG production rate. However, the author does not believe that this would have had a substantial effect on calculated VLDL<sub>1</sub>-TG production rates for a number of reasons. Firstly, evidence from the literature suggests that for large TG-rich particles, particularly chylomicron-like particles, lipolysis and particle removal from the plasma is likely to occur simultaneously, rather than by sequential mechanisms (Olivecrona & Olivecrona,

1998; Hultin *et al.*, 1996), with the majority of particles removed from the plasma before conversion to smaller VLDL-sized remnant particles (Karpe *et al.*, 1997). Secondly, as Intralipid contains TG but not apoB, appearance of Intralipid remnants in the VLDL<sub>1</sub> fraction would lead to a disproportionate rise in VLDL<sub>1</sub>-TG compared with VLDL<sub>1</sub>-apoB, leading to an increase in the VLDL<sub>1</sub> TG/apoB ratio. There was no significant increase in this ratio during the infusion ( $P = 0.21$ ). Thirdly, if the rise in VLDL<sub>1</sub>-TG was influenced by the appearance of Intralipid remnant particles, then a positive correlation between Intralipid clearance and VLDL<sub>1</sub>-TG production would be evident (i.e. increased Intralipid clearance would lead to increased VLDL<sub>1</sub>-TG production). Instead, a negative relationship between Intralipid clearance and VLDL<sub>1</sub>-TG production (expressed in  $\text{mg.h}^{-1}$ ) was observed (i.e. subjects with slow Intralipid clearance also had high VLDL<sub>1</sub>-TG production) ( $r = -0.67$ ,  $P = 0.04$ ). Furthermore, the relationship between VLDL<sub>1</sub>-apoB production, which would be unaffected by the presence of Intralipid remnant particles, and VLDL<sub>1</sub>-TG production was very strong, with 71% of the variance in the VLDL<sub>1</sub>-TG production rate explained by the VLDL<sub>1</sub>-apoB production rate ( $r = 0.85$ ,  $P = 0.002$ ).

Furthermore, it is important to ascertain whether the results obtained are comparable with data obtained using the ‘gold standard’ stable isotope tracer method. An internal validation of this method was previously undertaken by Björkegren and colleagues (Björkegren *et al.*, 1996) in 3 subjects. They reported VLDL<sub>1</sub>-apoB production rates of 20.0, 25.6 and 7.2  $\text{mg.h}^{-1}$  calculated from the Intralipid infusion method with corresponding rates calculated from a stable isotope method of 23.8, 21.6 and 8.0  $\text{mg.h}^{-1}$ , respectively, indicating that data obtained from the two methods were comparable. In addition, from **Table 3.4**, it is clear that the values for VLDL<sub>1</sub>-apoB production in the present study are of the same order as those obtained from a number of studies which determined VLDL<sub>1</sub>-apoB production using stable isotope techniques. Determination of VLDL<sub>1</sub>-TG production rates using stable isotope tracer methods is technically more difficult than determination of VLDL<sub>1</sub>-apoB production, and the author is only aware of one group of workers who have evaluated this (Adiels *et al.*, 2005a; Adiels *et al.*, 2006b). The values obtained for VLDL<sub>1</sub>-TG production in the present study are of the same order as those published by Adiels *et al.* (Adiels *et al.*, 2005a; Adiels *et al.*, 2006b).

This Intralipid method enabled the Intralipid-TG clearance rate to be calculated in two different ways: from the steady state concentration of Intralipid-TG during the infusion, which was defined in the present study as the mean of the final 3 values if these differed by less than 13.8% (i.e. two times the CV for the separation of the Intralipid fraction and measurement of the TG), and from the exponential decrease in Intralipid-TG post-infusion (Rossner, 1974). In subjects where a steady-state Intralipid-TG concentration was achieved, the Intralipid-TG clearance rates calculated from the steady state concentration and from the post-infusion exponential decrease agreed closely (see **Table 3.2**). However, not all subjects achieved a steady-state Intralipid-TG concentration in 75 min of infusion and it is not possible to determine whether a steady-state was achieved for a given subject until sample analysis was completed. Therefore, in practice, it may be easier to use the post-infusion values to determine Intralipid-TG clearance rates, as this ensures that the Intralipid infusion can be kept as short as is necessary to enable calculation of VLDL<sub>1</sub>-TG and -apoB production rates.

Moreover, the author sought to determine whether this Intralipid method revealed the physiologically expected differences in TRL kinetics between subjects with differing physical and metabolic profiles. As expected, there were strong positive correlations between fasting TG concentrations and VLDL<sub>1</sub>-TG production rates expressed in either absolute terms or normalized according to body mass and between fasting TG and VLDL<sub>1</sub>-apoB, production expressed in mg.h<sup>-1</sup> units, with a strong negative correlation between fasting TG and the Intralipid-TG clearance rates, indicating that those with high TG concentrations exhibited a combination of enhanced VLDL<sub>1</sub> production and diminished TG clearance. VLDL<sub>1</sub>-TG FCR in the fasted state (i.e. with no Intralipid present) was ~45% of the Intralipid-TG clearance rate ( $30.2 \pm 5.7$  pools.d<sup>-1</sup> vs.  $66.2 \pm 11.7$  pools.d<sup>-1</sup>, see **Table 3.2**) and there was a very strong correlation between the two variables ( $r = 0.90$ ,  $P < 0.0005$ ), indicating that clearance rates for VLDL<sub>1</sub> and chylomicron-like particles within an individual are very tightly linked, consistent with the fact that these particles are cleared by the same pathway. The expected positive correlations between indices of body fatness (waist circumference and BMI) and insulin resistance (HOMA<sub>IR</sub>) and VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB production rates were also observed, in agreement with findings previously reported using stable isotope tracer methods (Gill *et al.*, 2004a). In

addition, significant negative relationships were observed between Intralipid-TG clearance and HOMA<sub>IR</sub> and waist circumference. Thus, the Intralipid method appears to be sensitive enough to detect physiologically relevant differences in TRL kinetics between individuals across the normal and moderately hypertriglyceridaemic range.

Finally, liver fat has been reported to be independently associated with insulin (Westerbacka *et al.*, 2004) and predicted VLDL<sub>1</sub> production (Adiels *et al.*, 2006b). It was proposed that insulin resistance induces an increase in the fatty acid flux from adipose tissue to the liver, which leads to the accumulation of fat in the liver, resulting in overproduction of VLDL<sub>1</sub> particles (Adiels *et al.*, 2006b). As serum ALT concentrations, within normal range, have been shown to be a marker for liver fat (Westerbacka *et al.*, 2004) and has been shown to predict the onsets of type 2 diabetes (Sattar *et al.*, 2004; Sattar *et al.*, 2007), the metabolic syndrome (Schindhelm *et al.*, 2007b) and CVD events (Schindhelm *et al.*, 2007a), the author investigated whether serum ALT concentrations, would predict hepatic VLDL<sub>1</sub> production rate in the present study. Indeed, ALT concentrations correlated significantly with VLDL<sub>1</sub>-apoB production rate and Intralipid-TG clearance rate, explaining 53% and 62% of their variance; respectively. This suggests that ALT concentrations are significant predictors of hepatic VLDL<sub>1</sub> production and further validates the sensitivity of the ‘Intralipid Method’.

In conclusion, this chapter describes the development of a novel method to determine TRL kinetics. The ‘Intralipid method’ provides a relatively straightforward and cost-effective way of determining VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB production rates, and the clearance rate of chylomicron-like particles, which does not require specialised equipment or skill in multicompartmental modelling. The author believes that this method will increase the scope for the study of TRL kinetics, particularly in circumstances where issues related to funding or equipment availability preclude the use of more traditional isotopic tracer methods.

**Table 3.4: Comparison of values for VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB production rates (range) calculated in the present study (Intralipid Method) and in previously published studies using the stable-isotope method.**

Study	Subjects			Production rates			
	N (m/f)	BMI	TG	VLDL <sub>1</sub> -ApoB		VLDL <sub>1</sub> -TG	
		kg.m <sup>-2</sup>	mmol.l <sup>-1</sup>	mg.h <sup>-1</sup>	mg.kg <sup>-1</sup> .d <sup>-1</sup>	mg.h <sup>-1</sup>	mg.kg <sup>-1</sup> .d <sup>-1</sup>
(Bjorkegren <i>et al.</i> , 1996)	16 (m)	20.0 - 25.8	0.56 – 1.85	8.0 - 23.8			-
(Demant <i>et al.</i> , 1996)	6 (m)	-	1.00 – 2.40	21.2 - 51.8			-
(Pietzsch <i>et al.</i> , 1996)	6 (3/3)	20.5 - 25.0	0.70 – 1.46		22.9 - 50.7		-
(Gill <i>et al.</i> , 2004a)	16 (8/8)	19.6 - 32.9	1.00 – 3.15	8.5 - 67.8			-
(Adiels <i>et al.</i> , 2005a)	17	22.4 - 30.1	0.99 – 2.59		2.9 - 12.5		107 - 347
(Zheng <i>et al.</i> , 2006)	5 (f)	22.0 – 27.0	0.89 – 1.58		8.4 ± 5.6*		-
(Adiels <i>et al.</i> , 2006b)	18	22.0 – 30.0	0.67 – 3.14		2.9 - 12.5		107 - 352
<b>Intralipid Method</b>	<b>10 (7/3)</b>	<b>20.8 - 34.7</b>	<b>0.40 – 4.43</b>	<b>12.0 - 50.9</b>	<b>4.4 - 14.5</b>	<b>446.0 - 2563.2</b>	<b>199 - 610</b>

\*Data are mean ± SD.

## **4. Effects of Acute Hyperinsulinaemia and Hyperglycaemia on VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-ApoB Production and Intralipid-TG Clearance in Normoglycaemic Subjects**

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### **4.1 Introduction**

Diabetic dyslipidaemia is believed to be initiated by high concentrations of large VLDL<sub>1</sub> particles, which, consequently, cause the accumulation of remnant atherogenic particles and the generation of small dense LDL and small dense HDL (Taskinen, 2003). Interestingly, this profile of atherogenic lipoprotein abnormalities along with insulin resistance and hyperinsulinaemia is detectable in both the fasting (Tilly-Kiesi *et al.*, 1996) and postprandial (Johanson *et al.*, 2004) states years before the diagnosis of diabetes and appears to be related more to insulin resistance than hyperglycaemia. Therefore, it is important to understand the role of insulin in the regulation of hepatic VLDL<sub>1</sub> production.

Insulin plays a pivotal role in the regulation of hepatic VLDL production and acute hyperinsulinaemia has been shown to suppress VLDL production in normal subjects (Lewis *et al.*, 1993; Malmstrom *et al.*, 1997b). Insulin directly inhibits VLDL assembly and secretion in the liver through a number of mechanisms; e.g. it enhances intracellular apoB degradation (Sparks & Sparks, 1990) and inhibits the maturation process of VLDL assembly (Brown & Gibbons, 2001). In addition, it suppresses the NEFA flux to the liver, thereby reducing substrate availability for VLDL formation (Coppack *et al.*, 1994). However, these suppressive effects may be attenuated or lost in insulin resistant conditions. It has been reported that type 2 diabetic (Malmstrom *et al.*, 1997a; Bioletto *et al.*, 2000) and obese individuals (Lewis *et al.*, 1993; Bioletto *et al.*, 2000) fail to suppress hepatic VLDL production and concentrations (Bioletto *et al.*, 2000) to the same extent as their normal-weight, non-diabetic counterparts.

It is important to recognise that VLDL is a heterogeneous particle consisting of at least two major subclasses - large TG-rich VLDL<sub>1</sub> ( $S_f$  60-400) and small, denser VLDL<sub>2</sub> ( $S_f$  20-40) (Packard & Shepherd, 1997) - which have been shown to be independently regulated (Packard & Shepherd, 1997; Malmstrom *et al.*, 1998; Gill *et al.*, 2004a). Thus, the apparent effects of regulators on VLDL<sub>1</sub> metabolism may be masked or attenuated in studies examining the kinetics of total VLDL without subdivision into VLDL<sub>1</sub> and VLDL<sub>2</sub> subclasses. Indeed, VLDL<sub>1</sub>, but not VLDL<sub>2</sub>, particles have been shown to be the major determinant of plasma TG (Tan *et al.*, 1995) and the target for insulin-mediated VLDL suppression (Gill *et al.*, 2004a; Malmstrom *et al.*, 1998).

Furthermore, it is important to differentiate between VLDL-apoB and VLDL-TG production rates. As there is only one apoB molecule per VLDL particle (Elovson *et al.*, 1988), VLDL-apoB reflects particle number, whereas VLDL-TG is a measure of the major lipid component of VLDL particle. VLDL-apoB production rate may be dissociated from that of VLDL-TG (Kissebah *et al.*, 1981), giving rise to variable TG-to-apoB ratios and, hence, VLDL particle size.

To the best of the author's knowledge, studies examining the effects of acute hyperinsulinaemia on VLDL have either (1) addressed VLDL heterogeneity but considered only the production of VLDL<sub>1</sub>-apoB, but not VLDL<sub>1</sub>-TG (Malmstrom *et al.*, 1997b; Malmstrom *et al.*, 1998; Bioletto *et al.*, 2000; Annuzzi *et al.*, 2001); (2) investigated both apoB and TG production rates, but in total VLDL (Lewis *et al.*, 1993; Lewis *et al.*, 1994; Lewis & Steiner, 1996; Mittendorfer *et al.*, 2003b; Lewis *et al.*, 1995), rather than VLDL<sub>1</sub> specifically; or (3) studied the composition of VLDL subfractions rather than production rates (Annuzzi *et al.*, 2001; Bioletto *et al.*, 2000). However, at the start of the present study, no information existed on the acute effects of insulin on both VLDL<sub>1</sub>-TG and -apoB production rates in normoglycaemic subjects. Recently, a non-steady-state stable isotope study by Adiels *et al.* (Adiels *et al.*, 2007) was published, which investigated the acute effect of insulin on the production rates and pool sizes of VLDL<sub>1</sub>-TG and -apoB in relation to liver fat content in normal and diabetic subjects.

Therefore, the aims of the present study initially were to investigate (1) whether acute hyperinsulinaemia inhibits VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB production to the same extent and (2) factors that might influence the extent to which VLDL<sub>1</sub> is suppressed in a group of normoglycaemic individuals with a wide range of adiposity, fasting TG and insulin resistance. This was conducted using a novel, non-stable isotope method to determine VLDL<sub>1</sub>-TG and -apoB kinetics (Al-Shayji *et al.*, 2007), the development of which was previously described in [Chapter 3](#).

## 4.2 Subjects and Methods

### 4.2.1 Subjects

Eight non-smoking healthy subjects (6 men and 2 women) were included in this study after giving written informed consent. All subjects had normal thyroid, liver and renal function and none had acute illness, a history of known cardiovascular disease or hypertension, nor were under medication known to influence carbohydrate or lipid metabolism. The subject information sheet, consent form and health questionnaire were the same as those used for the ‘Intralipid Method’ study (Appendices A1 & B1). Two subjects had the E3/2 phenotype, five had the E3/3 phenotype and one had the E4/4 phenotype. The subjects’ characteristics are shown in **Table 4.1**. The study protocol was approved by the Research Ethics Committee of the North Glasgow University Hospitals NHS Trust.

### 4.2.2 Study Design

Each subject participated in 2 Intralipid trials (Control and Glucose) in random order at an interval of 2-3 weeks for men and 4 weeks (to control for menstrual cycle) for women. In the Glucose trial, subjects were given an oral bolus of a glucose drink (30 g in 120 ml) an hour before an Intralipid infusion (see below) followed by 10 g of glucose (in 40 ml) at 15-min intervals through out the duration of the trial. This was substituted with equivalent volumes of water in the Control trial (**Figure 4.1**). The dose and frequency of the glucose drink were previously determined, as explained in [section 2.10](#), to elucidate sufficient suppression in NEFA concentrations, before administering the Intralipid infusion, and maintain, as far as possible, a steady state of hyperinsulinaemia and hyperglycaemia throughout the trial.

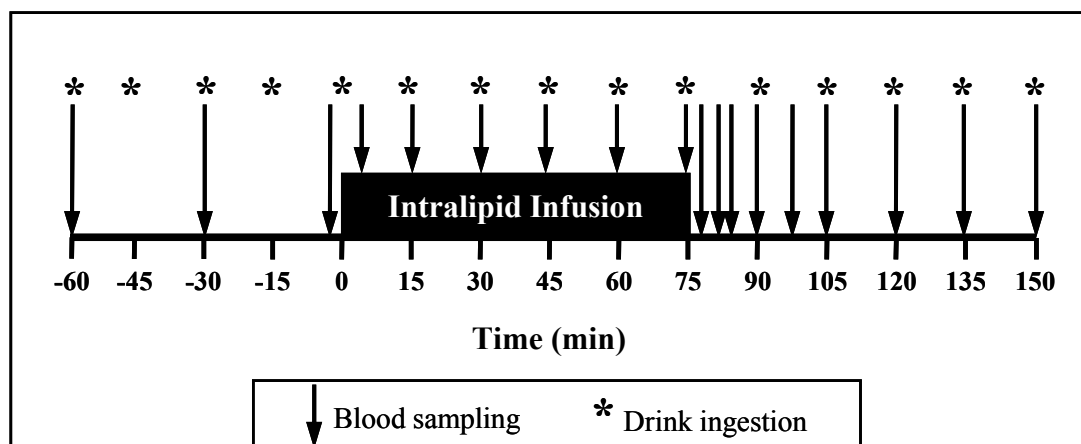


Subjects were requested not to exercise for three days before their study days as this is known to affect TRL metabolism (Gill & Hardman, 2003). In addition, they were asked to weigh and record their dietary intake for two days prior to the first Intralipid test and this diet was replicated for the second trial.

**Table 4.1: Subjects' physical characteristics and mean fasting concentrations (n=8).**

	Mean	Range
Age (years)	34	(22-55)
Body Mass (kg)	73.9	(56.0-96.5)
Body mass index (kg.m <sup>-2</sup> )	25.8	(21.0-34.7)
Waist circumference (cm)	83.8	(73.0-104.5)
Waist/hip ratio	0.82	(0.77-0.88)
ALT (U.l <sup>-1</sup> )	20	(15-26)
Triglycerides (mmol.l <sup>-1</sup> )*	1.17	(0.52-2.36)
Total cholesterol (mmol.l <sup>-1</sup> )*	4.21	(3.37-5.65)
HDL cholesterol (mmol.l <sup>-1</sup> )*	1.37	(0.81-1.81)
LDL cholesterol (mmol.l <sup>-1</sup> )*	2.29	(1.47-3.63)
Glucose (mmol.l <sup>-1</sup> )*	5.32	(4.50-5.91)
Insulin (mU.l <sup>-1</sup> )*	7.15	(2.48-15.84)
HOMA <sub>IR</sub> *	1.71	(0.50-3.59)
NEFA (mmol.l <sup>-1</sup> )*	0.50	(0.28-0.77)
Fasting VLDL <sub>1</sub> -TG (mmol.l <sup>-1</sup> )*	0.47	(0.12-1.31)
Fasting VLDL <sub>1</sub> -apoB (mg.dl <sup>-1</sup> )*	1.51	(0.38-4.04)

\*Values are the mean of Control and Glucose trials



**Figure 4.1: Study Protocol.** In the Glucose trial a bolus drink of 30 g of glucose in 120 ml water was given at -60 mins, followed by 10 g of glucose in 40 ml of water at 15-min intervals thereafter. In the Control trial, equivalent volumes of water were given at the same intervals.

### 4.2.3 Intralipid Infusion

TRL kinetics were determined using the ‘Intralipid Method’ (Al-Shayji *et al.*, 2007) as previously described in [Chapter 3](#). Briefly, subjects reported to the Clinical Investigation Suite in the Department of Vascular Biochemistry in Glasgow Royal Infirmary after an overnight fast of 12 h where they were given a bolus dose of 0.1 g.kg<sup>-1</sup> of 20% Intralipid followed immediately by an intravenous infusion of 0.1 g.kg<sup>-1</sup>.h<sup>-1</sup> of 10% Intralipid for 75 min. Multiple EDTA blood samples were drawn at baseline, before and at 15-min intervals during infusion as well as at 2.5, 5, 10, 15, 20, 30, 45, 60 and 75 min post-infusion.

### 4.2.4 Analytical Methods

Plasma total and HDL cholesterol concentrations were measured in the fasted state and glucose, TG and NEFA concentrations were measured in all time points. LDL cholesterol was calculated in the fasted state using the Friedewald equation (Friedewald *et al.*, 1972). Plasma insulin concentrations were analysed in the fasted state and every 30 min throughout the trial. Serum ALT concentrations were measured at screening and the apoE phenotype was determined for each subject. All analyses were performed as previously described in [sections 2.4.2](#) to 2.4.4.

Intralipid (S<sub>f</sub> >400) and VLDL<sub>1</sub> (S<sub>f</sub> 60-400) fractions were separated from whole plasma as previously described in [sections 3.2.4](#) and [2.4.1](#), respectively (Al-Shayji *et al.*, 2007). TG concentrations (corrected for glycerol, [section 3.2.7](#)) were measured in the Intralipid and VLDL<sub>1</sub> fractions in all time points and apoB concentrations were measured directly in the VLDL<sub>1</sub> fraction by immunoturbidimetry as previously described in [section 2.9.2.3](#).

### 4.2.5 Calculations and Statistical Analyses

VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB productions rates were calculated from their linear rises in the VLDL<sub>1</sub> fraction during infusion. Intralipid-TG clearance rate was calculated from its exponential decay post-infusion as described in [Chapter 3](#) (Al-Shayji *et al.*, 2007). The production ratio of TG/apoB (mol:mol) was calculated as an index of the size of the VLDL<sub>1</sub> particle being produced.

Time-averaged concentrations for glucose, insulin and NEFA were calculated from their respective areas under the curve from the start of the Intralipid infusion to 75 min post-infusion.

Statistical analyses were performed using MINITAB for Windows (Version 14.0, MINITAB Inc., State College, PA) and STATISTICA (Release 6.0, StatSoft, Inc, USA). Normality was checked for all the data using the Anderson-Darling test. When data did not approximate a normal distribution, these were log-transformed, specifically fasting TG, kinetic data and the time-averaged concentrations of glucose and insulin. Time-trends were tested using two-way ANOVA (trial  $\times$  time) with repeated measures on both factors. *Post hoc* Fisher LSD tests were used to indentify exactly where any differences lay. Paired *t*-tests were used to compare fasting concentrations and kinetic variables between the Glucose and Control trials. HOMA<sub>IR</sub> was used as a validated surrogate measure of insulin resistance (Matthews *et al.*, 1985) and was calculated as  $\text{insulin (mU.l}^{-1}) \times \text{glucose (mmol.l}^{-1}) / 22.5$ . To determine factors which predicted the change in VLDL<sub>1</sub>-TG and -apoB production and Intralipid-TG clearance rates between Control and Glucose trials, Pearson product-moment correlations were performed between the changes (Control minus Glucose) in kinetic variables on one hand and HOMA<sub>IR</sub>, ALT, NEFA and the differences in the time-averaged concentrations of glucose and insulin on the other hand. Significance was accepted at the  $P < 0.05$  level. Data are presented as mean  $\pm$  SEM unless otherwise stated. There were no data available to base a formal power calculation on for the present study. However, in a previous study investigating the effects of hyperinsulinaemia on VLDL<sub>1</sub>-apoB production using stable isotope methods, a significant effect was seen with five subjects (Malmstrom *et al.*, 1998), suggesting that studying eight subjects in the present study would be sufficient to detect a significant effect of hyperinsulinaemia on VLDL<sub>1</sub> metabolism, should such an effect exist.

### 4.3 Results

#### 4.3.1 Fasting Plasma Concentrations

There was no significant difference in fasting concentrations between the Glucose and Control trials. **Table 4.1** therefore shows the mean of fasting plasma lipid, glucose, insulin, HOMA<sub>IR</sub> and NEFA concentrations in both trials.

#### 4.3.2 Glucose, Insulin and NEFA Concentrations during and Post-Infusion

Mean plasma glucose concentrations were significantly higher throughout the Glucose trial compared with the Control trial ( $P = 0.005$ , **Figure 4.2A**). In the Glucose trial, glucose concentrations increased significantly within 30 min of the initial glucose ingestion (i.e. 30 min prior to the Intralipid infusion) (from  $5.35 \pm 0.18$  to  $7.92 \pm 0.59$  mmol.l<sup>-1</sup>,  $P < 0.001$ ) and averaged  $7.80 \pm 0.12$  mmol.l<sup>-1</sup> during infusion. In the Control trial, glucose concentrations decreased slightly from baseline during the first hour before infusion (from  $5.28 \pm 0.15$  to  $5.06 \pm 0.11$  mmol.l<sup>-1</sup>,  $P = 0.01$ ), but remained unchanged at an average of  $5.08 \pm 0.08$  mmol.l<sup>-1</sup> ( $P = 0.44$ ) during infusion.

As expected, insulin concentrations were significantly higher in the Glucose trial compared to the Control trial ( $P = 0.03$ , **Figure 4.2B**). Like glucose concentrations, insulin concentrations increased significantly 30 min after the start of the glucose drinks in the Glucose trial (from  $7.25 \pm 1.80$  mU.l<sup>-1</sup> to  $47.61 \pm 8.78$  mU.l<sup>-1</sup>,  $P = 0.02$ ) and remained elevated ( $P = 0.47$ ) throughout the trial with an average concentration of  $64.91 \pm 9.7$  mU.l<sup>-1</sup>. This is in contrast to the Control trial, where plasma insulin concentrations did not change significantly from baseline ( $P = 0.91$ ) with an average of  $6.87 \pm 0.17$  mU.l<sup>-1</sup>.

In the hour prior to the start of the Intralipid infusion, mean plasma NEFA concentrations declined rapidly in the Glucose trial from  $0.46 \pm 0.07$  at baseline to  $0.16 \pm 0.04$  mmol.l<sup>-1</sup> immediately before infusion ( $P < 0.001$ , **Figure 4.2C**), but did not change significantly from baseline in the Control trial ( $0.55 \pm 0.01$  mmol.l<sup>-1</sup>,  $P = 0.27$ ). When the Intralipid infusion started in both trials, NEFA concentrations

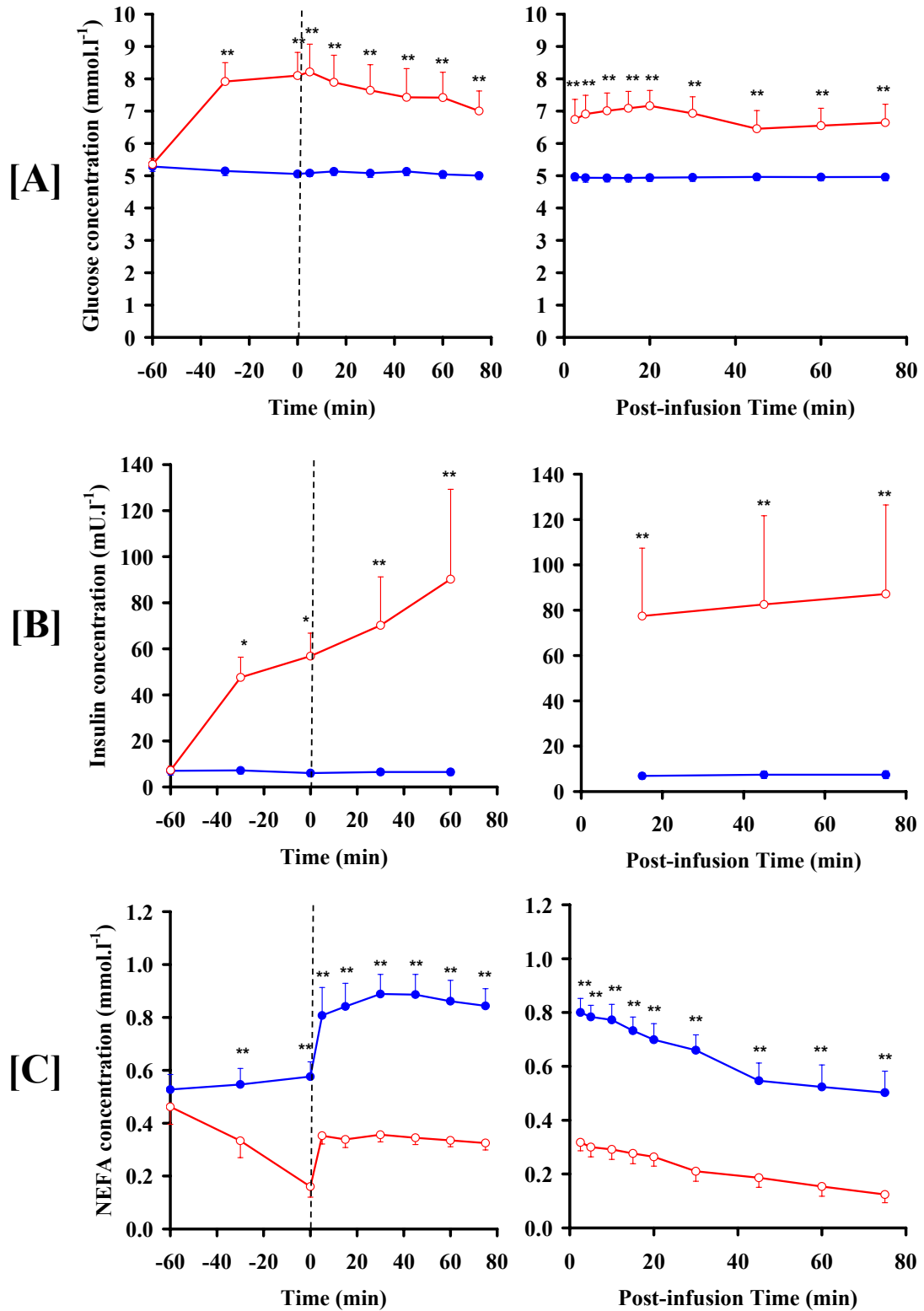


Figure 4.2: Mean [A] glucose, [B] insulin and [C] NEFA concentrations before and during Intralipid infusion (left panel) and post-infusion (right panel) between the Control (●) and Glucose (○) trials. The dotted line represents the beginning of a bolus Intralipid dose of 0.1 g.kg<sup>-1</sup> followed immediately by a constant Intralipid infusion of 0.1 g.kg<sup>-1</sup>.h<sup>-1</sup>. Significant differences between the two trials were tested using two-way ANOVA with (\*) representing  $P < 0.05$  and (\*\*)  $P < 0.001$ .

increased about 2.5-fold and leveled during infusion at an average concentration of  $0.85 \pm 0.04 \text{ mmol.l}^{-1}$  in the Control trial and  $0.34 \pm 0.02 \text{ mmol.l}^{-1}$  in the Glucose trial. Overall, NEFA concentrations were significantly lower throughout the Glucose trial compared to the Control trial ( $P < 0.001$ ).

#### **4.3.3 The Change in Production and Clearance Rates**

Individual and mean VLDL<sub>1</sub>-TG and -apoB production and Intralipid-TG clearance rates in the Control and Glucose trials according to apoE phenotypes are shown in **Figure 4.3**. The VLDL<sub>1</sub>-TG production rate was lower in the Glucose trial in 6 out of 8 subjects with an overall mean production rate of  $805.7 \pm 264.0 \text{ mg.h}^{-1}$  compared to  $969.3 \pm 181.6 \text{ mg.h}^{-1}$  in the Control trial ( $P = 0.05$ , **Figure 4.3A**). Similarly, VLDL<sub>1</sub>-apoB production rate was lower in the Glucose trial in the same 6 subjects with an overall mean production rate of  $21.8 \pm 5.3 \text{ mg.h}^{-1}$  in the Glucose trial versus  $25.8 \pm 4.1 \text{ mg.h}^{-1}$  in the Control trial ( $P = 0.04$ , **Figure 4.3B**). Overall, there was a 25.4% and 20.9% reduction in VLDL<sub>1</sub>-TG and -apoB production rates, respectively, in the Glucose trials. It is unclear why these two subjects responded differently. However, of note, one of them had the highest HOMA<sub>IR</sub> value (3.59) among all subjects.

On the other hand, the Intralipid-TG clearance response differed widely amongst subjects (**Figure 4.3C**). While four subjects showed a slight increase in Intralipid-TG clearance rate in the glucose trial, one subject showed a dramatic increase and the remaining three subjects showed a decreased Intralipid-TG clearance rate. The reason for these different responses is unclear. However, it is possibly due to differences in insulin resistance: the subject who had the highest increase in Intralipid-TG clearance rate, had the lowest HOMA<sub>IR</sub> of 0.5 (mean between the Control and Glucose trials), while the three subjects who had decreased clearance rates in the glucose trial, exhibited the highest HOMA<sub>IR</sub> concentrations amongst the 8 subjects (1.97, 3.47 and 3.59). This suggestion is supported by a strong correlation between HOMA<sub>IR</sub> and the change in Intralipid-TG clearance rate (see below). There was no significant difference in Intralipid-TG clearance between the Control and Glucose trials ( $69.4 \pm 12.7$  and  $103.5 \pm 49.8 \text{ pools.d}^{-1}$  respectively,  $P = 0.95$ ), although clearance was 16.9% higher in the Glucose trial. However, this numerical

increase in Intralipid-TG clearance was entirely due to one subject who showed a dramatic increase in Intralipid-TG clearance following glucose ingestion. If this subject was excluded from analysis, the Intralipid-TG clearance rates between the Control and Glucose trials were almost identical ( $58.4 \pm 7.3$  and  $54.8 \pm 11.9$  pools.d<sup>-1</sup>, respectively,  $P = 0.600$ ).

#### **4.3.4 VLDL<sub>1</sub> Particle Size**

The size of the VLDL<sub>1</sub> particle being produced (estimated as TG/apoB molar ratio) in response to hyperinsulinaemia and hyperglycaemia varied widely among subjects: five subjects exhibited a decrease in VLDL<sub>1</sub> particle size (range 7.4 - 50.6%), whereas 3 subjects showed an increased VLDL<sub>1</sub> particle size (range 10.9 - 38.2%). Unlike the Intralipid-TG clearance rate, the reason for this variable response in VLDL<sub>1</sub> particles size is unclear and could not be explained by differences in HOMA<sub>IR</sub>. In total, there was a mean reduction of  $7.6 \pm 10.2\%$  in VLDL<sub>1</sub> particle size with glucose ingestion. However, this change was not statistically significant ( $P = 0.29$ ).

#### **4.3.5 Predictors of the Change in Production and Clearance Rates**

The change (Control minus Glucose) in NEFA concentrations at zero min (i.e. after the first hour of the start of the drink, and just before the Intralipid infusion) correlated significantly with the change in VLDL<sub>1</sub>-TG production rate ( $r = 0.79$ ,  $P = 0.02$ ), but not with the change in VLDL<sub>1</sub>-apoB production rate ( $r = 0.59$ ,  $P = 0.13$ , **Figure 4.4A**). Overall, the change in NEFA concentrations explained 62% ( $0.79^2$ ) of the variance in the change in VLDL<sub>1</sub>-TG production rate.

In addition, the author investigated whether ALT concentrations within normal range would predict the change in hepatic VLDL<sub>1</sub>-TG and -apoB production in response to hyperinsulinaemia and hyperglycaemia. It was noted that serum ALT concentrations correlated inversely and significantly with the change in VLDL<sub>1</sub>-TG ( $r = -0.83$ ,  $P = 0.01$ ) and VLDL<sub>1</sub>-apoB ( $r = -0.74$ ,  $P = 0.04$ ) production rates, explaining 69% and 55% of the variance in change in these responses, respectively (**Figure 4.4B**).

HOMA<sub>IR</sub> did not correlate significantly with the change in VLDL<sub>1</sub>-TG and -apoB production rates. However it correlated significantly with the change in Intralipid-

TG clearance rates ( $r = 0.72$ ,  $P = 0.04$ , **Figure 4.4C**), explaining 52% of the variance of the change in response. This indicates that subjects who had the lowest HOMA<sub>IR</sub> (i.e. the more insulin sensitive individuals) had the biggest increase in Intralipid-TG clearance rate in response to hyperinsulinaemia.

There was no apparent effect of apoE phenotype on the change in VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB production rates or Intralipid-TG clearance rate (**Figure 4.3**).



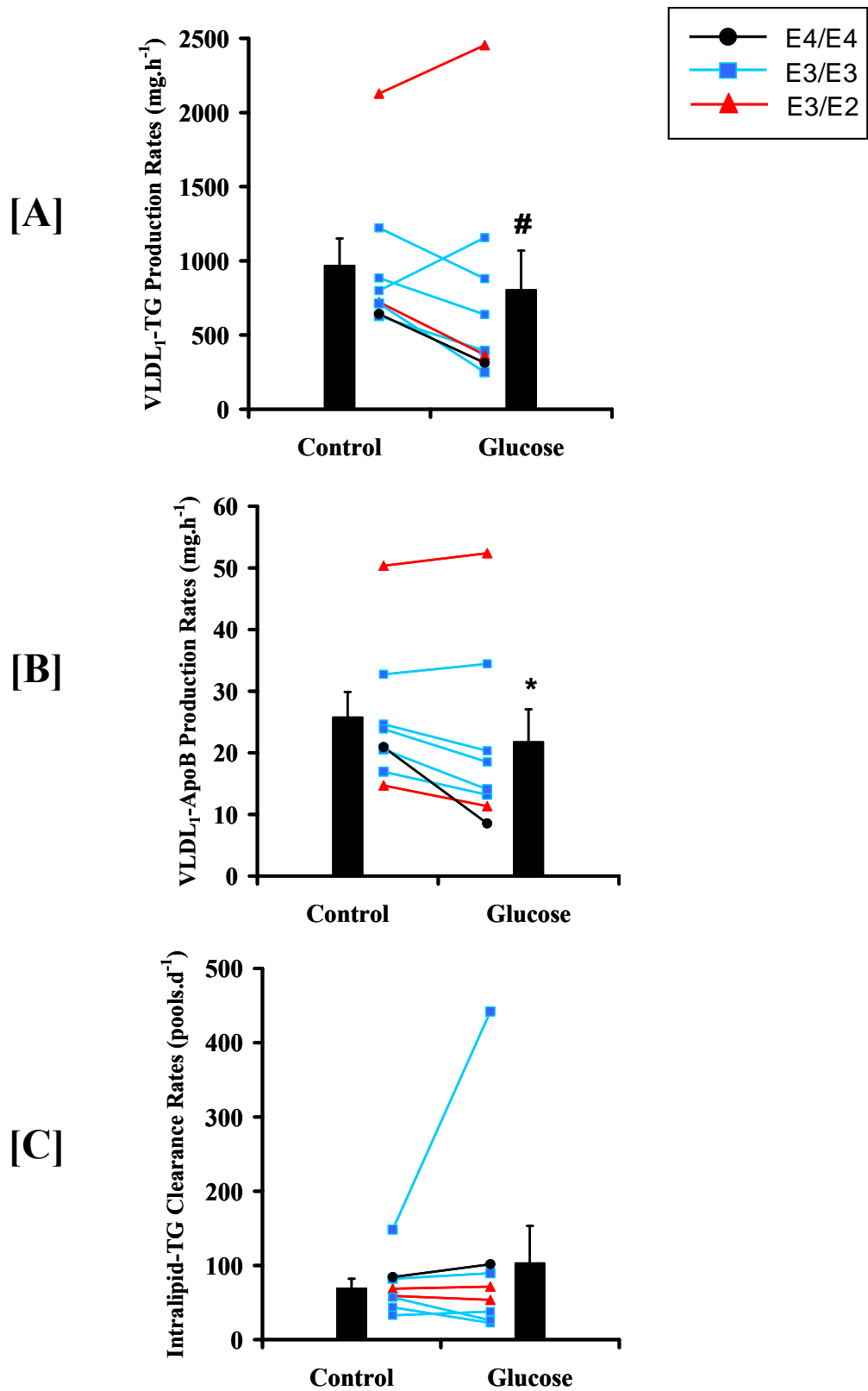


Figure 4.3: Individual (according to apoE phenotype) and mean (bars) [A] VLDL<sub>1</sub>-TG and [B] VLDL<sub>1</sub>-apoB production rates and [C] Intralipid-TG clearance rate in the Control and Glucose trials (n = 8). (#)  $P = 0.05$ , (\*)  $P < 0.05$ .

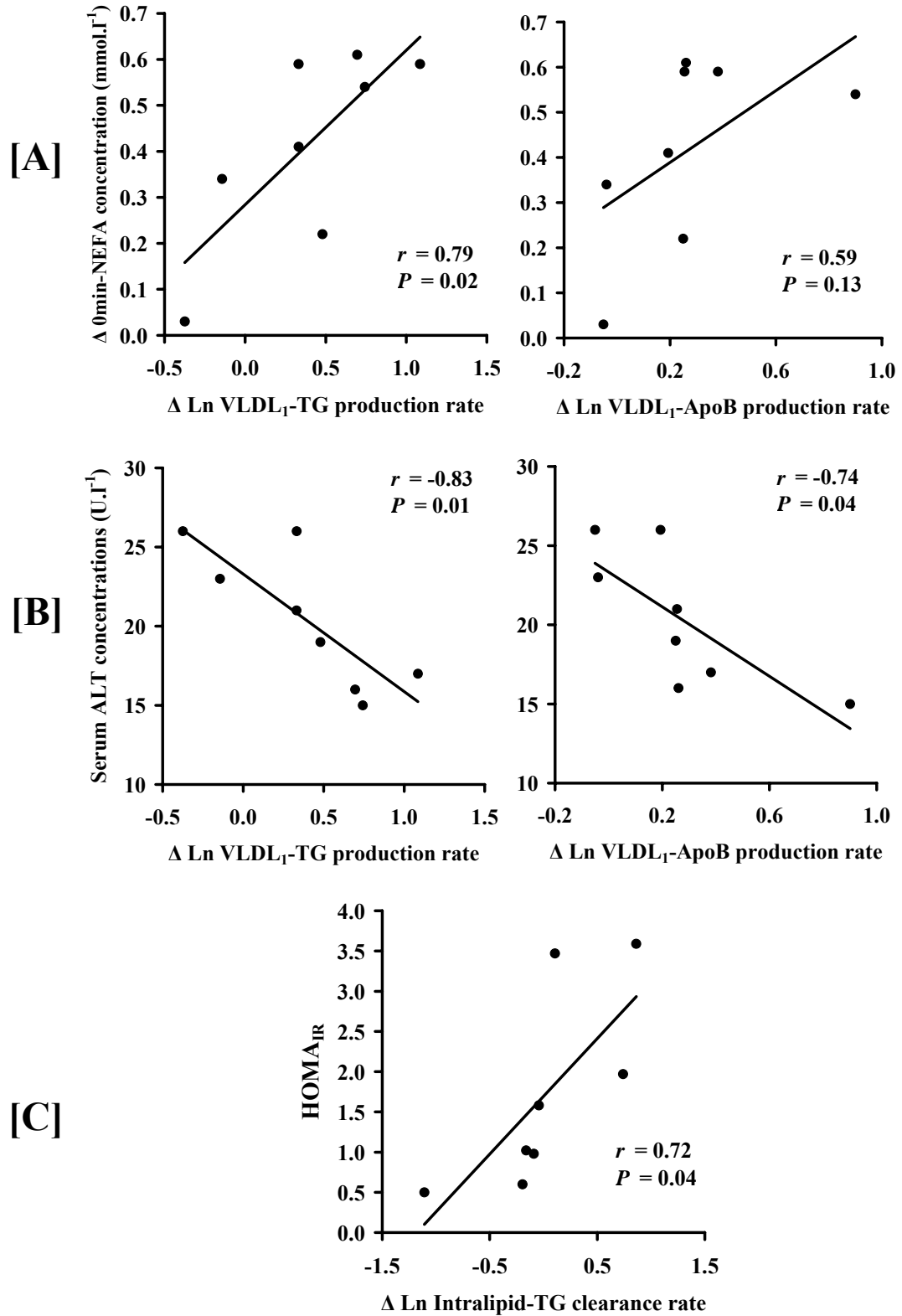


Figure 4.4: Scatterplots (with linear-regression lines of ‘best-fit’) illustrating the relationships between the change (Control minus Glucose) in [A] NEFA concentrations at 0 min (just before the start of the Intralipid infusion) and the change in VLDL<sub>1</sub>-TG (left) and VLDL<sub>1</sub>-apoB production rates (right); between [B] serum ALT concentrations and the change in VLDL<sub>1</sub>-TG (left) and VLDL<sub>1</sub>-apoB production rates (right); and [C] HOMA<sub>IR</sub> and the change in Intralipid-TG clearance rate. N = 8, *r* and *P* values are for Pearson product-moment correlations between variables.

#### 4.4 Discussion

This study aimed to investigate the acute effects of hyperinsulinaemia and hyperglycaemia on the production rates of both VLDL<sub>1</sub>-TG and -apoB and Intralipid-TG clearance rate in healthy normoglycaemic subjects. Firstly, the present results indicate that insulin acutely suppressed VLDL<sub>1</sub>-TG and -apoB production rates by 25% and 21%, respectively. However, as there was no significant difference in the size of the VLDL<sub>1</sub> particle being produced (i.e. TG/apoB ratio), this suggests equal suppression of VLDL<sub>1</sub>-TG and -apoB production. This is consistent with the recent findings of Adiels *et al.* (Adiels *et al.*, 2007), who showed that insulin acutely induced a 60% suppression in both VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB production rates in normal, non-diabetic subjects. The work of Adiels *et al.* was concomitant with the present study and, to-date, these are the only two studies that specifically investigated the effect of hyperinsulinaemia on VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB production rates.

The suppression in VLDL<sub>1</sub> production in response to insulin in the present study (at 21-25%) was less than the 50-60% reduction reported in VLDL<sub>1</sub>-TG (Adiels *et al.*, 2007) and VLDL<sub>1</sub>-apoB (Malmstrom *et al.*, 1997b; Malmstrom *et al.*, 1998) production rates. This could be due to the time of exposure to acute hyperinsulinaemia: in these studies, exogenous insulin was administered for 5-8.5 h, whereas in the present study, VLDL<sub>1</sub> production rates were determined during the first 135 min of exposure to hyperinsulinaemia. This is supported by Malmstrom *et al.* (Malmstrom *et al.*, 1997b), who reported a significant detectable reduction in VLDL<sub>1</sub>-apoB synthesis after 0.5 h of insulin infusion at an average of ~9% per h (range 0% – 24%).

Hepatic assembly of VLDL is a complex process (see [section 1.3.2](#)). Briefly, it starts in the rough ER with the co-translational lipidation of apoB-100 forming a ‘lipid pocket’ and the subsequent interaction with MTP, which catalyses the transfer of TG to the continually synthesised apoB-100 and allowing apoB to fold on a core of neutral lipids. This gives rise to a partially lipidated primordial particle (pre-VLDL), which is retained in the cell by interaction with chaperones and other ER proteins until it is further lipidated to form VLDL<sub>2</sub> or sorted to degradation. If apoB fails to be lipidated and misfolds, it is unfolded, retracted to the cytosol, ubiquitinated, and

sorted to proteasomal degradation (Adiels *et al.*, 2006a). VLDL<sub>2</sub> is converted to VLDL<sub>1</sub> in the Golgi apparatus by the uptake of a defined lipid load from a fairly rapidly mobilised cytosolic TG pool (Gibbons *et al.*, 2000). Insulin controls a number of enzymatic steps in the assembly and secretion of VLDL by the hepatocyte (Gibbons *et al.*, 2002; Sparks & Sparks, 1994). It downregulates MTP gene expression (Lin *et al.*, 1995), enhances apoB degradation (Sparks & Sparks, 1990), inhibits the secretion of newly synthesised apoB as well as inhibiting VLDL maturation by suppressing transfer of cytosolic TG to pre-VLDL particles (Brown & Gibbons, 2001). This is likely to explain the reduction in VLDL<sub>1</sub>-TG and -apoB production rates observed in the present study. In addition, insulin decreases NEFA flux to the liver, likely due to inhibition of hormone-sensitive lipase (Lewis *et al.*, 1993) and adipose tissue triglyceride lipase (Kershaw *et al.*, 2006). This consequently decreases substrate availability for VLDL assembly (Coppack *et al.*, 1994), as the influx of NEFA from adipose tissue to the liver normally accounts for 60-80% of hepatic TG (Donnelly *et al.*, 2005). Indeed, in this study, a rapid decrease in plasma NEFA concentrations occurred in the first hour, shortly after induction of hyperinsulinaemia in response to glucose ingestion, which correlated significantly and positively with the suppression in VLDL<sub>1</sub>-TG and -apoB production in response to insulin: subjects who had the biggest reduction in NEFA concentrations, had the biggest suppression in VLDL<sub>1</sub>-TG and -apoB production. It should be noted that although NEFA concentrations were significantly lower throughout the Glucose trial compared to the Control trial in response to insulin, NEFA concentrations increased during Intralipid infusion in both trials due to the 'spill-over' from Intralipid produced by LPL-mediated TG hydrolysis taking place directly into the circulation (Evans *et al.*, 1999). Therefore, NEFA concentrations during the infusion were not included in the correlations with changes in VLDL<sub>1</sub> production rates as they do not reflect the 'true' insulin-mediated suppression of NEFA.

Secondly, in the present study, hyperinsulinaemia did not significantly influence Intralipid-TG clearance rate. This is in contrast to the findings of Preiss-Landl *et al.* who reported that insulin upregulates LPL activity in adipose tissue (Preiss-Landl *et al.*, 2002). One likely explanation could be due to the heterogeneity of subjects' insulin resistance in the present study as there was a significant correlation between HOMA<sub>IR</sub> and the change in Intralipid-TG clearance rate. While glucose ingestion

caused an upregulation in clearance in the more insulin sensitive subjects, it induced a paradoxical downregulation of TG clearance in the more insulin resistant individuals. Indeed, it was previously reported that the more resistant an individual is to insulin-mediated glucose uptake, the lower will be the plasma post-heparin LPL activity and adipose tissue LPL mRNA levels (Maheux *et al.*, 1997).

Finally, the author investigated whether serum ALT concentrations, a marker for liver function and hepatic fat accumulation (Westerbacka *et al.*, 2004), would also predict the change in VLDL<sub>1</sub> production in response to hyperinsulinaemia. The present results demonstrate that serum ALT is a strong predictor of the suppression in VLDL<sub>1</sub>-TG and -apoB production, accounting for 69% and 55% of the variance in the change in production rates, respectively. This finding was recently supported by Adiels *et al.* (Adiels *et al.*, 2007) who showed that individuals with high liver fat failed to suppress VLDL<sub>1</sub> production in response to insulin. Whether this reduced suppression of VLDL<sub>1</sub> production by insulin is a result of liver fat, a consequence of hepatic insulin resistance or both remains to be uncertain (Adiels *et al.*, 2007). Similarly, it is unclear, in the present study, whether ALT (i.e. liver function) per se is the mediator of the change in VLDL<sub>1</sub> production or whether ALT is just a marker for liver fat.

In summary, this study demonstrated that, in normoglycaemic apparently healthy individuals, (1) insulin suppressed VLDL<sub>1</sub>-TG and -apoB production; and (2) the more insulin sensitive the subject is, the bigger the increase in Intralipid-TG clearance rates due to hyperinsulinaemia and hyperglycaemia. In addition, ALT concentration – a marker of liver function and associated with liver fat content – strongly predicted individuals' ability to suppress VLDL<sub>1</sub>-TG and -apoB in response to hyperinsulinaemia. Not only do these findings help the basic scientific understanding of the effect of insulin on VLDL<sub>1</sub> production, they may also help identifying individuals at higher risk of developing atherogenic dyslipidaemia which is believed to be initiated by insulin resistance, liver fat and hepatic overproduction of VLDL<sub>1</sub> particles.

## 5. Effects of Moderate Exercise on VLDL<sub>1</sub> Kinetics in Overweight/Obese Middle-Aged Men

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### 5.1 Introduction

CVD is the main cause of death in the UK, with CHD, being the most common form, accounting for about 101,000 deaths each year (Allender *et al.*, 2007). Elevated plasma concentrations of TG are considered an independent risk factor for CHD (Cullen, 2000). High concentrations of large very low density lipoprotein (VLDL<sub>1</sub>, S<sub>f</sub> 60-400) are the major determinant of plasma TG levels (Tan *et al.*, 1995; Hiukka *et al.*, 2005) and believed to initiate a chain of reactions that generate the atherogenic lipoprotein phenotype associated with insulin resistance conditions (e.g. obesity, type 2 diabetes and the metabolic syndrome) (Bloomgarden, 2007; Taskinen, 2003; Ginsberg *et al.*, 2005). Thus, reducing VLDL<sub>1</sub> concentration is likely to induce clinically important changes to the atherosclerotic risk profile.

While patient groups often undergo pharmacological treatments for these lipoprotein disorders, obesity and insulin resistance are growing problems in the population: over half of UK adults are overweight (BMI > 25 kg.m<sup>-2</sup>) and over 20% are obese (BMI > 30 kg.m<sup>-2</sup>) (Allender *et al.*, 2006) and estimated prevalence of the metabolic syndrome in Scottish middle-aged men is ~25% (Sattar *et al.*, 2003). Therefore, it is unfeasible and probably undesirable to subject large sections of the population to long-term pharmacological therapies. Moderate exercise is one potential non-pharmacological therapy for elevated VLDL<sub>1</sub> concentrations. There is clear evidence that exercise of this nature can lower fasting and postprandial TG concentrations by 20-25%, mostly in the VLDL<sub>1</sub> fraction (Gill *et al.*, 2006), in population groups at increased risk of cardiovascular disease, such as centrally obese middle-aged men (Gill *et al.*, 2004b) and postmenopausal women (Gill & Hardman, 2000). This effect is seen following a single exercise session, so is not mediated by weight loss (Gill *et al.*, 2004b; Gill & Hardman, 2000; Gill *et al.*, 2006). However the mechanism(s)

responsible for this TG-reduction are unclear and require further elucidation as it could reflect reduced hepatic VLDL<sub>1</sub> production and/or increased LPL-mediated VLDL<sub>1</sub> clearance. Interestingly, a prior session of moderate exercise induces a larger TG-reduction in the VLDL<sub>1</sub> fraction than the chylomicrons fraction (Gill *et al.*, 2001a; Gill *et al.*, 2006). Because chylomicrons are preferred substrates to LPL above VLDL<sub>1</sub> (Karpe *et al.*, 1993a; Fisher *et al.*, 1995), this reduction in VLDL<sub>1</sub> was hypothesised to be the result of an exercise-induced suppression of hepatic VLDL<sub>1</sub> production rather than increased clearance. However, two recent stable-isotope kinetic studies have demonstrated that, in a group of lean recreationally active young men, a single session of a 90-120 min of moderate intensity exercise (at 60% of  $\dot{V}O_{2max}$ ) resulted in an increased clearance of total VLDL-TG (Magkos *et al.*, 2006; Tsekouras *et al.*, 2007) and a significant decrease in hepatic VLDL-apoB production (Magkos *et al.*, 2006) with no significant change in hepatic VLDL-TG production (Magkos *et al.*, 2006; Tsekouras *et al.*, 2007). However, studies investigating the effects of moderate exercise on large VLDL<sub>1</sub> kinetics, especially in overweight/obese middle-aged men, a typical population at which exercise-for-health guidelines are targeted, are lacking. This is important as moderate exercise may have a considerable potential to be used as a first-line therapeutic option for preventing and treating the primary defect in obesity/insulin resistance-related dyslipidaemia.

Kinetics of VLDL are usually determined using stable-, or radio-, isotopic tracer methods, but these techniques are expensive and time-consuming which limits the scope of studies to which these methods can be applied. Earlier in this thesis ([Chapter 3](#)) the development of a relatively easy method to determine VLDL<sub>1</sub>-TG and -apoB production and clearance rates, as well as Intralipid-TG clearance rate (as a surrogate measure of chylomicron-TG clearance) was described (Al-Shayji *et al.*, 2007), which is used in the present study to elucidate the mechanisms by which moderate exercise induces TG lowering.

The aim of the present study was to investigate the effects of a 120-min session of prior moderate exercise on VLDL<sub>1</sub>-TG and -apoB kinetics and Intralipid-TG clearance rate in a group of overweight/obese middle-aged adults using the Intralipid method.

## 5.2 Materials and Methods

### 5.2.1 Subjects

Twelve overweight/obese, middle-aged men participated in this study; their characteristics are shown in **Table 5.1**. All subjects were apparently healthy, normotensive, normoglycaemic, non-smokers who displayed no symptoms of coronary heart disease during a clinical exercise stress test ([section 2.5](#)). None was taking any drugs known to affect lipid or carbohydrate metabolism. The subject information sheet, consent form and health questionnaire are shown in Appendices A2 & B2. One subject had E3/2 apoE phenotype, 4 had E3/3 phenotype, 5 had E4/3 phenotype and 2 had E4/4 phenotype. The study was conducted with the approval of North Glasgow University Hospitals NHS Trust Ethics Committee and subjects gave written informed consent prior to participation.

**Table 5.1: Characteristics of the participants (n=12).**

	Mean $\pm$ SD
Age (years)	44 $\pm$ 8.4
Body Mass (kg)	95.0 $\pm$ 17.1
Body mass index (kg.m <sup>-2</sup> )	31.3 $\pm$ 5.1
Waist circumference (cm)	106.9 $\pm$ 13.2
Waist/hip ratio	0.98 $\pm$ 0.05
Maximal oxygen uptake (ml.kg <sup>-1</sup> .min <sup>-1</sup> )	37.9 $\pm$ 7.2
ALT (U.l <sup>-1</sup> )	39 $\pm$ 10

### 5.2.2 Study Design

Each subject underwent two Intralipid infusions (see below) within an interval of 2 weeks. Preconditions (i.e. control and exercise) were different and administered in random order. On the afternoon prior to one Intralipid infusion subjects walked 120 min on a treadmill at an intensity of  $\sim 50\%$   $\dot{V}O_{2\max}$  (Exercise trial). In the other trial, subjects performed no exercise on the day preceding the other Intralipid infusion (Control trial). They were asked to weigh and record their dietary intake for two days prior to their first Intralipid infusion and replicate this prior to the second test and were instructed to refrain from alcohol consumption on those two days. No



exercise other than the treadmill walk in the Exercise trial was performed during the three days prior to each Intralipid infusion.

### **5.2.3 Treadmill Walk**

A preliminary submaximal incremental treadmill test was performed as previously described in [section 2.6](#) at least 1 week prior to the first Intralipid test to estimate  $\dot{V}O_{2\max}$  and determine the speed and gradient required to elicit 50%  $\dot{V}O_{2\max}$ . The 120-min walk in the Exercise trial was completed ~16-18 h prior to the Intralipid test. Heart rate, O<sub>2</sub> uptake and CO<sub>2</sub> production were measured as previously described in [section 2.8](#). Ratings of perceived exertion (Borg, 1973) were obtained at 15-min intervals during the walk.

### **5.2.4 Intralipid Infusion**

VLDL<sub>1</sub> kinetics were determined using the 'Intralipid Method' as previously described in [Chapter 3](#) (Al-Shayji *et al.*, 2007) with a slight modification in the frequency of blood sampling during infusion. Briefly, subjects reported to the Clinical Investigation Suite in the Department of Vascular Biochemistry at Glasgow Royal Infirmary in the morning after an overnight fast of 12 h and they were given a bolus dose of 0.1 g.kg<sup>-1</sup> of 20% Intralipid followed immediately by an intravenous infusion of 0.1 g.kg<sup>-1</sup>.h<sup>-1</sup> of 10% Intralipid for 75 min. Multiple EDTA blood samples were drawn at baseline, before and at 10-min intervals during infusion as well as at 2.5, 5, 10, 15, 20, 30, 40 and 60 min post-infusion. The post-infusion period was reduced from 75 to 60 min when it became clear that 60 min was sufficient to calculate the Intralipid-TG clearance rate.

### **5.2.5 Analytical Methods**

Plasma glucose, insulin, total and HDL cholesterol and apos C-II, C-III and E concentrations were analysed in the fasted state. TG and NEFA concentrations were analysed in all time points. LDL cholesterol was calculated using the Friedewald equation (Friedewald *et al.*, 1972). Serum ALT concentrations were measured at screening. ApoE phenotype was determined for each subject by isoelectric focusing

using Western blot techniques. All analytical methods were performed as previously described in [sections 2.4.2](#) to 2.4.4.

Intralipid ( $S_f > 400$ ) was separated from whole plasma as previously described in [section 3.2.4](#) with a minor modification. In the present subjects (typical candidates for hyperlipidaemia), it was found that the Intralipid fraction was technically easier to separate at 23° C instead of 4° C. The VLDL<sub>1</sub> ( $S_f$  60-400) fraction was separated as previously described in [section 2.4.1](#). TG concentrations were measured in the Intralipid and VLDL<sub>1</sub> fractions in all time points and were corrected for glycerol as described in [section 3.2.7](#) (Al-Shayji *et al.*, 2007).

The composition of the VLDL<sub>1</sub> fraction (i.e. TG, cholesterol, FC and PL) was measured as described in [section 2.4.2](#). Total protein content was measured manually using a modified Lowry assay ([section 2.4.5](#)). CE concentration was determined by multiplying the difference between total cholesterol and free cholesterol concentrations (in mg.dl<sup>-1</sup>) by 1.68 to account for the difference in mass between cholesterol and CE. VLDL<sub>1</sub> concentration was calculated by summing the concentrations of these components in mg.dl<sup>-1</sup> (i.e. TG, FC, CE, PL and protein). ApoB concentrations were measured directly in the VLDL<sub>1</sub> fraction by automated immunoturbidimetry (see [section 2.9.2.3](#)). In four subjects, where VLDL<sub>1</sub>-apoB was not detectable by the immunoturbidimetry method, apoB concentrations were measured manually, as described in [section 2.9.2.2](#) by precipitation with Isopropanol and subsequent protein measurement by Lowry method ([section 2.4.5](#)).

### **5.2.6 Calculations and Statistics**

VLDL<sub>1</sub>-TG and -apoB production rates and FCRs, as well as Intralipid-TG clearance rates were determined as previously described in [Chapter 3](#) (Al-Shayji *et al.*, 2007).

Net energy expenditure during the 120-min walk was calculated using indirect calorimetry assuming no protein oxidation (Frayn, 1983). HOMA was used as a validated surrogate measure of insulin resistance (IR) (Matthews *et al.*, 1985).

The number of apoC-II, apoC-III and apoE molecules per VLDL<sub>1</sub> particle were calculated, in the fasted state, by dividing the apolipoprotein concentrations in

mmol.l<sup>-1</sup> by the apoB concentration in mmol.l<sup>-1</sup>. In addition, VLDL<sub>1</sub> TG/apoB ratio (mol:mol), representing the size of the particle in the circulation; CE/TG ratio (mol:mol), representing the composition of its core; CE/apoB ratio (mol:mol), representing the CE content per VLDL<sub>1</sub> particle; and FC/PL ratio (mg:mg), representing the composition of its surface layer, were also calculated in the fasted state.

Statistical analyses were performed using MINITAB for Windows (Version 14.0, MINITAB Inc., State College, PA) and STATISTICA (Release 6.0, StatSoft, Inc, USA). Normality was checked for all the data using the Anderson-Darling test. When data did not approximate a normal distribution, these were log-transformed, namely insulin, HOMA<sub>IR</sub>, VLDL<sub>1</sub>-apoB fasting concentrations, TG/apoB ratios, FC/PL ratio, Intralipid-TG clearance rate, VLDL<sub>1</sub>-apoB FCR and production rate, VLDL<sub>1</sub>-TG production rate, and the exercise-induced change in VLDL<sub>1</sub>-apoB FCR and production rate required transformation. Comparisons of fasting values, summary responses and kinetic data between Control and Exercise trials were made using paired *t*-tests and changes over the trial period were assessed by two-way ANOVA (trial × time) with repeated measures on both factors. *Post hoc* Fisher LSD tests were used to identify exactly where any differences lay. Significance was accepted at the *P* < 0.05 level. Data are presented as mean ± SEM unless otherwise stated.

To determine factors which predicted the change in VLDL<sub>1</sub>-TG and -apoB production and clearance rates as well as Intralipid-TG clearance rates between Control and Exercise trials, Pearson product-moment correlations were performed between the exercise-induced changes in kinetic variables on one hand and ALT, the exercise-induced changes in glucose, insulin, HOMA<sub>IR</sub> and NEFA on the other hand.

### 5.2.7 Power Calculations

The pilot data from 5 subjects who underwent two Intralipid infusions at the 0.1 g.kg<sup>-1</sup>.h<sup>-1</sup> Intralipid and 0.2 g.kg<sup>-1</sup>.h<sup>-1</sup> Intralipid doses (described in [Chapter 3](#)), indicate that the standard deviation for the within-subject variability in VLDL<sub>1</sub>-apoB and VLDL<sub>1</sub>-TG production are 12.7% and 20.1%, respectively, when preceding diet, exercise and alcohol intake are well-controlled. Based on these data, 12 subjects

would enable detection of an 11% difference in VLDL<sub>1</sub>-apoB production and an 18% difference in VLDL<sub>1</sub>-TG production, with 80% power at the 0.05 level. Groups of 8-12 subjects clearly show significant effects of a single moderate exercise session on postprandial TG metabolism where a 20-25% reduction in plasma TG concentration is typically observed (Gill & Hardman, 2003).

## 5.3 Results

### 5.3.1 Treadmill Walk

Subjects walked at a speed of  $4.5 \pm 0.2$  km.h<sup>-1</sup> up a gradient of  $6.0 \pm 0.6\%$ . All subjects completed the 120-min walk without difficulty, with an average perceived exertion of  $12.2 \pm 0.4$  (between ‘fairly light’ and ‘somewhat hard’) on the Borg scale of 6-20 (Borg, 1973). Mean  $\dot{V}O_2$  was  $18.6 \pm 0.8$  ml.kg<sup>-1</sup>.min<sup>-1</sup> ( $49.2 \pm 0.7\%$   $\dot{V}O_{2max}$ ), mean heart rate was  $123 \pm 3$  beat.min<sup>-1</sup> and net energy expenditure for the walk was  $3.5 \pm 0.1$  MJ ( $837 \pm 35$  kcal).

### 5.3.2 Fasting Concentrations in the Control and Exercise Trials

Mean plasma and VLDL<sub>1</sub> composition and concentrations in the fasted state are shown in **Table 5.2**. In four subjects, VLDL<sub>1</sub>-apoB was not detectable directly using the immunoturbidimetry method in at least one trial, and, therefore, was measured manually for both the Control and Exercise trials. In addition, fasting apoE concentrations were below limits of detection in 2 subjects in one or two trials, and, therefore, these two subjects were omitted from all apoE calculations. Exercise significantly ( $P < 0.05$ ) reduced fasting plasma TG and glucose concentrations by  $21.9 \pm 6.0\%$  and  $3.0 \pm 1.0\%$ , respectively, and increased NEFA by  $23.8 \pm 12.1\%$ . There was a reduction of  $12.6 \pm 6.7\%$  in insulin concentrations and of  $14.8 \pm 6.9\%$  in HOMA<sub>IR</sub> post-exercise, but these did not reach statistical significance ( $P = 0.080$  and  $0.059$ , respectively). Fasting VLDL<sub>1</sub> concentrations were significantly lower by  $32.6 \pm 9.7\%$  after exercise ( $P = 0.006$ ), as were fasting concentrations of VLDL<sub>1</sub>-TG ( $29.6 \pm 9.3\%$ ), VLDL<sub>1</sub>-Cholesterol ( $43.7 \pm 11.4\%$ ), VLDL<sub>1</sub>-apoB ( $38.2 \pm 9.8\%$ ), VLDL<sub>1</sub>-apoC-II ( $28.0 \pm 10.7\%$ ), VLDL<sub>1</sub>-apoC-III ( $25.0 \pm 12.1\%$ ) and VLDL<sub>1</sub>-apoE ( $27.6 \pm 11.4\%$ ,  $n = 10$ ), all  $P < 0.05$ . There were no significant differences in fasting concentrations of total, HDL- and LDL-cholesterol (**Table 5.2**).

### 5.3.3 TRL Kinetics in Responses to Exercise

**Figure 5.1** shows the mean percentage changes in VLDL<sub>1</sub>-TG and -apoB kinetics and Intralipid-TG clearance rate and **Figure 5.2** shows the absolute mean and individual production and clearance rates in response to exercise according to apoE phenotypes. There was a significant increase after exercise in Intralipid-TG clearance rate (Control:  $47.6 \pm 6.8$ , Exercise:  $68.1 \pm 9.7$  pools.d<sup>-1</sup>,  $P < 0.001$ ), VLDL<sub>1</sub>-TG FCR (Control:  $16.0 \pm 2.1$ , Exercise:  $29.1 \pm 4.4$  pools.d<sup>-1</sup>,  $P = 0.002$ ) and VLDL<sub>1</sub>-apoB FCR (Control:  $10.4 \pm 2.0$ , Exercise:  $25.6 \pm 5.1$  pools.d<sup>-1</sup>,  $P = 0.015$ ). However, there was no significant difference between the Control and Exercise trials in the production rates of VLDL<sub>1</sub>-TG (Control:  $1271.5 \pm 155.7$ , Exercise:  $1431.7 \pm 148.3$  mg.h<sup>-1</sup>,  $P = 0.104$ ) and VLDL<sub>1</sub>-apoB (Control:  $37.2 \pm 7.4$ , Exercise:  $41.5 \pm 5.4$  mg.h<sup>-1</sup>,  $P = 0.335$ ).

### 5.3.4 VLDL<sub>1</sub> Compositional Responses to Exercise

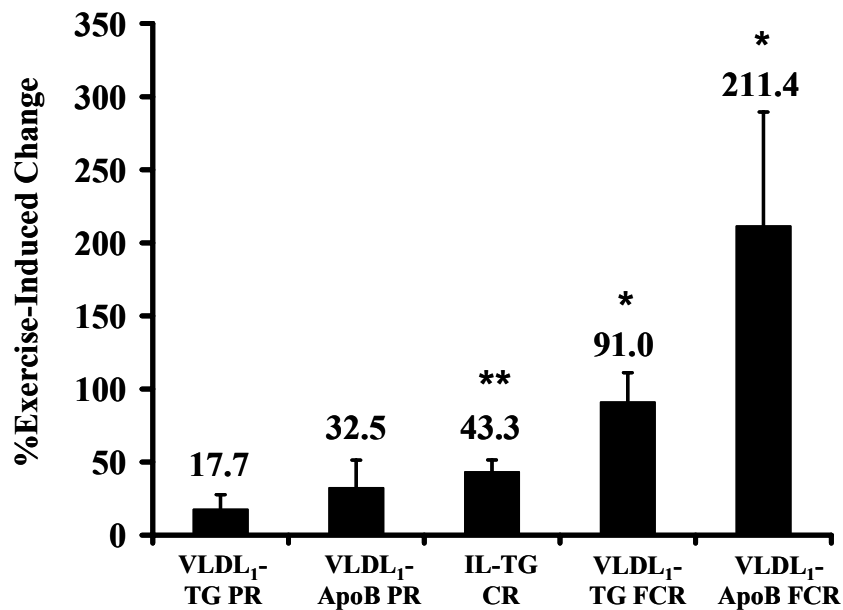
**Table 5.3** shows the exercise-induced compositional changes in VLDL<sub>1</sub> in the fasted state – reflecting the composition of circulating VLDL<sub>1</sub> particles. There was a tendency for a bigger VLDL<sub>1</sub> particle (TG/apoB ratio) in the circulation in the Exercise trial (Control:  $15392 \pm 2178$ , Exercise:  $19449 \pm 3150$ ). However, this did not reach statistical significance ( $P = 0.059$ ). On the other hand, the CE/TG ratio, reflecting the neutral lipid composition of the VLDL<sub>1</sub> core, was significantly lower in the Exercise trial (Control:  $0.17 \pm 0.02$ , Exercise:  $0.12 \pm 0.02$ ,  $P = 0.007$ ). Consequently, the CE/apoB ratio, reflecting CE content per VLDL<sub>1</sub> particle, was not significantly different between both trials (Control:  $2487 \pm 353$ , Exercise:  $2276 \pm 564$ ,  $P = 0.537$ ). PL was not detectable for one subject and, consequently, the FC/PL ratio in both trials for this subject were not included. There was no significant difference in the FC/PL ratio, which reflects the composition of the VLDL<sub>1</sub> surface layer, between the Control ( $0.32 \pm 0.01$ ) and Exercise ( $0.32 \pm 0.02$ ) trials ( $P = 0.606$ ,  $n = 11$ ).

Moreover, there was no significant difference in the fasting ratios of apoC-II/apoB (Control:  $32.6 \pm 5.0$ , Exercise:  $42.8 \pm 8.6$ ,  $P = 0.23$ ), apoC-III/apoB (Control:  $64.1 \pm 7.9$ , Exercise:  $83.8 \pm 13.9$ ,  $P = 0.10$ ) or apoE/apoB (Control:  $0.83 \pm 0.20$ , Exercise:  $1.12 \pm 0.33$ ,  $P = 0.409$ ,  $n = 10$ ).

**Table 5.2: Plasma and VLDL<sub>1</sub> concentrations in the fasted state**

	Control	Exercise	<i>P</i> -Value
Plasma TG (mmol.l <sup>-1</sup> )	1.54 ± 0.16	1.21 ± 0.15	0.004
Total Cholesterol (mmol.l <sup>-1</sup> )	5.17 ± 0.25	5.14 ± 0.24	0.843
HDL Cholesterol (mmol.l <sup>-1</sup> )	1.05 ± 0.06	1.08 ± 0.07	0.233
LDL Cholesterol (mmol.l <sup>-1</sup> )	3.38 ± 0.31	3.48 ± 0.29	0.409
Glucose (mmol.l <sup>-1</sup> )	5.76 ± 0.09	5.58 ± 0.09	0.010
Insulin (mU.l <sup>-1</sup> )	6.99 ± 1.30	6.29 ± 1.46	0.080
HOMA <sub>IR</sub>	1.81 ± 0.35	1.58 ± 0.37	0.059
NEFA (mmol.l <sup>-1</sup> )	0.55 ± 0.04	0.64 ± 0.03	0.026
VLDL <sub>1</sub> concentration (mg.dl <sup>-1</sup> )	94.9 ± 14.1	62.9 ± 11.4	0.006
VLDL <sub>1</sub> -TG (mmol.l <sup>-1</sup> )	0.67 ± 0.10	0.46 ± 0.08	0.007
VLDL <sub>1</sub> -C (mmol.l <sup>-1</sup> )	0.47 ± 0.07	0.28 ± 0.06	0.006
VLDL <sub>1</sub> -apo B (mg.dl <sup>-1</sup> )	2.77 ± 0.44	1.55 ± 0.30	0.004
VLDL <sub>1</sub> -apo CII (mg.dl <sup>-1</sup> )	1.22 ± 0.18	0.88 ± 0.16	0.017
VLDL <sub>1</sub> -apo CIII (mg.dl <sup>-1</sup> )	2.52 ± 0.40	1.84 ± 0.37	0.031
VLDL <sub>1</sub> -apo E (mg.dl <sup>-1</sup> )*	0.16 ± 0.06	0.11 ± 0.05	0.049

\* n = 10. All values are mean ± SEM. *P* values are for the difference between the Control and Exercise trials



**Figure 5.1: The mean percentage change in VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB production rates (PR) and fractional catabolic rates (FCR) and Intralipid (IL)-TG clearance rate (CR) in response to a single session of 120-min brisk walking at ~50% VO<sub>2max</sub>. (\* *P* < 0.05, \*\* *P* < 0.001).**

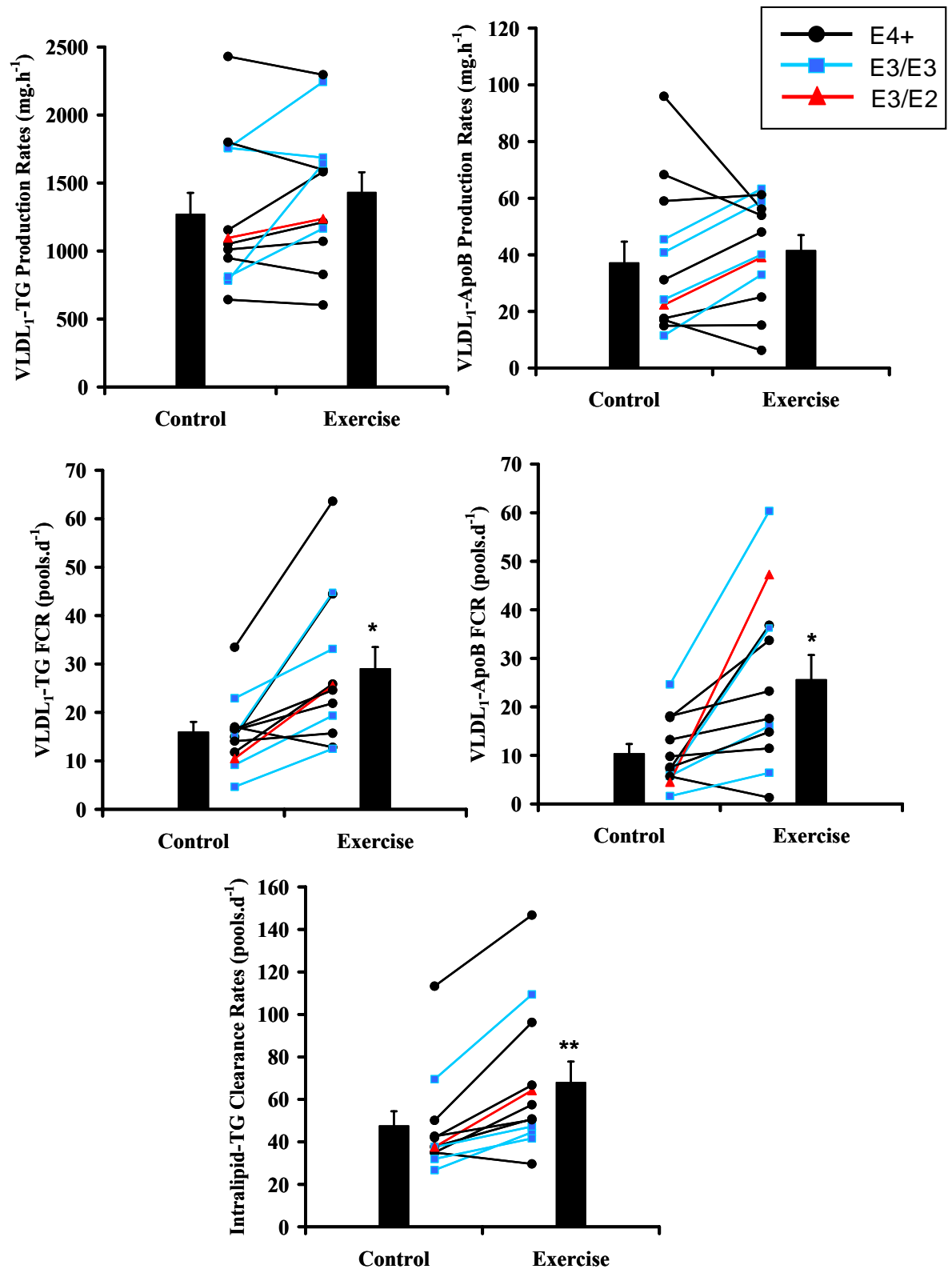


Figure 5.2: Individual (according to apoE phenotype) and mean (bars) VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB production rates (top panel) and fractional catabolic rates (FCR, middle panel) and intralipid-TG clearance rate (bottom) in response to a single session of 120-min brisk walking at ~50% VO<sub>2max</sub> (\*  $P < 0.05$ , \*\*  $P < 0.001$  for the difference in mean response between the Control and Exercise trials).

**Table 5.3: Exercise-induced compositional changes to circulating VLDL<sub>1</sub> particles**

	Control	Exercise	P-Value
TG/apoB ratio (mol:mol)	15392 ± 2178	19449 ± 3150	0.059
CE/TG ratio (mol:mol)	0.17 ± 0.02	0.12 ± 0.02	0.007
CE/apoB ratio (mol:mol)	2487 ± 353	2276 ± 564	0.537
FC/PL ratio (mg:mg)*	0.32 ± 0.01	0.32 ± 0.02	0.606

\* n = 11. All values are mean ± SEM.

### 5.3.5 Predictors of the Exercise-Induced Changes in TRL Kinetics

There was a significant correlation ( $r = 0.61$ ,  $P = 0.035$ ) between the exercise-induced change in Intralipid-TG clearance rate and the exercise-induced change in VLDL<sub>1</sub>-TG FCR (**Figure 5.3A**), which suggests that a single mechanism (likely LPL-mediated clearance) was likely to be responsible for both effects.

Although there was no statistically significant effect of exercise on fasting insulin concentrations and HOMA<sub>IR</sub>, a significant inverse relationship was found between the exercise-induced change in VLDL<sub>1</sub>-apoB FCR and the exercise-induced change in insulin concentrations ( $r = -0.65$ ,  $P = 0.022$ ), the exercise-induced change in glucose ( $r = -0.58$ ,  $P = 0.049$ ), and, consequently, with the exercise-induced change in HOMA<sub>IR</sub> ( $r = -0.68$ ,  $P = 0.016$ , **Figure 5.3B**). This shows that those who had the biggest reduction in HOMA<sub>IR</sub> had the biggest increase in VLDL<sub>1</sub>-apoB FCR.

No significant correlations were found between ALT or the exercise-induced change in NEFA and any of the exercise-induced changes in VLDL<sub>1</sub>-TG and -apoB FCR or production rates or Intralipid-TG clearance rate (all  $P > 0.05$ ). In addition, there was no significant effect of apoE phenotype on the response of the above kinetics to moderate exercise.



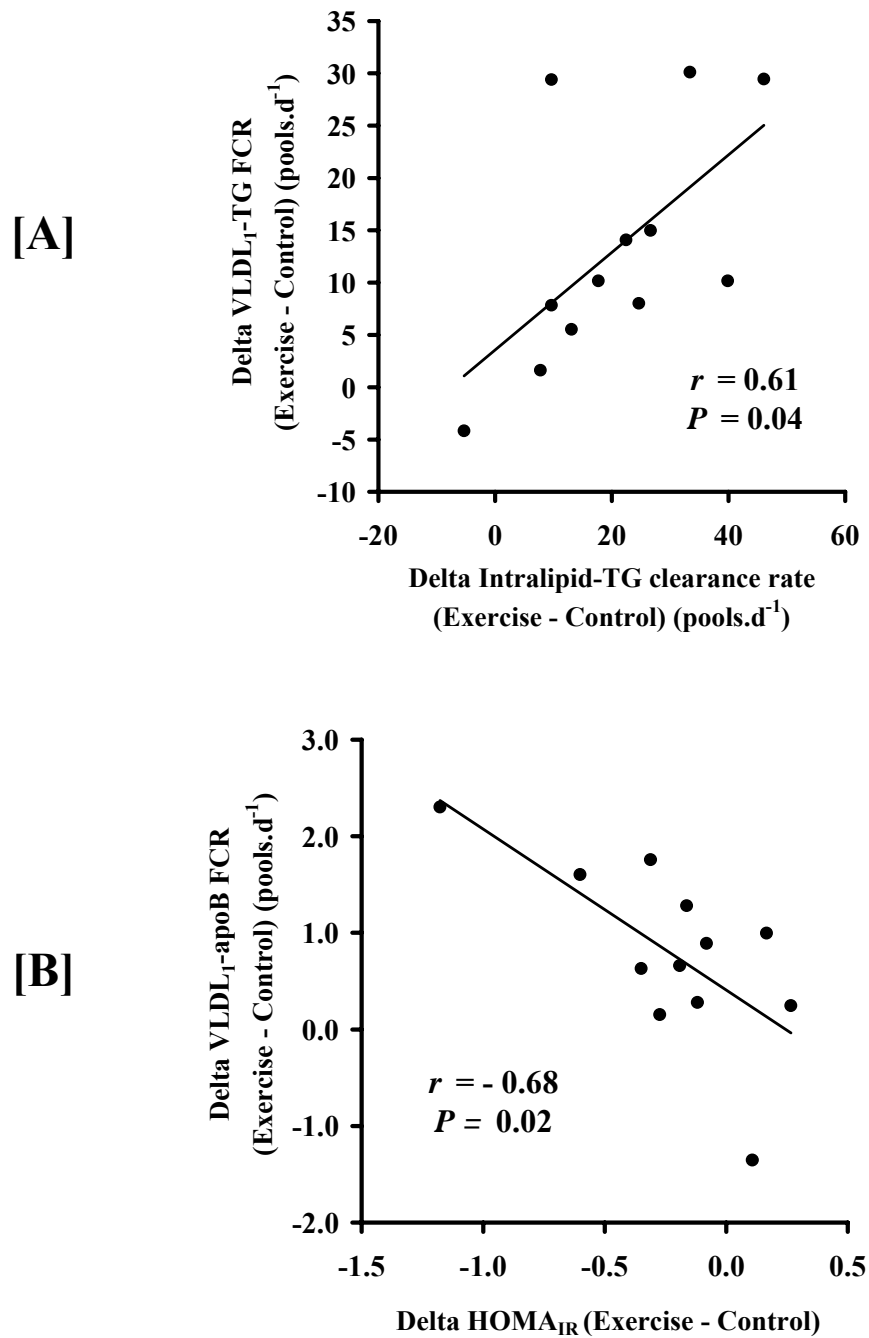


Figure 5.3: Scattergrams indicating the relationships between the exercise-induced changes (Exercise minus Control) in [A] Intralipid-TG and VLDL<sub>1</sub>-TG FCR (pools.d<sup>-1</sup>) and [B] VLDL<sub>1</sub>-apoB FCR (pools.d<sup>-1</sup>) and HOMA<sub>1R</sub>.

## 5.4 Discussion

To the best of the author's knowledge, this is the first study to investigate the possible mechanism(s) underlying the hypotriglyceridaemic effect of a prior session of moderate exercise (120-min brisk walk at  $\sim 50\%$   $\dot{V}O_{2\max}$ ) in overweight/obese middle-aged men, a group typically targeted in exercise-for-health guidelines.

The results of the present study confirm and expand on earlier observations that a prior session of moderate exercise induces a significant reduction in plasma TG and influences VLDL<sub>1</sub> to a greater extent than chylomicrons (Gill *et al.*, 2006). Indeed, in the present study, the change in VLDL<sub>1</sub>-TG clearance was greater than the change in Intralipid change (91.0% *vs.* 43.3%, respectively). Because chylomicrons are preferred substrates for LPL (Fisher *et al.*, 1995) and, therefore, compete with VLDL<sub>1</sub> particles for a LPL-mediated clearance (Karpe *et al.*, 1993b), it was hypothesised earlier that the exercise-induced TG reduction was likely to be the result of a reduced hepatic VLDL<sub>1</sub> production rather than increased clearance (Malkova & Gill, 2006). However, this does not seem to be the case. The present results demonstrate that this exercise-induced reduction in TG is due to a significant 91% increase in VLDL<sub>1</sub>-TG FCR (i.e. clearance), with no significant effect on VLDL<sub>1</sub>-TG production rate. This is consistent with two recent stable-isotope studies by Magkos *et al.* (Magkos *et al.*, 2006) and Tsekouras *et al.* (Tsekouras *et al.*, 2007) who reported an increased total VLDL-TG clearance, rather than decreased VLDL-TG production following a session of 120-min cycling and 90-min walking (at 60%  $\dot{V}O_{2\max}$ ), respectively.

The novelty of this study, however, is that it allowed the simultaneous measurements of VLDL<sub>1</sub>-TG FCR and Intralipid-TG clearance rate in response to a prior session of moderate exercise. This is important in providing a better understanding of the major possible mechanism(s) involved in the exercise-induced hypotriglyceridaemia as it was difficult to measure the clearance of VLDL<sub>1</sub>-TG in the presence of chylomicrons/Intralipid because of their competition for the same catabolic pathway (Bjorkegren *et al.*, 1996; Karpe & Hultin, 1995). Importantly, the present results show a significant correlation ( $r = 0.61$ ,  $P = 0.035$ ) between the exercise-induced

changes in Intralipid-TG and in VLDL<sub>1</sub>-TG indicating that exercise is upregulating their clearance by the same mechanism/pathway (i.e. LPL). However, it is unclear why VLDL<sub>1</sub> appears to develop a higher affinity following exercise for clearance by LPL relative to Intralipid, but this could be due to the compositional changes in the VLDL<sub>1</sub> particle post-exercise: the particles in the circulation tended to be ~26% bigger following exercise and more TG-enriched, which may render them more favourable substrates for LPL-mediated lipolysis (Fisher *et al.*, 1995). There are two possible explanations as to why this might be the case. Firstly, the liver might be producing a bigger, more TG-enriched VLDL<sub>1</sub> particle following exercise. Consistent with this, circulating NEFA concentrations were 16% higher in the exercise trial than the Control trial, increasing fatty acid availability to the liver, which could lead to larger secreted VLDL<sub>1</sub> particle. Secondly, under normal conditions, once in the circulation, the TG content of the newly secreted VLDL<sub>1</sub> particle, and consequently its size, are being reduced rapidly by hydrolysis by LPL (Karpe *et al.*, 2007) and the action of CETP, which promotes the transfer of TG from apoB-containing particles (VLDL and LDL) in exchange for CE from HDL (Barter, 2002). The present results show a significant ~29% reduction in VLDL<sub>1</sub> CE/TG ratio in the Exercise trial, which is supported by a similar 18% post-exercise reduction in CE/TG ratio reported by Gill *et al.* (Gill *et al.*, 2006). This may be suggestive of a reduced *in vivo* CETP activity following moderate exercise, rendering the VLDL<sub>1</sub> particle more TG-enriched. However, this explanation is less likely to occur as the CE/apoB ratio, representing the CE content per VLDL<sub>1</sub> particle, in the present study was not significantly different between the two trials (Control:  $2487 \pm 353$ , Exercise:  $2276 \pm 564$ ,  $P = 0.537$ ). Unfortunately, conclusive results about the effect of moderate exercise on CETP activity or mass are currently lacking in the literature. While some studies found an increase in CETP mass following exercise (Thomas *et al.*, 2001), others reported an exercise-induced decrease in CETP mass (Magkos *et al.*, 2006). Two other studies found no significant difference in CETP activity post-moderate exercise (Thomas *et al.*, 2004) even after 24 h (Zhang *et al.*, 2002).

Furthermore, although there was no significant difference in the number of apoC-II, C-III and E per VLDL<sub>1</sub> between the Exercise and Control trials in this study, a reduced apoC-III and apoE enrichment of VLDL<sub>1</sub> was reported in a similar group of

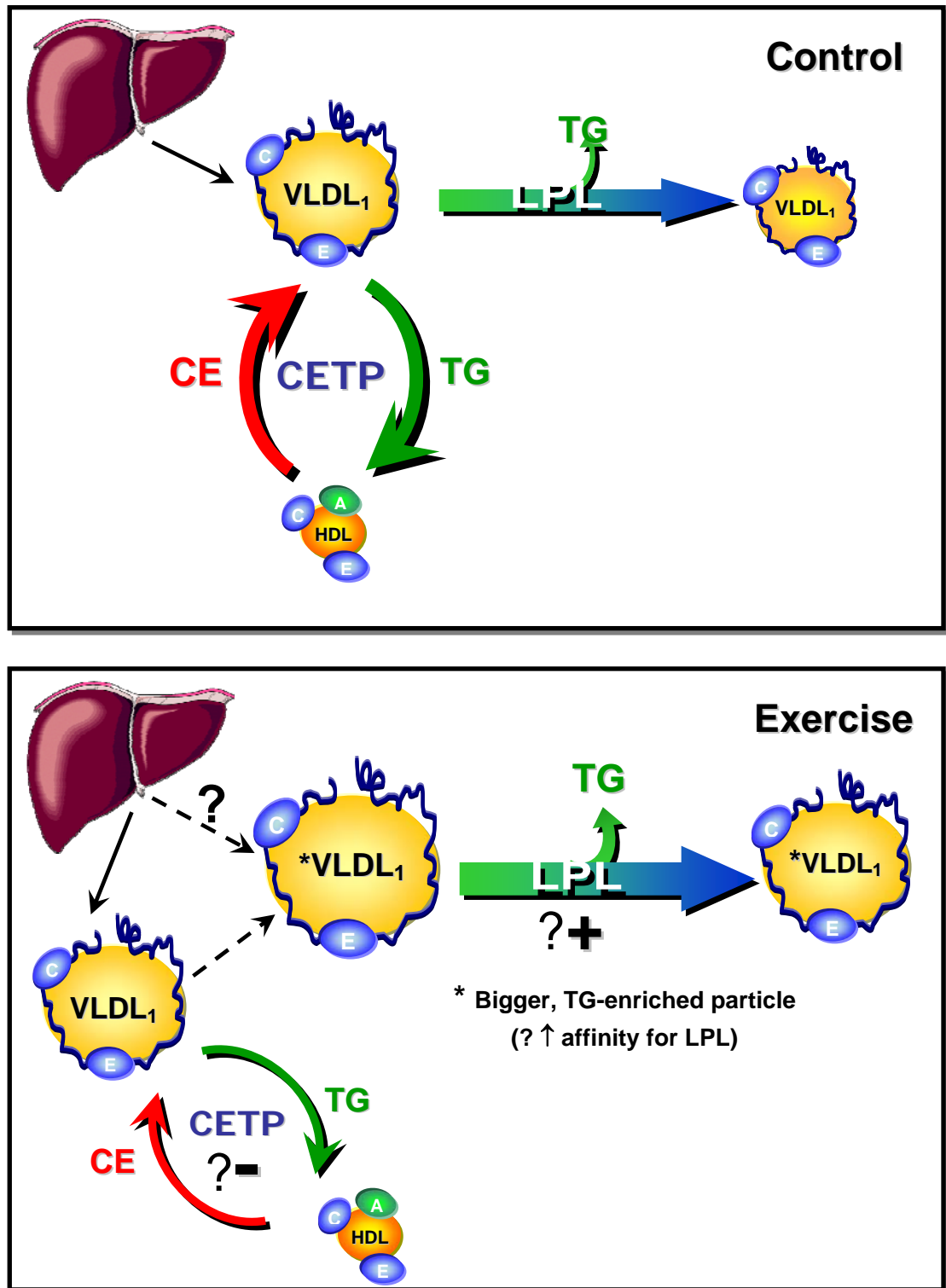
participants following moderate exercise (Gill *et al.*, 2006). These apolipoproteins are acquired by VLDL from circulating HDL and apoC-III (Wang *et al.*, 1985) and apo E (Jong *et al.*, 1997) both act to inhibit lipolysis by LPL. This could conceivably have increased the potential of post-exercise VLDL<sub>1</sub> particle as an LPL substrate. It is unclear why the present findings differ from the earlier report.

The present results also show a significant 43% increase in Intralipid-TG clearance rate post-exercise, which is likely to reflect increased LPL activity and/or mass. Although early studies of prolonged and vigorous exercise, such as marathon running, clearly demonstrated that this type of exercise increased LPL-mediated TG-clearance (Sady *et al.*, 1986), recent studies of a more moderate exercise are conflicting. While some studies show no significant difference in muscle (Herd *et al.*, 2001) or post-heparin plasma LPL activity (reflecting overall LPL activity from all body tissues) ~16h after a moderate session of exercise (Gill *et al.*, 2003), others reported a significant increase in plasma (Zhang *et al.*, 2002), muscle and adipose tissue LPL activity post-exercise (Perreault *et al.*, 2004). In addition, although Magkos *et al.* (Magkos *et al.*, 2006) found no significant difference in muscle LPL mass, the group reported a significant ~20% increase in plasma LPL concentrations in response to acute moderate exercise. It should be noted, though, that even in the absence of a significant increase in post-heparin LPL activity, the exercise-induced changes in LPL activity was reported to correlate significantly with the exercise-induced changes in fasting and postprandial TG (Gill *et al.*, 2003). Furthermore, if exercise increased the affinity of VLDL<sub>1</sub> for clearance, this could have occurred without an increase in measured LPL action using a standard substrate. This further highlights the importance of LPL as a mediator of the TG-lowering effect of moderate exercise.

In addition to increased VLDL<sub>1</sub>-TG FCR, exercise significantly increased VLDL<sub>1</sub>-apoB FCR by 211% with no significant influence on VLDL<sub>1</sub>-apoB production rate. This is in contrast with Magkos *et al.* (Magkos *et al.*, 2006) who reported a significant reduction in total VLDL-apoB production rate with no significant change in VLDL-apoB clearance. A likely explanation for these different findings could be due to the different characteristics of participants. Unlike the normal weight subjects who participated in Magkos *et al.* study (Magkos *et al.*, 2006), the participants in the

present study were overweight/obese men with increased abdominal obesity, which makes them likely candidates for insulin resistance and high liver fat content (Karpe & Tan, 2005; Haffner, 2007; Després, 2007; Adiels *et al.*, 2006b). Similar group of subjects with increased liver fat failed to suppress VLDL<sub>1</sub> production in response to insulin (Adiels *et al.*, 2007). Indeed, the participants in the present study had higher insulin concentrations compared with the study subjects of Magkos *et al.* (Magkos *et al.*, 2006) (7.0 vs. 5.4 mU.l<sup>-1</sup>, respectively) and no significant reductions in insulin or HOMA<sub>IR</sub> concentrations were observed following exercise in the present study. Interestingly, however, there was a significant correlation in the present study between the exercise-induced change in HOMA<sub>IR</sub> and the exercise-induced change in VLDL<sub>1</sub>-apoB FCR, indicating that subjects who had a bigger decrease in HOMA<sub>IR</sub> following exercise had a bigger increase in VLDL<sub>1</sub>-apoB FCR (clearance). These exercise-induced changes in insulin sensitivity may augment changes in VLDL<sub>1</sub>-apoB clearance, but this relationship was not observed for VLDL<sub>1</sub>-TG.

In summary, a single session of moderate intensity exercise significantly reduced fasting plasma TG, mainly in the VLDL<sub>1</sub> fraction, by increasing the VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB FCR (i.e. clearance). The increase in VLDL<sub>1</sub> was greater than the increase in Intralipid clearance, suggesting that exercise may have increased the affinity of VLDL<sub>1</sub> for LPL-mediated lipolysis (**Figure 5.4**). In contrast, exercise did not decrease hepatic VLDL<sub>1</sub>-TG or VLDL<sub>1</sub>-apoB production. However, further studies are needed elucidate the mechanism(s) underlying this increased affinity for clearance of VLDL<sub>1</sub>, such as *in vitro* studies investigating the affinity of VLDL<sub>1</sub> for LPL as well as the change in CETP activity both under control and exercise conditions. This may help improve understanding of the nature of these exercise-induced changes.



**Figure 5.4: Possible mechanisms involved in moderate exercise-induced reduction in VLDL<sub>1</sub>.** **Control:** once in the circulation, the TG content of the newly secreted VLDL<sub>1</sub> particle from the liver, and consequently its size, are being reduced rapidly by hydrolysis by lipoprotein lipase (LPL) and the action of cholesteryl ester transfer protein (CETP), which promotes the transfer of TG from VLDL (and LDL) in exchange for cholesteryl ester (CE) from HDL. **Exercise:** A prior session of moderate exercise significantly reduces VLDL<sub>1</sub> concentration by increasing its clearance from plasma, possibly by compositional changes to the VLDL<sub>1</sub> particle which render it bigger in size and more TG-enriched, thereby increase its affinity for LPL-mediated lipolysis. The bigger particles are either produced directly by the liver and/or reduced CETP activity. Also, increased LPL activity is likely to play a role.

## 6. General Discussion

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### 6.1 Summary

There is a growing body of evidence implicating elevated VLDL concentrations, particularly the larger VLDL<sub>1</sub> (S<sub>f</sub> 60-200) particles, as the primary cause of the atherogenic lipoprotein phenotype in insulin resistance conditions; such as type 2 diabetes (Taskinen, 2003), metabolic syndrome (Grundy, 2006), obesity (Aguilera *et al.*, 2008) and, eventually, coronary heart disease (Abbasi *et al.*, 2002; Tanaka *et al.*, 2001). The fact that this atherogenic dyslipidaemia is detectable years before the onset of the disease (Tilly-Kiesi *et al.*, 1996; Johanson *et al.*, 2004) makes the measurement and regulation of hepatic VLDL<sub>1</sub> production of great importance. However, mere measurements of VLDL<sub>1</sub> concentrations provide limited information and do not reflect the dynamics of the continual synthesis and clearance of VLDL<sub>1</sub> particles. Radioactive and stable-isotope tracers have been successfully used to measure the kinetics of different lipoproteins, including VLDL<sub>1</sub>. Although such methods yield a large amount of useful information, they are expensive, labour-intensive, time consuming and require the use of specialised laboratories, equipment and skills in multicompartmental modelling. Of note, these methods measure generally either total VLDL-TG and -apoB, or VLDL<sub>1</sub>- and VLDL<sub>2</sub>-apoB, but not VLDL<sub>1</sub>- or VLDL<sub>2</sub>-TG. It is only recently that Adiels and colleagues (Adiels *et al.*, 2005a) have developed a multicompartmental stable-isotope method to determine both VLDL<sub>1</sub>- and VLDL<sub>2</sub>-TG and -apoB kinetics. Thus, this thesis was designed to develop and validate a simpler and cost-effective method – the ‘Intralipid Method’ – to determine VLDL<sub>1</sub> kinetics and use this method to study the effects of insulin and moderate exercise on the regulation of hepatic VLDL<sub>1</sub> kinetics.

The method builds on the work of Björkegren *et al.* (Björkegren *et al.*, 1996) and relies on the fact that Intralipid (chylomicron-like particles) compete with VLDL<sub>1</sub> for the same catalytic pathway (Björkegren *et al.*, 1996; Karpe & Hultin, 1995), which allows the accumulation of VLDL<sub>1</sub> in the circulation (Björkegren *et al.*, 1996) during a constant Intralipid infusion. By measuring the linear rise in VLDL<sub>1</sub>-TG and -apoB concentrations, it is possible to determine their production rates. In addition, because

the method is employed in the fasted state in which there is a steady state where production equals clearance, it is also possible to determine the fractional catabolic rate (FCR, i.e. clearance) of VLDL<sub>1</sub>-TG and -apoB by dividing their linear rises during the Intralipid infusion by their fasting concentrations. Furthermore, the Intralipid method combines the findings of Rössner (Rossner, 1974), who showed that Intralipid is cleared following first-order kinetics, which allows the determination of Intralipid-TG clearance rate as a surrogate measure of chylomicron-TG clearance.

In addition to the fact that the kinetic data obtained using the Intralipid method are similar to that obtained by the ‘gold standard’ stable-isotope method (Bjorkegren *et al.*, 1996), this newly developed method has a number of strengths. First, it provides valuable information about TRL kinetics: both VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB production rates *and* Intralipid-TG clearance rate. In addition, under steady state conditions, it allows the simultaneous determination of large VLDL<sub>1</sub>-TG (rather than total VLDL-TG) and Intralipid-TG clearance rates. This is important for two reasons: (1) it is the high concentrations of large VLDL<sub>1</sub> particles that are implicated in atherogenic dyslipidaemia and the progression of atherosclerosis (Taskinen, 2003; Packard, 2003) and, unlike VLDL<sub>2</sub>, VLDL<sub>1</sub> are the target for insulin-induced hepatic suppression (Gill *et al.*, 2004a); (2) due to the fact that Intralipid/chylomicrons and VLDL<sub>1</sub> compete for the same LPL-mediated catalytic pathway, it has previously been difficult to determine the clearance rate of VLDL<sub>1</sub> in the presence of Intralipid or chylomicrons. As a result, the Intralipid method helped to elucidate the hypotryglyceridaemic effects of exercise on TRL kinetics ([Chapter 5](#)). Second, the Intralipid method is sensitive enough to detect physiological changes in hepatic VLDL<sub>1</sub> production and Intralipid-TG clearance, such as the effect of insulin on TRL kinetics ([Chapter 4](#)). Third, the method is relatively easy and simple enough to be used in any laboratory equipped with only one ultracentrifuge without the need for any other specialised equipment (such as a mass spectrometer). Fourth, estimation of production and clearance rates using the Intralipid method is relatively easy to carry out and can be automated using Excel. Fifth, the protocol for the Intralipid method takes less time to complete than the stable-isotope method, which is beneficial both for the subject and the researcher. While a standard stable-isotope method to measure VLDL<sub>1</sub> kinetics requires ~48 h of the subject’s time, the Intralipid infusion



and post-infusion period of the present method lasts only for about 3.5 h. In addition, unlike the time required (at least 2 weeks) for laboratory techniques and sample and kinetic analyses using the stable-isotope method, the Intralipid method protocol, including lipoprotein separation, sample analysis and kinetic calculations, takes approximately 2-3 days. Sixth, the Intralipid method is cheaper than the stable-isotope method and suitable for laboratories with limited funding or resources as each trial (including investigator/technician time at £20 per hour) costs approximately £750 and 2-3 days to complete compared to the ~£2500 and about two weeks using the stable-isotope method. Finally, because the Intralipid method is safe as well as both time- and cost-effective, it can be used for a large number of subjects, with even multiple trials for each subject.

As a first application, the Intralipid method was used to study the effect of hyperinsulinaemia and hyperglycaemia on VLDL<sub>1</sub>-TG and -apoB production rates in 8 normoglycaemic subjects ([Chapter 4](#)). This was done using a simple frequent oral administration of glucose to provide a pseudo-steady hyperinsulinaemic state as the use of a conventional hyperinsulinaemic normoglycaemic clamp was not possible due to lack of constant medical cover. The result demonstrated that insulin acutely suppressed VLDL<sub>1</sub>-TG and -apoB production, a finding recently supported by Adiels *et al.* (Adiels *et al.*, 2007). Although hyperinsulinaemia had no apparent effect on Intralipid-TG clearance rate, the change seemed to be related to insulin sensitivity, i.e. the more insulin sensitive the subject was, the bigger the increase in Intralipid-TG clearance rate. One possible limitation of this study is the number of subjects, especially with such a wide range of insulin resistance, which may have diluted the effect of insulin on Intralipid-TG clearance rate. However, the technique would make it very easy to increase the numbers in future studies.

Because moderate exercise has been shown to decrease plasma TG by 20-25%, mostly in the VLDL<sub>1</sub> fraction (Gill *et al.*, 2006), the second application of the Intralipid method was to investigate whether this decrease is due to decreased hepatic VLDL<sub>1</sub> production, increased clearance or a combination of both ([Chapter 5](#)). The present results showed that a session of moderate exercise significantly increased the clearance of VLDL<sub>1</sub>-TG and -apoB rather than suppress their hepatic production. Although there is no information in the literature about the effect of exercise on

VLDL<sub>1</sub> kinetics specifically, these results are consistent with recent findings using stable-isotopes which investigated the effect of moderate exercise on total VLDL-TG (Magkos *et al.*, 2006; Tsekouras *et al.*, 2007) and VLDL-apoB (Magkos *et al.*, 2006) using the stable-isotope method. In addition, because of the nature of the Intralipid method, it provided the opportunity to simultaneously study both Intralipid and VLDL<sub>1</sub> clearance, which was previously unfeasible. As a result, a novel finding is observed that this increased exercise-induced clearance of VLDL<sub>1</sub> is likely to be the result of increased affinity of VLDL<sub>1</sub> to LPL-mediated lipolysis. Furthermore, this study is the first to investigate the kinetics of such effects of exercise in overweight/obese middle-aged men, a group typically targeted in exercise-for-health guidelines.

Adequate liver function is important for VLDL<sub>1</sub> homeostasis. Serum ALT concentrations are used clinically as an indication of liver function and, within normal range, have been shown to be a marker for liver fat (Westerbacka *et al.*, 2004). It was recently shown that hepatic overproduction of VLDL<sub>1</sub> particles is driven by liver fat accumulation (Adiels *et al.*, 2006b). Results from this thesis show a significant relationship between serum ALT concentrations and hepatic VLDL<sub>1</sub> production and regulation. Even within normal range, ALT concentrations correlated significantly with VLDL<sub>1</sub> production ([Chapter 3](#)) and inversely with the extent to which VLDL<sub>1</sub> is suppressed in response to hyperinsulinaemia and hyperglycaemia ([Chapter 4](#)): subjects who had higher ALT concentrations, produced more VLDL<sub>1</sub> particles and failed to suppress VLDL<sub>1</sub> secretion in response to insulin. It is unclear, however, whether this correlation with ALT concentrations reflects liver function *per se* or hepatic fat content, as ALT is a marker for both.

In all studies described in this thesis, 3 subjects were E2/3 phenotype, 11 subjects were E3/3 phenotype, 6 subjects were E3/4 phenotype and 3 subjects were E4/4 phenotype. Previous studies showed that apoE phenotypes influence plasma lipids (Eto *et al.*, 1986) and particularly large VLDL<sub>1</sub> concentrations (Bioletto *et al.*, 1998) as well as VLDL-apoB kinetics (Watts *et al.*, 2000; Welty *et al.*, 2000; Demant *et al.*, 1991). However, no apparent effect of apoE phenotype on VLDL<sub>1</sub>-TG and -apoB production rates or Intralipid-TG clearance rate was observed in this thesis (Appendix C). It is uncertain, however, whether this is a true lack of influence of

apoE on VLDL<sub>1</sub>-TG and -apoB production rates and Intralipid-TG clearance rate or whether it may be due to lack of sufficient numbers for each phenotype. Fortunately, because of the relative ease and cost-effectiveness of the Intralipid method, a bigger study can be conducted which may have not been feasible otherwise using the stable-isotope method.

## 6.2 Limitations of the Intralipid Method

There are two main limitations to the Intralipid method. The first is that it only allows the determination of VLDL<sub>1</sub> kinetics but not VLDL<sub>2</sub>. This is because VLDL<sub>2</sub> is hydrolysed by HL; i.e. Intralipid does not compete with VLDL<sub>2</sub> for LPL-mediated lipolysis and, consequently, VLDL<sub>2</sub> will not accumulate in the circulation like VLDL<sub>1</sub> particles. Although VLDL<sub>1</sub> and VLDL<sub>2</sub> are regulated independently (Packard & Shepherd, 1997; Gill *et al.*, 2004a), it has been recently reported that while insulin acutely reduced VLDL<sub>1</sub> secretion, it increased VLDL<sub>2</sub> secretion (Adiels *et al.*, 2007). Although this acute suppression of VLDL<sub>1</sub> secretion by insulin was detected using the Intralipid method, it was not possible to investigate this effect on VLDL<sub>2</sub>. Thus, the addition of VLDL<sub>2</sub> kinetics might have helped to complete the picture of the effect of insulin on total VLDL hepatic regulation. The second limitation of the Intralipid method is that the FCR can only be calculated if the subject is in a steady state. For example, it was not possible to determine VLDL<sub>1</sub>-TG and VLDL<sub>1</sub>-apoB FCRs in [Chapter 4](#) where VLDL<sub>1</sub> production was investigated in response to hyperinsulinemia and hyperglycemia. This is because the steady state value used for FCR calculation is made in the fasted state before the glucose ingestion, which does not represent the steady state value for VLDL<sub>1</sub> concentration during glucose ingestion.

Furthermore, the lack of direct comparison with either the stable-isotope or radioactive tracer methods within the same subjects represents a limitation to the validation of the method. Although Björkegren *et al.* (1996) reported a close agreement between the two methods, ideally, further studies on a larger scale should be performed in the future to validate the Intralipid method against the currently used 'gold-standard' methods.

### 6.3 Future Experiments and Applications of the Method

Results from [Chapter 5](#) suggest that moderate exercise is likely to increase VLDL<sub>1</sub> clearance by increasing its size and TG-enrichment, thereby increasing its affinity for LPL-mediated lipolysis. Reports from the literature with respect to the effect of moderate exercise on LPL and CETP activities, which are responsible for VLDL<sub>1</sub> TG-enrichment, are conflicting. Therefore, it would be helpful to conduct the following two studies. The first is an *in vitro* study investigating the suggested higher affinity of VLDL<sub>1</sub> to LPL post-exercise. This could possibly be done by incubating LPL (obtained commercially or post heparin from a donor) with different concentrations (and possibly at different durations) of the subject's native VLDL<sub>1</sub> particles separated from the Control and Exercise trials. VLDL<sub>1</sub> lipolysis is then determined by measuring the released NEFA from VLDL<sub>1</sub> by LPL over time. In addition, Intralipid could also be incubated with native VLDL<sub>1</sub> and LPL to investigate the relative competition between Intralipid and VLDL<sub>1</sub> particles for LPL-mediated lipolysis in both the Control and Exercise trials.

Secondly, it is important to compare the activity of CETP in control and post-exercise as data from [Chapter 5](#) suggest an increased TG-enrichment of VLDL<sub>1</sub> particles post-exercise. However, current commercial methods that measure CETP activity use the same concentration for donor and acceptor particles in both exercise and control trials, which does not necessarily reflect physiological conditions as *in vivo* concentrations of VLDL particles differ between both conditions. Therefore, it would be better to measure CETP activity in an assay in which native VLDL<sub>1</sub> from the Exercise and Control trials are used as donor and acceptor particles.

These two experiments would aid a better understanding of the effect of exercise on VLDL<sub>1</sub> and test the hypothesis that the increased clearance of VLDL<sub>1</sub> post-exercise is due to the increased size and TG-enrichment of the VLDL<sub>1</sub> particle.

Finally, previous studies have shown that the relationships between obesity, insulin resistance, metabolic syndrome and risk for cardiovascular disease are subject to ethnic variations (Joshi *et al.*, 2007; Lear *et al.*, 2003; Whincup *et al.*, 2002). For example, compared with Caucasian men, South Asian men have excess subcutaneous abdominal adiposity, in spite of similar BMI, which renders them more susceptible to

insulin resistance than white men (Chandalia *et al.*, 2007). In addition, although Afro-Caribbean men in the United Kingdom are reported to be as insulin-resistant as South Asian men, they have been found to be less susceptible to the dyslipidaemia that accompanies insulin resistance (Zoratti *et al.*, 2000). Interestingly, Zoratti *et al.* (2000) reported that this favourable lipoprotein profile in Afro-Caribbean may relate to an effective VLDL metabolism. However, there is currently no information available on the effect of ethnicity on VLDL<sub>1</sub> kinetics. Therefore, an important application of the Intralipid method would be to investigate VLDL<sub>1</sub> kinetics in a different ethnic population. In addition, the method could be used to study VLDL<sub>1</sub> kinetics in response to moderate exercise in Kuwaiti subjects and compare that with results from those obtained in Scottish men. Kuwait (the author's homeland) has a high prevalence of obesity (Al-Kandari, 2006), insulin resistance (Al-Shaibani *et al.*, 2004) and type 2 diabetes (Al-Adsani *et al.*, 2004) and ischemic heart diseases are a major cause of death in the country (El-Shazly *et al.*, 2004).

## **6.4 Conclusion**

In conclusion, this thesis describes the development and validation a novel method to determine the kinetics of large VLDL<sub>1</sub>, and the application of this method to investigate the effects of hyperinsulinaemia/hyperglycaemia and moderate exercise on VLDL<sub>1</sub> kinetics. The method is relatively straightforward and cost-effective which means it has potential for future widespread use in situations where isotopic tracer methods are not feasible because of technical or cost limitations.

## 7. References

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## 8. Appendices

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Appendix A1: Subject information sheet and consent form – Chapters 3 and 4

Appendix A2: Subject information sheet and consent form – Chapter 5

Appendix B1: Health screen questionnaire – Chapters 3 & 4

Appendix B2: Health screen questionnaire – Chapter 5

Appendix C: Kinetic results for all subjects according to apoE phenotypes



**Appendix A1: Subject Information Sheet and Consent Form –  
Chapters 3 & 4**

Version 3, 18<sup>th</sup> March 2004



**VOLUNTEER INFORMATION SHEET**

**Title: Development of novel method to determine very low  
density lipoprotein kinetics**

**Lay title: A new method of measuring the metabolism of blood fats**

You are being invited to take part in a research study. Before you decide it is important for you to understand why the research is being done and what it will involve. Please take time to read the following information carefully and discuss it with others if you wish. Ask us if there is anything that is not clear or if you would like more information. Take time to decide whether or not you wish to take part.

*Thank you for reading this.*



*Please reply to :* Division of Neuroscience and Biomedical Systems,  
Institute of Biomedical and Life Sciences,  
University of Glasgow, West Medical Building,  
University Avenue, Glasgow G12 8QQ  
*Tel:* 0141-330-2916 *Fax:* 0141-330-2915 *E-mail:* j.gill@bio.gla.ac.uk

## *Frequently Asked Questions*

### *What is the purpose of the study?*

The aim of this study is to develop a new method of measuring how fast your liver produces fatty particles (called lipoproteins) and how fast these particles are cleared from your bloodstream. This will help us to determine how effective different treatments are at influencing fat metabolism in future studies.

### *Why have I been chosen?*

You have been chosen because you are a healthy adult aged between 18 and 65 years.

### *Do I have to take part?*

It is up to you to decide whether or not to take part. If you do decide to take part you will be given this information sheet to keep and be asked to sign a consent form. If you decide to take part you are still free to withdraw at any time and without giving a reason. A decision to withdraw at any time, or a decision not to take part, will not affect the standard of care you receive.

### *What will happen to me if I take part?*

In the first instance you will be asked to attend for a screening visit in which we will discuss with you and complete confidential questionnaires regarding your health and family history, measure your blood pressure and provide an opportunity for you to ask questions about the study. We will then ask you to help us in one of two different ways:

***To provide a fasting blood sample:*** We will ask you to come to the Clinical Investigation Suite after an overnight fast and provide a simple blood sample. We will take up to 60 ml (4 tablespoons) of blood from you. Your blood will be used to help us to develop laboratory methods for analysing the fat in your bloodstream.

**OR**

***To help us to develop a new method for assessing the metabolism of blood fats:*** We will ask you to perform either one, two or three experimental trials to help us to develop a new method for assessing the metabolism of lipoproteins. Each trial will involve coming to our Clinical Investigation Suite after an overnight fast. We will then introduce two small plastic tubes called cannulas into a vein in your arms (one in each arm). This is no more painful than giving a normal blood sample. We will take blood from one cannula and use the other to pump small amounts of a fatty substance called Intralipid® into your blood stream. Intralipid® is used in hospitals to feed people through a 'drip' when they are unable to eat normally. The Intralipid® we give you will cause the level of fat in your blood to rise about the same amount as it does after you eat a fatty meal and it will be pumped into your bloodstream for between one and three hours. We will take two blood samples from the cannula before giving you the Intralipid® and then take blood samples at intervals during the Intralipid® infusion and for a maximum of two hours after the Intralipid® pump is stopped. We may ask you to sip small amounts of a glucose

drink throughout the trial. We will take a maximum of 250 ml of blood during each trial – this is half the amount of blood that you give when you donate a ‘pint’ of blood. During the experimental trial you will only be able to drink water (or the glucose drink), but we will provide you with some food to eat when the trial is finished. The total duration of each trial will vary between 2½ and 6 hours but we will tell you specifically how long each of your own trials will take. You will be able to watch TV or videos, listen to music or read to keep yourself occupied during the experiment.

### *What do I have to do?*

We will ask you to weigh and record everything that you eat and drink and to avoid alcohol and planned exercise for the two days before each Intralipid® test. We will provide you with scales and record sheets to enable you to record your diet easily. If you do more than one test, we will ask you to eat the same diet on the two days leading up to each test. (You will not need to do this if you are just giving us a fasting blood sample).

### *What are the possible disadvantages and risks of taking part?*

Blood sampling via the cannula may cause minor bruising, an inflammation of the vein or haematoma (a small accumulation of blood under the skin). Good practice, however, minimises this risk. Some people may feel faint when they give blood. There is a small risk that you could have an allergic reaction to Intralipid® (less than one chance in a million). This usually only occurs in people are allergic to soya protein or eggs, so if you have these allergies you will not be able to participate in this study. In the unlikely event that you do have an allergic reaction to Intralipid®, the appropriate medicines and equipment are in place for you to be treated immediately.

There is a small possibility that taking part in this study will reveal a health problem that you already have such as high cholesterol or high blood pressure. If such a problem is revealed, we will inform your GP to ensure that you receive appropriate treatment.

### *What are the possible benefits of taking part?*

The purpose of this study is to develop a new method of determining how fast lipoprotein particles are produced by the liver and how fast they are broken down and cleared from the bloodstream. We will be able to use this method in future research to help to determine how effective different treatments are at improving the way in which the body handles fat.

### *What if something goes wrong?*

The chances of something going wrong are extremely small. All of the procedures involved in this study are low risk and our screening tests are designed to ensure that you will only participate if it is safe for you to do so. In the unlikely event that you are harmed due to someone's negligence, then you may have grounds for a legal action but you may have to pay for it. Regardless of this, if you wish to complain, or have any concerns about any aspect of the way you have been approached or treated during the course of this study, the normal National Health Service complaints mechanisms may be available to you.

***Will my taking part in this study be kept confidential?***

All information which is collected about you during the course of the research will be kept strictly confidential. Any information about you which leaves the University or hospital will have your name and address removed so that you cannot be recognised from it.

***Who has reviewed the study?***

This study has been reviewed and approved by the North Glasgow NHS Trust Research Ethics Committee.

***Contact for Further Information***

Any questions about the procedures used in this study are encouraged. If you have any doubts or questions, please ask for further explanations by contacting either:

**Iqbal AlShayji**

Mobile: 07799353689

Tel: 0141 2114595

E-mail: [ialshayji@yahoo.com](mailto:ialshayji@yahoo.com)

**Dr Jason Gill**

Tel: 0141 3302916

E-mail: [j.gill@bio.gla.ac.uk](mailto:j.gill@bio.gla.ac.uk)

*You will be given a copy of this information sheet and a signed consent form to keep for your records.*

Volunteer Identification Number for this trial: \_\_\_\_\_



## CONSENT FORM

**Title of Project: Development of novel method to determine very low density lipoprotein kinetics**

**Lay title: A new method of measuring the metabolism of blood fats**

Name of Researcher: \_\_\_\_\_

**Please initial box**

1. I confirm that I have read and understand the information sheet dated 18 March 2004 (version 3) for the above study and have had the opportunity to ask questions. ☐
2. I understand that my participation is voluntary and that I am free to withdraw at any time, without giving any reason, without my medical care or legal rights being affected. ☐
3. I agree to take part in the above study. ☐

_____ Name of Patient	_____ Date	_____ Signature
_____ Name of Person taking consent (if different from researcher)	_____ Date	_____ Signature
_____ Researcher	_____ Date	_____ Signature

*Blue for patient*  
*Pink for researcher*

## **Appendix A2: Subject Information Sheet and Consent Form - Chapter 5**



### **VOLUNTEER INFORMATION SHEET**

#### **Title: Effects of Moderate Exercise on Very Low Density Lipoprotein Kinetics**

**Lay title: How does exercise affect the metabolism of the liver's fatty  
particles?**

You are being invited to take part in a research study. Before you decide it is important for you to understand why the research is being done and what it will involve. Please take time to read the following information carefully and discuss it with others if you wish. Ask us if there is anything that is not clear or if you would like more information. Take time to decide whether or not you wish to take part.

*Thank you for reading this.*

### *What is the purpose of the study?*

Heart disease is the leading cause of death in Scotland and being even slightly overweight, particularly when fat is stored in the tummy region, increases your risk. Many factors contribute to a person's risk of heart disease, but it is thought that the ability to cope with fat contained in food plays a role. A single session of moderate exercise decreases blood fat concentrations following a meal, but is unclear whether this is due to an increase of the clearance of fatty particles by the blood, or their reduced production from the liver. This study will help to investigate the effects of a single brisk walking session lasting from 90 to 120 minutes on the production and clearance of fatty particles. This is important as exercise could be used as a treatment option, rather than drugs, to prevent and treat disturbances in blood fat metabolism.

### *Why have I been chosen?*

You have been chosen because you are a man or a postmenopausal woman aged between 30-60 years who is heavier than the ideal weight for your height.

### *Do I have to take part?*

It is up to you to decide whether or not to take part. If you do decide, you will be given this information sheet to keep and be asked to sign a consent form. If you do this you are still free to withdraw at any time and without giving a reason. A decision to withdraw at any time, or a decision not to take part, will not affect the standard of care you receive.

### *What will happen to me if I take part?*

In the first instance you will be asked to attend for a screening visit in which we will:

- ✚ discuss with you and complete confidential questionnaires regarding your health, family history and physical activity level
- ✚ measure your blood pressure
- ✚ take your height, weight and waist measurements
- ✚ take two small blood samples (20 ml or 4 teaspoons) to check the fat and sugar levels in your blood. We will also check if you are anaemic or have any abnormalities in the function of your liver, kidneys and thyroid gland
- ✚ provide an opportunity for you to ask questions

These preliminary procedures will enable us to determine whether you fall into the group of people we wish to study and will also ensure that it is perfectly safe for you to take part. In addition, we will inform your GP of your participation in the study and send them a copy of your screening results.

## *Experimental procedures*

### **A. Preliminary Exercise Test**

At the beginning of the study, an exercise test will be undertaken. This will involve walking on a motorised treadmill. If you are not used to walking on a treadmill, we will familiarise you with this before any 'real' sessions are performed. The test is designed to estimate your body's ability to use oxygen and enables us to find the correct speed and gradient for you to walk at during your treadmill walks. This will **not** require a maximal effort and the test will last for about 20 minutes. Heart rate


will be monitored and recorded throughout using a heart rate monitor and expired air will be collected at intervals using a mouthpiece and respiratory valve. For safety reasons, the test will be stopped if your heart rate exceeds 85% of your predicted maximum.


### **B. Body Composition**

The amount and distribution of your body fat will be determined by measuring body girths and by using callipers to measure skin fold thickness at four different sites (a sophisticated version of "pinch an inch"). Your height, weight and waist and hip circumferences will also be recorded. You will need to wear only underclothing for these measurements which will be made in private. These measurements only take a few minutes and can be made on the same day as other tests.

### **C. Main Trials**

We will ask you to undertake 2 Intralipid tests, approximately 1-2 weeks apart, in random order. On the day prior to one Intralipid test, a controlled exercise session of 90- to 120-minutes walk will be undertaken (Exercise test). Other than this, conditions (such as alcohol consumption, food eaten, etc) in the days leading up to each trial will be EXACTLY the same. This is explained in detail in the "What do I have to do?" section.

 **Control Test** – We will ask you to come to our Clinical Investigation Suite at Glasgow Royal Infirmary after an overnight fast (i.e. having eaten nothing and with only water to drink for 12 hours). We will then introduce two small plastic tubes called cannulas into a vein in your arms (one in each arm). This is no more painful than giving a normal blood sample. We will take blood from one cannula and use the other to pump small amounts of a fatty substance called Intralipid into your blood stream. Intralipid is used in hospitals to feed people through a 'drip' when they are unable to eat normally. This is a quick way for us to give you a fatty meal. The Intralipid we give you will cause the level of fat in your blood to rise about the same amount as it does after you eat a fatty meal and it will be pumped into your bloodstream for 60 to 75 minutes. We will take a maximum of 250 ml of blood during each trial – this is half the amount of blood that you give when you donate a 'pint' of blood. During the test you will not be able to eat or drink anything except water, but we will provide you with some food to eat when the trial is finished. The total duration of each test will be about 3 hours. You will be able to watch TV or videos, listen to music or read to keep yourself occupied during the experiment.

 **Exercise Test** – This will be identical to the control test, except that we will ask you to come to the University to walk on the treadmill for 90 to 120 minutes on the day before the test.

### ***What do I have to do?***

We also ask you to maintain your usual lifestyle (i.e. don't change your diet or exercise habits) for the duration of this study. However, before each Intralipid test, we ask you to do the following:



1. For 3 days before each Intralipid test, refrain from planned or strenuous exercise, other than for personal transportation and the 90- to 120-minute treadmill walk in our laboratory..
2. Weigh and record everything you eat and drink for 2 days before each test. We will provide you with weighing scales and diet sheets to do this.
3. Refrain from alcohol consumption on the day before each test.
4. For the 2 days leading up to the second test, we will ask you to eat the same diet as you did on the two days leading up to the first test. Therefore, we would advise you to eat meals that you will be able to easily repeat during the days preceding both tests.

### ***What are the possible disadvantages and risks of taking part?***

- ✚ Exercise testing will not be at a maximal level but the possibility exists that, very occasionally, certain changes may occur during or shortly after the tests. They include abnormal blood pressure, fainting or a change in the normal rhythm of the heartbeat.
- ✚ Blood sampling via the cannula may cause minor bruising or an inflammation of the vein. Good practice, however, minimises this risk. Some people may feel faint when they give blood.
- ✚ There is a small risk that you could have an allergic reaction to Intralipid (less than one chance in a million). This usually only occurs in people who are allergic to soya protein or eggs, so if you have these allergies you will not be able to participate in this study. In the unlikely event that you do have an allergic reaction to Intralipid, the appropriate medicines and equipment are in place for you to be treated immediately.
- ✚ There is a small possibility that taking part in this study will reveal a health problem that you already have such as high cholesterol or high blood pressure. If such a problem is revealed, we will inform your GP to ensure that you receive appropriate treatment.

### ***What are the possible benefits of taking part?***

There may be no benefits to you but as a result of being involved in this study you will receive health and fitness information about yourself including fitness tests, dietary assessment, body fat measurement and your cholesterol and blood sugar levels. This study will help us to determine how exercise can improve risk factors for heart disease and diabetes. The findings will be published in scientific journals so that understanding of the way in which exercise decreases the risk of heart disease and diabetes can be increased. This information may help make up better exercise guidelines, particularly for people who are overweight or obese.

We will provide you with feedback about the main study findings and also about your own results and would be delighted to explain results and discuss the implications with you.

### ***What if something goes wrong?***

The chances of something going wrong are extremely small. We have recently conducted a similar project and there were no problems. All of the procedures involved in this study are low risk and our screening tests are designed to ensure that you will only participate if it is safe for you to do so. In the unlikely event that you

are harmed due to someone's negligence, then you may have grounds for a legal action but you may have to pay for it. Regardless of this, if you wish to complain, or have any concerns about any aspect of the way you have been approached or treated during the course of this study, the normal National Health Service complaints mechanisms may be available to you.

### *Will my taking part in this study be kept confidential?*

All information which is collected about you during the course of the research will be kept strictly confidential. Any information about you which leaves the University or hospital will have your name and address removed so that you cannot be recognised from it. In addition, your records, samples and results will be identified by a number and not your name.

### *What will happen to the results of the research study?*

The results from this study will be presented at scientific meetings and published in scientific journals. A copy of the published results will be sent to you upon request. You will be informed which part of the study you were in, as this information will be confidential and no one else will know your name and which part you participated in.

### *What will happen to my samples after the study has finished?*

The blood samples that you provide for this study may be useful for future research into the prevention and treatment of diabetes and heart disease; this may involve investigating new biochemical markers that are not yet identified. Samples will be analysed anonymously and will require a new ethics application before they would be used for future research. If you do not wish your samples to be used for future research, please indicate this on the consent form.

### *Who has reviewed the study?*

This study has been reviewed and approved by the North Glasgow NHS Trust Research Ethics Committee.

### *Who is organising and funding the research?*

This study is being funded by the British Heart Foundation and the Glasgow Royal Infirmary R&D Endowment Fund (Chest, Heart and Stroke).

### *Contact for Further Information*

Any questions about the procedures used in this study are encouraged. If you have any doubts or questions, please ask for further explanations by contacting either:

#### **Iqbal AlShayji**

Mobile: 07799353689

Tel: 0141 211 4595 or 4596

E-mail: [i.alshayji.1@research.gla.ac.uk](mailto:i.alshayji.1@research.gla.ac.uk)

#### **Dr Jason Gill**

Tel: 0141 3302916

E-mail: [j.gill@bio.gla.ac.uk](mailto:j.gill@bio.gla.ac.uk)

*You will be given a copy of this information sheet and a signed consent form to keep for your records.*

*Thank You for Your Time and Participation*

Volunteer Identification Number: \_\_\_\_\_



## CONSENT FORM

**Title of Project: Effects of moderate exercise on very low density lipoprotein kinetics**

**Lay title: How does exercise affect the metabolism of the liver's fatty particles?**

Name of Researcher: \_\_\_\_\_

**Please initial box**

- |  |                              |
|--|------------------------------|
| 1. I confirm that I have read and understand the information sheet dated 21/02/2006 (Version 3) for the above study and have had the opportunity to ask questions.                                 | <input type="checkbox"/>     |
| 2. I understand that my participation is voluntary and that I am free to withdraw at any time, without giving any reason, without my medical care or legal rights being affected.                  | <input type="checkbox"/>     |
| 3. I agree to take part in the above study.  | <input type="checkbox"/>     |
| 4. I agree that my GP is to be informed of my participation in this study.   | <input type="checkbox"/>     |
| 5. I agree for my samples to be used for future research into the prevention and treatment of diabetes and heart disease. This may involve analysis of new biochemical markers not yet identified. | <input type="checkbox"/>     |
|  | Yes <input type="checkbox"/> |
|  | No <input type="checkbox"/>  |

\_\_\_\_\_  
Name of Subject

\_\_\_\_\_  
Date

\_\_\_\_\_  
Signature

\_\_\_\_\_  
Name of Person taking consent  
(if different from researcher)

\_\_\_\_\_  
Date

\_\_\_\_\_  
Signature

\_\_\_\_\_  
Researcher

\_\_\_\_\_  
Date

\_\_\_\_\_  
Signature

Copy for subject  
Copy for researcher

## Appendix B1: Health Screen Questionnaire – Chapters 3 & 4

### HEALTH SCREEN FOR STUDY VOLUNTEERS

Name: \_\_\_\_\_

It is important that volunteers participating in research studies are currently in good health and have had no significant medical problems in the past. This is to ensure (i) their own continuing well-being and (ii) to avoid the possibility of individual health issues confounding study outcomes.

Please complete this brief questionnaire to confirm fitness to participate:

**1. At present, do you have any health problem for which you are:**

- |  |            |           |
|--|------------|-----------|
| (a) on medication, prescribed or otherwise | yes [    ] | no [    ] |
| (b) attending your general practitioner    | yes [    ] | no [    ] |
| (c) on a hospital waiting list             | yes [    ] | no [    ] |

**2. In the past two years, have you had any illness which required you to:**

- |   |            |           |
|---|------------|-----------|
| (a) (a) consult your GP                     | yes [    ] | no [    ] |
| (b) attend a hospital outpatient department | yes [    ] | no [    ] |
| (c) be admitted to hospital                 | yes [    ] | no [    ] |

**3. Have you ever had any of the following:**

- |  |            |           |
|--|------------|-----------|
| (a) Convulsions/epilepsy                 | yes [    ] | no [    ] |
| (b) Asthma                               | yes [    ] | no [    ] |
| (c) Eczema                               | yes [    ] | no [    ] |
| (d) Diabetes                             | yes [    ] | no [    ] |
| (e) A blood disorder                     | yes [    ] | no [    ] |
| (f) Digestive problems                   | yes [    ] | no [    ] |
| (g) Hearing problems                     | yes [    ] | no [    ] |
| (h) Disturbance of balance/co-ordination | yes [    ] | no [    ] |
| (i) Numbness in hands or feet            | yes [    ] | no [    ] |
| (j) Disturbance of vision                | yes [    ] | no [    ] |
| (k) Thyroid problems                     | yes [    ] | no [    ] |
| (l) Kidney or liver problems             | yes [    ] | no [    ] |
| (m) Chest pain or heart problems         | yes [    ] | no [    ] |
| (n) Any other health problems            | yes [    ] | no [    ] |
| (o) An allergy to soya protein or eggs   | yes [    ] | no [    ] |

(a) Are you pregnant or think that you might be pregnant      yes [    ]      no [    ]

(b) Do you take the contraceptive pill or other hormone-based contraceptives  
yes [    ]      no [    ]

(c) Are you postmenopausal      yes [    ]      no [    ]

(d) Are you receiving Hormone Replacement Therapy (HRT)  
yes [    ]      no [    ]

(a) Any heart problems	yes [    ]	no [    ]
(b) Diabetes	yes [    ]	no [    ]
(c) Stroke	yes [    ]	no [    ]
(d) Any other family illnesses	yes [    ]	no [    ]

If so, for how long did you smoke and when did you stop? .....

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177

## Appendix B2: Health Screen Questionnaire – Chapter 5

### HEALTH SCREEN FOR STUDY VOLUNTEERS

Name: \_\_\_\_\_

It is important that volunteers participating in research studies are currently in good health and have had no significant medical problems in the past. This is to ensure (i) their own continuing well-being and (ii) to avoid the possibility of individual health issues confounding study outcomes.

Please complete this brief questionnaire to confirm fitness to participate:

**1. At present, do you have any health problem for which you are:**

- |  |            |           |
|--|------------|-----------|
| (a) on medication, prescribed or otherwise | yes [    ] | no [    ] |
| (b) attending your general practitioner    | yes [    ] | no [    ] |
| (c) on a hospital waiting list             | yes [    ] | no [    ] |

**2. In the past two years, have you had any illness which required you to:**

- |   |            |           |
|---|------------|-----------|
| (a) consult your GP                         | yes [    ] | no [    ] |
| (b) attend a hospital outpatient department | yes [    ] | no [    ] |
| (c) be admitted to hospital                 | yes [    ] | no [    ] |

**3. Have you ever had any of the following:**

- |  |            |           |
|--|------------|-----------|
| (a) Convulsions/epilepsy                 | yes [    ] | no [    ] |
| (b) Asthma                               | yes [    ] | no [    ] |
| (c) Diabetes                             | yes [    ] | no [    ] |
| (d) A blood disorder                     | yes [    ] | no [    ] |
| (e) Digestive problems                   | yes [    ] | no [    ] |
| (f) Disturbance of balance/co-ordination | yes [    ] | no [    ] |
| (g) Numbness in hands or feet            | yes [    ] | no [    ] |
| (h) Disturbance of vision                | yes [    ] | no [    ] |
| (i) Thyroid problems                     | yes [    ] | no [    ] |
| (j) Kidney or liver problems             | yes [    ] | no [    ] |
| (k) Chest pain or heart problems         | yes [    ] | no [    ] |
| (l) Any other health problems            | yes [    ] | no [    ] |
| (m) An allergy to soya protein or eggs   | yes [    ] | no [    ] |
| (n) An allergy to nuts                   |            |           |

(a) Are you pregnant or think that you might be pregnant      yes [    ]      no [    ]

(b) Do you take the contraceptive pill or other hormone-based contraceptives  
yes [    ]      no [    ]

(c) Are you postmenopausal      yes [    ]      no [    ]

(d) Are you receiving Hormone Replacement Therapy (HRT)  
yes [    ]      no [    ]

(a) Any heart problems	yes [    ]	no [    ]
(b) Diabetes	yes [    ]	no [    ]
(c) Stroke	yes [    ]	no [    ]
(d) Any other family illnesses	yes [    ]	no [    ]

- If so, for how long did you smoke and when did you stop? .....

- If YES to any question, please describe briefly if you wish (e.g. to confirm whether problem was short-lived, insignificant or well controlled.) (Use a separate sheet if necessary)**

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## Appendix C: Kinetic Results For All Subjects According To ApoE Phenotypes

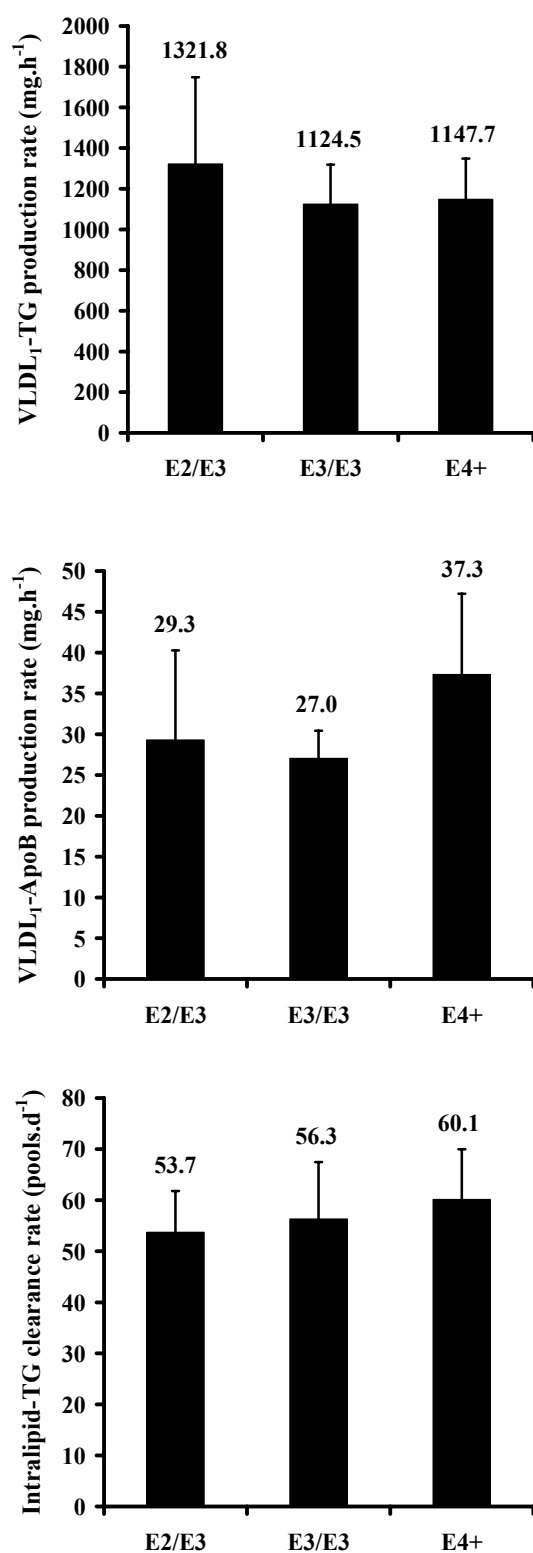


Figure 8.1: Production rates of [A] VLDL<sub>1</sub>-TG and [B] VLDL<sub>1</sub>-apoB and [C] Intralipid-TG clearance rate for all subjects (n = 23) according to apoE phenotypes: E2/E3 (n = 3), E3/E3 (n = 11) and E4+ (n = 9). *P* > 0.05 for differences between phenotypes.