The immediate fate of dietary fat is important to health. Indeed, atherogenesis has been described as a postprandial phenomenon(1), and postprandial lipaemia (the rise in plasma TAG concentrations after a meal) is thought to be involved(2,3). Although the exact mechanisms that link postprandial lipaemia remain to be elucidated, small chylomicron remnants have been implicated in the progression of coronary artery disease(4). Postprandial lipaemia has also been shown to be associated with oxidative stress and inflammation as recently reviewed(5). Therefore, it is important to understand factors that influence the duration and magnitude of postprandial lipaemia. A key tissue in the disposal of meal fatty acids is adipose tissue. The importance of adipose tissue in this respect is outlined in the so-called adipose tissue expandability hypothesis(6).

This hypothesis proposes that ‘a failure in the capacity for adipose tissue expansion, rather than obesity per se is the key factor linking positive energy balance and type 2 diabetes’. With increasing adiposity in some individuals, it is proposed that the capacity of adipose tissue to store further TAG is reduced, and lipids begin to accumulate in other tissues. The aim of this paper is to summarise the metabolic studies in human subjects performed by ourselves and others that have helped to understand the way in which adipose tissue metabolises and stores dietary fat.

The single meal model
A typical protocol for studying postprandial lipaemia is to study volunteers after an overnight fast followed by a
While fasting the role of adipose tissue is to release fatty acids into the systemic plasma in order to supply tissues with a high requirement for fatty acids, such as skeletal muscle and the heart. Thus, fasting plasma NEFA concentrations are high (Fig. 1). After a test meal is given, adipose tissue metabolism is coordinated in order to deal with the nutrient load that is given. A typical metabolic response to a high-fat meal in a healthy non-obese male is shown in Fig. 1. As fat enters the bloodstream in the form of chylomicron-TAG the concentration of plasma TAG increases. The increase in plasma TAG is also partly due to an increase in the concentration of endogenous TAG in VLDL (synthesised in the liver) and plasma taken after a meal containing fat is often cloudy because these large lipoproteins scatter light. The concentration of plasma TAG starts to fall as it is cleared from the plasma (Fig. 1). This is largely mediated by the action of adipose tissue lipoprotein lipase (LPL), situated at the capillary endothelium. LPL hydrolyses chylomicron-TAG, releasing fatty acids to be taken up by adipose tissue. This pathway is up-regulated by insulin, which increases rapidly in response to the carbohydrate content of the meal (Fig. 1). It has long been known that LPL is inhibited by apoCII but it has recently been shown that LPL is also inhibited physiologically by angiopoietin-like protein-4. The expression of this protein appears to be decreased in response to food, thus lifting the inhibition and allowing TAG hydrolysis to proceed at a higher rate. Studies from knock-out mice have shown that LPL also appears to require GPIHBP1, a cell-surface glycoprotein synthesised by the endothelium.

In contrast to the increase in concentration of plasma TAG after a mixed meal, the concentration of plasma NEFA rapidly decreases, but often rebounds above post-absorptive values at the end of the postprandial period (Fig. 1). The initial decrease is due to the action of plasma insulin on the suppression of intracellular lipases. Thus, insulin is a key mediator of changes in adipose tissue fatty acid trafficking in the transition from fasting to fed states; any meal given with no carbohydrate (e.g. pure fat load) would fail to illicit the metabolic responses that depend on the increase in plasma-insulin concentrations. However, adipose tissue is very sensitive to insulin. Therefore, even a small insulin excursion can reduce plasma insulin concentrations (Fig. 2, after fructose ingestion).

The two-meal model

The single meal model is very useful in understanding the key pathways involved in the metabolism of dietary fatty acids, but in reality, most people on a Western diet would eat a second meal less than 5–8 h later. It became clear that postprandial TAG metabolism became complicated by...
a second, later meal, giving an unusual plasma TAG profile with more than one peak. We further investigated plasma TAG concentrations in response to two meals by using naturally occurring fatty acids in foods as tracers of metabolism. After an overnight fast, we gave healthy volunteers a high-fat breakfast that was enriched in linoleic acid (18:2n-6). Five hours later, the volunteers consumed a lunch meal that was enriched in oleic acid (18:1n-9). After the second meal, there was a rapid peak in chylomicron-TAG, and the fatty acid composition of this early peak was remarkably similar to that of the breakfast meal. This suggested that the breakfast fat had been residing in a storage pool before being released by a stimulus from lunch. In an elegant study, Robertson et al. obtained biopsy material from volunteers who had consumed fat 5h previously, and used electron microscopy to show that the fat was in fact residing in the enterocyte, and that it was released in response to a nutrient stimulus.

Spillover fatty acids

An interesting observation from the two-meal study described above was that the plasma NEFA profile was quite unusual. While we normally expect plasma NEFA concentrations to decrease after a meal, an increase was observed in response to lunch. Moreover, the composition of plasma NEFA after lunch was similar to that of the breakfast meal. This suggested that the breakfast fat had been residing in a storage pool before being released by a stimulus from lunch. In an elegant study, Robertson et al. obtained biopsy material from volunteers who had consumed fat 5h previously, and used electron microscopy to show that the fat was in fact residing in the enterocyte, and that it was released in response to a nutrient stimulus.

**Spillover fatty acids**

An interesting observation from the two-meal study described above was that the plasma NEFA profile was quite unusual. While we normally expect plasma NEFA concentrations to decrease after a meal, an increase was observed in response to lunch. Moreover, the composition of plasma NEFA after lunch was similar to that of the breakfast meal. This suggested that the breakfast fat had been residing in a storage pool before being released by a stimulus from lunch. In an elegant study, Robertson et al. obtained biopsy material from volunteers who had consumed fat 5h previously, and used electron microscopy to show that the fat was in fact residing in the enterocyte, and that it was released in response to a nutrient stimulus.

**Spillover fatty acids**

An interesting observation from the two-meal study described above was that the plasma NEFA profile was quite unusual. While we normally expect plasma NEFA concentrations to decrease after a meal, an increase was observed in response to lunch. Moreover, the composition of plasma NEFA after lunch was similar to that of the breakfast meal. This suggested that the breakfast fat had been residing in a storage pool before being released by a stimulus from lunch. In an elegant study, Robertson et al. obtained biopsy material from volunteers who had consumed fat 5h previously, and used electron microscopy to show that the fat was in fact residing in the enterocyte, and that it was released in response to a nutrient stimulus.

**Spillover fatty acids**

An interesting observation from the two-meal study described above was that the plasma NEFA profile was quite unusual. While we normally expect plasma NEFA concentrations to decrease after a meal, an increase was observed in response to lunch. Moreover, the composition of plasma NEFA after lunch was similar to that of the breakfast meal. This suggested that the breakfast fat had been residing in a storage pool before being released by a stimulus from lunch. In an elegant study, Robertson et al. obtained biopsy material from volunteers who had consumed fat 5h previously, and used electron microscopy to show that the fat was in fact residing in the enterocyte, and that it was released in response to a nutrient stimulus.
Adipose tissue metabolism can be quantified in specific adipose tissue depots by the technique of arteriovenous difference. Blood is sampled simultaneously from an artery (or arterialised vein) and a small vein draining the adipose tissue. The difference in concentration of plasma or blood metabolites between the two sites represents metabolism during one pass through the tissue. We have used this technique to study adipose tissue metabolism of meal fatty acids, initially in human abdominal subcutaneous tissue(21) and more recently in the femoral depot(22). In order to quantify flux through the tissue, adipose tissue blood flow (ATBF) must be measured allowing the net transcapillary flux of fatty acids across the tissue to be calculated. This represents the net balance of different pathways of fatty acid trafficking through the tissue. In the postabsorptive state, there is a large concentration gradient between plasma NEFA entering the tissue (artery) and plasma NEFA leaving the tissue (adipose venous drainage). In a group of healthy men (BMI 23–34), the mean values were 591\textsubscript{176}/C\textsubscript{41} and 1208\textsubscript{131}/C\textsubscript{176}, respectively(23), and the mean net flux of NEFA from the tissue was 1080 nmol per 100 g tissue per min. Using an average adipose tissue fatty acid composition(24), this equates to approximately 250 mg adipose tissue TAG hydrolysed per 100 g tissue/min. For a person in energy balance with a 20 kg total fat mass this represents approximately 24 g TAG to be replaced (0.12 % of adipose tissue TAG).

**Adipose tissue uptake of meal fatty acids**

Using the technique of arteriovenous difference and measurements of mass balance, adipose tissue has been found to be a net importer of fatty acids for 5 h following a mixed meal in a number of studies as reviewed(25). The uptake of meal fatty acids can be studied more specifically using either radioactive(26) or stable isotope tracer methodology. With the addition of stable isotope tracers to our protocol of arteriovenous difference described above, we were able to develop a very powerful model to study adipose tissue metabolism in vivo. In this model, [U-\textsuperscript{13}C]palmitic acid is given orally, in order to represent exogenous fatty acids, and [2H\textsubscript{2}]palmitate is given as a continuous intravenous infusion, in order to label endogenous pools (plasma NEFA and VLDL-TAG), see Fig. 3. Using a specific antibody to apoB100, combined with immunoaffinity chromatography, we are able to separate VLDL from chylomicrons(27). This represents a very important advancement in the study of postprandial fatty acids as there is a large overlap in the size and density of these two lipoprotein classes.

**The technique of arteriovenous difference**

Adipose tissue metabolism can be quantified in specific adipose tissue depots by the technique of arteriovenous difference. Blood is sampled simultaneously from an artery (or arterialised vein) and a small vein draining the adipose tissue. The difference in concentration of plasma or blood metabolites between the two sites represents metabolism during one pass through the tissue. We have used this technique to study adipose tissue metabolism of meal fatty acids, initially in human abdominal subcutaneous tissue(21) and more recently in the femoral depot(22). In order to quantify flux through the tissue, adipose tissue blood flow (ATBF) must be measured allowing the net transcapillary flux of fatty acids across the tissue to be calculated. This represents the net balance of different pathways of fatty acid trafficking through the tissue. In the postabsorptive state, there is a large concentration gradient between plasma NEFA entering the tissue (artery) and plasma NEFA leaving the tissue (adipose venous drainage). In a group of healthy men (BMI 23–34), the mean values were 591±41 and 1208±131, respectively(23), and the mean net flux of NEFA from the tissue was 1080 nmol per 100 g tissue per min. Using an average adipose tissue fatty acid composition(24), this equates to approximately 250\textsubscript{125} μg adipose tissue TAG hydrolysed per 100 g tissue/min. For a person in energy balance with a 20 kg total fat mass this represents approximately 24 g TAG to be replaced (0.12 % of adipose tissue TAG).
In the study of healthy men mentioned above (23), we traced 100 mg [U-13C]palmitic acid into the chylomicron fraction where it was measurable at 60 min and peaked at 240 min. We calculated TAG extraction in adipose tissue as (arteriovenous difference of [U-13C]palmitic acid in plasma TAG) × ATBF, and this peaked at 120 min. The time course of fractional TAG extraction is very interesting. At 60 min after the test meal, it is as high as 30% in some individuals but falls to about 10% at 180 min. Presumably, this is because it is only during the early time points that newly released chylomicrons are available for hydrolysis. These newly released chylomicrons would be comparatively large, and therefore a good substrate for LPL (28). In addition, at later time points, plasma TAG-[U-13C]palmitic acid is no longer confined to chylomicrons, and may represent newly synthesised VLDL-TAG. This was less than the fractional extraction of carbohydrate diet for 3 d (31). This rather extreme triacylglycerolaemia promoted by the ingestion of a high carbohydrate diet (75% energy from carbohydrate) led to a doubling in fasting plasma TAG from 1.0 to 2.0 mmol/l compared with a diet that was only 45% energy from carbohydrate. In a diet that is high in extrinsic sugars, leads to an increase in fasting plasma TAG concentrations (30). We explored the capacity of subcutaneous adipose tissue to ‘cope’ with clearing meal fatty acids in the context of hypertriglycerolaemia promoted by the ingestion of a high carbohydrate, low fat diet for 3 d (31). This rather extreme diet (75% energy from carbohydrate) led to a doubling in fasting plasma TAG from 1.0 to 2.0 mmol/l compared with a diet that was only 45% energy from carbohydrate. In response to a standard test meal, the iAUC (incremental area under the curve) for postprandial lipaemia was similar after both diets. However, after the high carbohydrate diet, postprandial plasma TAG concentrations did not return to baseline by the end of the study (360 min). Fasting and postprandial plasma NEFA concentrations were identical after the two dietary regimens. We traced the meal fat with [13C]palmitic acid and found that plasma [13C]palmitate-TAG extraction and TAG clearance in adipose tissue were unaffected by the background diet. Thus, it would seem that despite an increase in the fasting concentration of plasma TAG after the high-fat diet, the ability of adipose tissue to dispose of dietary fat was unaffected. The metabolism of other tissues was, however, affected; TAG clearance (also measured by arteriovenous difference) was significantly lower across the forearm after the high-carbohydrate diet v. the high-fat diet. Also meal fatty acids tended to be repartitioned away from oxidation, towards esterification in the liver and muscle in response to short-term adaptation to the high carbohydrate diet.

Adipose tissue dietary fatty acid metabolism in response to a high carbohydrate diet

It is well known that a high carbohydrate diet, particularly one that is high in extrinsic sugars, leads to an increase in fasting plasma TAG concentrations (30). We explored the capacity of subcutaneous adipose tissue to ‘cope’ with clearing meal fatty acids in the context of hypertriglycerolaemia promoted by the ingestion of a high carbohydrate, low fat diet for 3 d (31). This rather extreme diet (75% energy from carbohydrate) led to a doubling in fasting plasma TAG from 1.0 to 2.0 mmol/l compared with a diet that was only 45% energy from carbohydrate. In response to a standard test meal, the iAUC (incremental area under the curve) for postprandial lipaemia was similar after both diets. However, after the high carbohydrate diet, postprandial plasma TAG concentrations did not return to baseline by the end of the study (360 min). Fasting and postprandial plasma NEFA concentrations were identical after the two dietary regimens. We traced the meal fat with [13C]palmitic acid and found that plasma [13C]palmitate-TAG extraction and TAG clearance in adipose tissue were unaffected by the background diet. Thus, it would seem that despite an increase in the fasting concentration of plasma TAG after the high-fat diet, the ability of adipose tissue to dispose of dietary fat was unaffected. The metabolism of other tissues was, however, affected; TAG clearance (also measured by arteriovenous difference) was significantly lower across the forearm after the high-carbohydrate diet v. the high-fat diet. Also meal fatty acids tended to be repartitioned away from oxidation, towards esterification in the liver and muscle in response to short-term adaptation to the high carbohydrate diet.

Twenty-four hour studies of human adipose tissue metabolism

To follow-on from the arteriovenous difference and two-meal models described above, we investigated postprandial fatty acid metabolism over a 24 h period, in order to study the response to three typical Western-style mixed meals (18). We used a continuous intravenous infusion of [3H2]palmitate as before, to label systemic fatty acids and used three unique uniformly labelled fatty acid tracers to trace the meal fatty acids. To label the breakfast meal, we used [U-13C]linoleic acid, and to label lunch and dinner, we used [U-13C]oleic acid and [U-13C]palmitic acid, respectively. The uniformly labelled notation indicates that all the carbon atoms are stable isotopes and an advantage of this analytically is that there is minimal background (when using GC-MS) compared with fewer carbons being labelled. Additionally, the high density of labelling offers greater potential for tracing metabolic pathways (32). The [U-13C]fatty acid tracers appeared in the plasma TAG pool after each respective meal but although the concentration started to fall after about 5 h, there always remained some tracer present. Thus, 24 h after the ingestion of [U-13C]linoleic acid, there was measurable [U-13C]linoleic acid in the plasma-TG pool. This was assumed to represent chylomicron remnants and tracers that had become incorporated into VLDL-TAG via hepatic recycling. Likewise, [U-13C]linoleic acid in the plasma NEFA pool, representing spillover fatty acids was still present at 24 h. Using a radioactive tracer of meal fatty acids, it has been estimated that 0.9% of meal fatty acids remain in the circulation after 24 h (33). In our study, we were able to calculate the total transcapillary flux of fatty acids across subcutaneous abdominal adipose tissue in healthy non-obese men. This calculation represents the total fatty acids, from all pools, crossing in/out of adipose tissue from the plasma. Over the 24 h period, there was no net uptake of fatty acids immediately after the first meal, and this continued until approximately 17 h after breakfast, i.e. during the whole of the daytime adipose tissue takes up and stores fatty acids. Although the plasma TAG taken up after a meal was mainly from chylomicrons, a proportion was taken up from VLDL-TAG. The quantitative significance of this pathway increased with each meal and made up one-third of the total of the transcapillary flux after the third meal. Plasma NEFA were also taken up directly, but this remained small. We estimated the amount of fat from the meal that was taken up into adipose tissue at a whole-body level. This increased from 15% of the meal fat after the first meal to 48% after last meal. The calculation is only approximate since it depends on an assumption that regional variation of adipose tissue metabolism is insignificant. However, the increase in stored fat with time is still a valid observation. This observation was dependent on an increase in LPL rate of action during the 24 h period, but even more so by an increase in fatty acid re-esterification (i.e. stepwise formation of TAG via the action of the enzymes monoaoylglycerol transferase and diacylglycerol transferase).
The implications for this are important, an excess of meal fatty acids would lead to increased exposure to other tissues and could lead to the possibility of ectopic fat deposition.

Metabolism of specific fatty acids

Some of the measurements described above have been made with the assumption that palmitic acid behaves as a typical fatty acid. Differences in the postprandial metabolism of specific fatty acids can be studied in one of two ways. Either different fatty acids are given within the same typical fatty acid. Differences in the postprandial metabolism of specific fatty acids can be studied in one of two ways. Either different fatty acids are given within the same meal, and compared, or the response of different meals (containing different fatty acids) is compared.
We have investigated the adipose tissue metabolism of different fatty acid species consumed in a single test meal using the technique of arteriovenous difference, using commercially produced structured TAG or natural fats. Perhaps surprisingly, we found that adipose tissue did not discriminate between the fatty acids that we were testing for either type of test meal. Using structured TAG containing oleic acid plus either stearic or palmitic acid, we found no difference in LPL-mediated hydrolysis of the different fatty acids or in their subsequent uptake by the tissue. The structural position of the individual fatty acids within the TAG molecule was also unimportant. We investigated the metabolism of different fatty acids given in the same test meal (14:0, 16:0, 16:1n-7, 18:0, 18:1n-9, 18:2n-6, 20:5n-3, 22:6n-3). Although the molar proportion of the fatty acids in the meal was not maintained in chylomicrons, net uptake into adipose tissue was entirely proportional to their presence in chylomicrons. Nevertheless, overall, this meant that the storage of meal fatty acids into adipose tissue was in the order n-3 polyunsaturated< saturated<n-6 polyunsaturated< monounsaturated; oleic acid was stored in the greatest amount. Note that these studies did not use stable isotope tracers; therefore represent net uptake. We have recently compared the metabolism of stable isotope tracers of 16:0 ([U-13C]palmitate), 18:1n-9 ([U-13C]oleate) and 18:2n-6 ([U-13C]linoleate); all tracers were given simultaneously in the same test meal. We found that their incorporation into chylomicrons was similar. However, there was a tendency for the incorporation of 18:1n-9 to be the highest. This pattern was maintained as the fatty acids became incorporated into plasma NEFA and VLDL-TAG. This was explained by the low partitioning of [U-13C]oleate into plasma cholesteryl ester and phospholipid fractions, and also erythrocyte phospholipid. Meal fatty acids are taken up into adipose tissue mainly via chylomicron TAG, VLDL-TAG and plasma NEFA. Therefore, the isotopic studies have shown that preferential uptake of meal oleate, compared with other meal fatty acids, could occur due to greater availability in these fractions.

Another aspect of comparing the metabolism of different fatty acids is to compare their postprandial metabolism when given separately. In this case, the fatty acids could behave differently due to different physico-chemical properties of the chylomicrons formed from the different test meals. This is difficult to study in human subjects because of the rapidity by which chylomicrons are cleared in the blood. However, animal studies in which the lymph has been accessed have shown that different dietary fatty acids form different sized chylomicrons. For example, linoleic acid forms larger chylomicrons than palmitic acid, and there is some evidence to suggest that this is the case in human subjects too, as reviewed\(^{40}\) but the relationship is not clear\(^{41,42}\). Paradoxically, greater accumulation of fat on the hips is beneficial in terms of risk of myocardial infarction\(^{43}\).

Because of the importance of body fat distribution, studies have attempted to look at mechanisms behind differential accumulation of fat in different depots. Regional meal fat storage has been studied by giving a \(^{3}H\) or \(^{14}C\) fatty acid tracer with a meal and then measuring specific activity in adipose tissue depots 24 h later. An advantage of radioactive tracers in this respect is that they are very sensitive; an important consideration since the tracer is dispersed into a large pool.

Regional differences in the uptake of meal fatty acids into adipose tissue depots has been comprehensively investigated by Jensen et al. using a \(^{3}H\)triolein tracer given with a liquid test meal\(^{33}\). Fatty-acid oxidation was determined by incorporation of the tracer into the plasma-water pool and it was estimated that approximately 50% of the tracer was oxidized over the 24 h. Intra-abdominal fat took up the tracer with the greatest avidity (Table 1) but the upper body depot, being the largest depot, accounted for over half of the uptake of fatty acids not oxidized. After a high-fat, high-energy meal, women store more dietary fatty acids in leg fat than men\(^{44}\). Moreover, meal fatty acid storage (mg meal fat/g adipose tissue lipid) was greater in women with more leg fat than those with less leg fat. Also, after a high-fat meal, the depots with more fat seemed to take up meal fat less avidly than those with less fat.

Mcguaid et al. have recently developed a technique to apply the technique of arteriovenous difference to femoral adipose tissue by cannulation of the saphenous vein\(^{22}\). In accordance with the work of Jensen et al. (Table 1), it was found that net uptake of meal fatty acids was lower in femoral than subcutaneous abdominal tissue. ATBF was also lower in the femoral depot. An interesting finding was that while chylomicron-TAG was the preferred source for postprandial fat deposition in the subcutaneous abdominal depot, the femoral depot discriminated less against VLDL-TAG. It was therefore hypothesized that femoral adipose tissue may accumulate dietary fatty acids that have been recycled as VLDL and NEFA.

### Table 1. Uptake of fatty acid tracer into different adipose tissue depots in lean men and women (n 6) (data taken from Jensen et al\(^{33}\)). The lower body subcutaneous fat is demarcated as all adipose tissue caudal to the inguinal ligament anteriorly and the ileac crest posteriorly\(^{45}\)

<table>
<thead>
<tr>
<th></th>
<th>% Body fat</th>
<th>% Taken up</th>
<th>Relative uptake*</th>
</tr>
</thead>
<tbody>
<tr>
<td>Intra-abdominal</td>
<td>11±7</td>
<td>17±18</td>
<td>1·5</td>
</tr>
<tr>
<td>Lower body subcutaneous</td>
<td>26±11</td>
<td>16±11</td>
<td>0·6</td>
</tr>
<tr>
<td>Upper body subcutaneous</td>
<td>63±15</td>
<td>57±30</td>
<td>0·9</td>
</tr>
</tbody>
</table>

*% Taken up/% body fat.

---

**Conclusions**

Healthy adipose tissue adapts rapidly to the ingestion of a mixed meal. Intracellular lipolysis is immediately...
suppressed, reducing the flux of fatty acids leaving the tissue. An increase in ATBF facilitates the entry of meal fat into the tissue, thus providing substrate for LPL which is activated by the action of insulin. The fractional extraction of chylomicrons is very efficient, as much as 30% in the early postprandial period. It is possibly higher than this at earlier time points. Not all fatty acids released by the action of LPL are taken up by the tissue; some are released into the plasma, particularly in the late postprandial period where as much as 50% of the plasma NEFA pool is composed of meal fatty acids. While these dietary fatty acids may be taken up by non-adipose tissues, there is the potential to recycle back to adipose tissue and be taken up directly from the plasma NEFA pool. Adipose tissue is a net importer of dietary fat for 5 h following a single test meal and for most of the day during a typical three-meal eating pattern. The action of LPL seems to increase sequentially after meal intake, but uptake of meal fatty acids into adipose tissue increases to a greater extent, suggesting up-regulation of pathways of esterification. Obesity and insulin resistance are associated with higher fasting and postprandial plasma TAG concentrations and reduced efficiency of adipose tissue meal-fat storage. This offers a possible explanation to explain ectopic fat deposition associated with obesity. Hypertriglyceridemia due to a high carbohydrate diet in non-obese individuals does not affect the ability of adipose tissue to carry dietary fat. There is a marked difference in the way that different adipose tissue depots handle dietary fat and this may be related to the metabolic phenotypes associated with different body fat distribution patterns.

Acknowledgements

Financial support is acknowledged from the project ‘Hepatic and adipose tissue and functions in the metabolic syndrome’ (HEPADIP, http://www.hepadip.org/), which is supported by the European Commission as an Integrated Project under the 6th Framework Programme (Contract LSHM-CT-2005-018734). The author declares no conflict of interests. I am indebted to all my work colleagues, past and present, for help, kindness and support. I would like to pay particular thanks to my mentors, Professor Keith Frayn and Fredrik Karpe, and colleagues Sandy Humphreys and Leanne Hodson. Thanks also to Mark Fielding for help with data handling.

References

22. McQuaid SE, Manolopoulos KN, Dennis AL et al. (2010) Development of an arterio-venous difference method to study the metabolic physiology of the femoral adipose tissue depot. Obesity (Silver Spring) 18, 1055–1058.


