Older adults have an altered chylomicron response to a high-fat meal

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Abstract
Ageing is associated with a prolonged and exaggerated postprandial lipaemia. This study aimed to examine the contribution of alterations in chylomicron synthesis, size and lipid composition to increased lipaemia. Healthy older (60–75 years; n 15) and younger (20–25 years; n 15) subjects consumed a high-fat breakfast. Chylomicron dynamics and fatty acid composition were analysed for 5 h in the postprandial state. Plasma TAG levels were elevated following the meal in the older subjects, relative to younger subjects (P<0·01). For older subjects compared with younger subjects, circulating chylomicron particle size was smaller (P<0·05), with greater apoB content (P<0·05) at all postprandial time points. However, total chylomicron TAG concentration between the groups was unaltered post-meal. Compared with younger subjects, the older subjects exhibited a greater proportion of oleic acid in the TAG and phospholipid (PL) fraction (P<0·05), plus lower proportions of linoleic acid in the TAG fraction of the chylomicrons (P<0·01). Thus, following the ingestion of a high-fat meal, older individuals demonstrate both smaller, more numerous chylomicrons, with a greater total MUFA and lower PUFA contents. These data suggest that the increased postprandial lipaemia of ageing cannot be attributed to increased chylomicron TAG. Rather, ageing is associated with changes in chylomicron particle size, apoB content and fatty acid composition of the chylomicron TAG and PL fractions.

Key words: Cholesterol; Diet and dietary lipids; Lipidomics; Lipoproteins; Nutrition; Phospholipids; TAG

Postprandial lipaemia, characterised by increased concentrations of TAG-rich chylomicron and remnant particles, exerts a significant and often unappreciated impact on CVD risk(11). Chylomicrons are synthesised within the small intestinal enteroctye, with a lipid composition highly reflective of the previous meal ingestion(2,3). Chylomicrons tend to increase in size with increasing quantities of ingested lipids, but are also larger when composed of proportionally more longer-chain and less saturated TAG(4,5). Although larger chylomicrons are cleared more slowly than smaller particles, the total number of particles competing for removal has been shown to be the rate-limiting factor in postprandial chylomicron and remnant clearance(6), with chylomicrons being the preferential lipolytic substrate over remnants(7,8).

Older individuals experience greater postprandial lipaemia(9–15). The current evidence suggests that ageing is associated with an increased TAG content of chylomicrons(14,15). Further, measurements of postprandial chylomicron dynamics in older adults have been inconsistent and are frequently reliant on proxy measures such as fat-soluble vitamin appearance(12–15) or apoB-48:TAG ratios(16). Direct measurement of chylomicron particle size with ageing is lacking and this can be quantified by dynamic light scattering methods(17). As chylomicron size dictates lipolytic clearance, age-related differences in chylomicron composition and size could contribute to the reported impaired chylomicron remnant clearance and hypertriacylglycerolaemia in elderly adults(18).

Fat meal loads used in previous studies to induce postprandial lipaemia have varied widely and have often been limited to simple fat boluses that are poorly representative of a typical meal(19). As macronutrients exert differential metabolic responses when ingested alone(19) or in unison(19,20), it is of considerable interest to understand the metabolic response to representative mixed meals. High-fat and high-carbohydrate meals representative of a typical Western diet have increasingly been used to study postprandial lipaemia in healthy(21–23) and metabolically compromised populations(24,25). Thus, in this study, direct measurement of the effect of age on chylomicron dynamics and composition following a high-fat (and high-carbohydrate) meal, representative of a Western diet, was conducted. Analysis was performed to examine chylomicron particle size, apoB concentration and fatty acid composition of both the chylomicron TAG and the phospholipid (PL) fractions in a healthy younger and older adult population. We hypothesised that an exaggerated postprandial lipaemic
Methods

Subject selection

In total, thirty healthy, community-dwelling subjects (n 7 young females, n 8 young males, n 9 older females, n 6 older males) from the Auckland region were recruited through newspaper advertisements and from the university community to participate in the study. Eligible subjects were required to have a BMI between 18 and 30 kg/m² and be between the ages of 20–25 and 60–75 years. Individuals with a history of CVD or metabolic disease/conditions including diabetes or thyroid conditions or who used medications that may interfere with study end points (i.e. anti-inflammatory drugs, statin drugs) were not eligible for participation. All subjects reported that they were non-smokers. This study was conducted according to the guidelines laid down in the Declaration of Helsinki, and all procedures involving human subjects were approved by the University of Auckland Human Participants and Ethics Committee (ref no. 8026). Written informed consent was obtained from all subjects. This study was registered prospectively at Australian New Zealand Clinical Trials Registry at anzctr.org.au (ID: ACTRN12612000515897).

Study procedures

The present study was a cross-sectional, postprandial trial using a high-fat breakfast, chosen as a standard test meal with high fat (62.2 g) and protein content, also representative of a high-fat restaurant breakfast meal (23) (Table 1). Subjects were asked to abstain from vigorous physical activity, high-fat foods, anti-inflammatory medications and nutritional supplements the day before their visit. Subjects arrived fasted on the morning of their visit; anthropometric data were collected before a catheter was inserted into an antecubital vein and a baseline sample (time 0 h) was obtained followed by consumption of the test breakfast. Subject physical activity was limited to sedentary activities within the clinical room of the study facility. Blood samples were collected after the meal into serum- and EDTA-containing blood collection tubes (Becton Dickinson) every hour for 5 h. Serum tubes were left to clot at room temperature for 15 min before all the tubes were centrifuged at 1500 g for 15 min at 4°C for serum and plasma separation. An aliquot of plasma was maintained at 4°C for chylomicron-rich fraction (CMRF) separation within 6 h, and the remaining plasma was collected in pyrogen-free microtubes and stored at −20°C until analysis.

Chylomicron-rich fraction isolation

The CMRF, containing chylomicrons and their large remnants, was separated in 4-ml OptiSeal tubes (Beckman Coulter) in an Optima MAX-XP ultracentrifuge using a TLA-110 rotor (Beckman Coulter), adapted from Oikawa et al. (26). Density-gradient saline solutions were prepared with NaCl and 0.005% EDTA (Sigma-Aldrich) according to Naito (27), and separation protocols were based on the studies of Kupke & Wörz-Zeugner (28). In brief, 3-5 ml plasma was overlaid with 1-2 ml saline solution (d = 1.006 g/ml) and centrifuged at 117 000 g for 10 min. The visible top layer was aspirated into microtubes and corrected to a final collection volume of 1-4 ml using saline solution. This provided a standardised dilution factor of the collected CMRF volume relative to initial plasma volume. The CMRF was stored in pyrogen-free microtubes at −80°C until use.

Chylomicron-rich fraction particle size analysis

CMRF particle size was measured using a NanoZetasizer S (Malvern Instruments) by dynamic light scattering. If more than one size peak was observed, the mean peak data for the peak within the range of large TAG-rich lipoproteins (TRL) were used (>70 nm) based on the lower range of mean diameters of small chylomicrons (Sf 400–1000), as reported by Fraser (29).

Biochemical analysis

Concentrations of plasma glucose, cholesterol, LDL, HDL, TAG, NEFA and CMRF TAG, apoB, serum alanine transaminase (ALT) and serum aspartate transaminase (AST) were measured using a Hitachi 902 autoanalyzer (Hitachi High Technologies Corporation) by enzymatic colorimetric assay (Roche). Plasma insulin concentration was measured using an Abbott AxsYM system (Abbott Laboratories) by microparticle enzyme immunoassay.

Fatty acid composition analysis of meals and chylomicron-rich fraction TAG and phospholipids

Meals were homogenised using a domestic blender with 600 g water. Samples were aliquoted into 50-ml centrifuge tubes,
frozen and stored at −80°C until use. Fatty acid extraction, methylation and GC of meal and CMRF samples were carried out according to protocols described previously (30–32). In short, an internal standard mixture of triheptadecanoin (Sigma-Aldrich) and dinonadecanoylphosphatidylcholine (Sigma-Aldrich) was added to the isolated CMRF. Subsequently, 1.5 ml methanol, 3 ml chloroform and 0.8 ml 0.88 % KCl in water were added and the blend was thoroughly vortexed after each addition. Samples were centrifuged at 2000 × g for 3 min to separate the layers, and the lower chloroform-rich layer was removed and the upper layer was extracted again (30). Food lipids were extracted just as CMRF after further homogenisation with an Ultra Turrax (IKA). CMRF TAG and PL were isolated from the extracted lipid mixture with solid-phase extraction based on Sep-Pak (Waters) (31). Fatty acid methyl esters (FAME) were prepared using the sodium methoxide method. In short, the lipids were suspended into 1 ml dry diethyl ether; subsequently, 25 μl methyl acetate and 25 μl sodium methoxide were added and the blend was incubated for 5 min while shaken at times. The reaction was stopped with 6 μl acetic acid. Samples were centrifuged at 2000 × g for 5 min, after which the supernatant was removed and gently evaporated to dryness. The resulting FAME were transferred to 100 μl inserts in hexane (32). The FAME were analysed using a gas chromatograph (Shimadzu GC-2010) equipped with an AOC-20i auto injector, a flame ionisation detector (Shimadzu Corporation) and a wall-coated open tubular column DB-23 (60 m × 0.25 mm i.d., liquid film 0.25 μm, Agilent Technologies, J.W. Scientific). Splitless/split injection was used and the split was opened after 1 min. Supelco 37 Component FAME Mix (Supelco), 68D (Nu-Chek-Prep) and GLC-490 (Nu-Chek-Prep) were used as external standards.

**Calculations**

The homoeostatic model assessment of insulin resistance (HOMA-IR) was calculated from fasting glucose and insulin concentrations using the equation from the study by Matthews et al. (33). Maximum concentrations (Cmax) were identified as the greatest value in an individual’s data set, with maximum peak times (Tmax) identified as the corresponding nominal sampling time. Arithmetic means were used for group Cmax comparisons, whereas median values were used for group Tmax comparisons. Plasma fatty acids were pooled for mathematical analysis into SFA, MUFA, PUFA and dairy-derived odd-chain fatty acids.

**Statistical analyses**

Statistical analyses were carried out using Statistical Package for the Social Sciences version 21 (SPSS; IBM Corporation). Sample size was estimated with a power analysis for a superiority design based on data from a previous study on lipaemia in the elderly (34). To achieve a significance level of P ≤ 0.05 with 80 % power to detect a between-subject difference of 4.77 mmol/l per h for TAG incremental AUC (iAUC), a sample size of 30 was necessary. Data are presented as means values with their standard errors except Tmax which is presented as the medians and interquartile ranges. iAUC was calculated after subtracting fasting values. Baseline subject characteristics, iAUC and Cmax were compared using unpaired Student’s t test. Tmax was compared using the Mann–Whitney U test using mean ranks. Two-factor repeated-measures ANOVA (time compared within-subject and age compared between subjects) followed by the Sidak-adjusted post hoc test were used for all multiple comparisons between different groups. Where Mauchly’s sphericity test failed, the Huynh–Feldt correction was applied. α was set at P < 0.05. Heat maps were created using R software version 2.15.2 using the packages gplots (heatmap.2), RColorBrewer and colorRamps (R Development Core Team).

**Results**

**Subject characteristics**

A total of thirty subjects completed the study (n 15 younger, n 15 older). There were no significant differences between the two age groups for BMI or fasting measurements of plasma glucose, plasma insulin, HOMA-IR, plasma or CMRF TAG or CMRF apoB content or particle size (Table 2). Fasting plasma lipid profiles were within the normal range for both age groups, although total, LDL and HDL-cholesterol levels were all higher in older subjects (P < 0.001, P = 0.002 and P < 0.001, respectively). Liver function measured by serum ALT and AST were within the normal range for all participants; however, the older participants displayed lower concentrations for each (P = 0.004 and P = 0.001, respectively).

**Plasma glucose, insulin and lipid responses**

Postprandial glucose (Fig. 1a i) and insulin (Fig. 1b i) responses to the meal did not differ between younger and older

<table>
<thead>
<tr>
<th>Measures</th>
<th>Younger subjects (n 15)</th>
<th>Older subjects (n 15)†</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age (years)</td>
<td>22.7 ± 0.4</td>
<td>67.3 ± 1.5</td>
</tr>
<tr>
<td>BMI (kg/m²)</td>
<td>23.7 ± 0.8</td>
<td>24.4 ± 1.0</td>
</tr>
<tr>
<td>HOMA-IR</td>
<td>2.0 ± 0.2</td>
<td>1.7 ± 0.3</td>
</tr>
<tr>
<td>Plasma</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Glucose (mmol/l)</td>
<td>5.1 ± 0.1</td>
<td>5.2 ± 0.1</td>
</tr>
<tr>
<td>Insulin (mmol/l)</td>
<td>9.0 ± 1.0</td>
<td>8.7 ± 1.9</td>
</tr>
<tr>
<td>Total cholesterol (mmol/l)</td>
<td>4.0 ± 0.2</td>
<td>5.0 ± 0.2†</td>
</tr>
<tr>
<td>LDL (mmol/l)</td>
<td>2.4 ± 0.2</td>
<td>3.0 ± 0.2†</td>
</tr>
<tr>
<td>HDL (mmol/l)</td>
<td>1.4 ± 0.1</td>
<td>1.8 ± 0.1†</td>
</tr>
<tr>
<td>TAG (mmol/l)</td>
<td>0.8 ± 0.07</td>
<td>0.91 ± 0.07</td>
</tr>
<tr>
<td>NEFA (mmol/l)</td>
<td>0.66 ± 0.06</td>
<td>0.57 ± 0.09</td>
</tr>
<tr>
<td>Serum</td>
<td></td>
<td></td>
</tr>
<tr>
<td>ALT (U/l)</td>
<td>10.0 ± 2.2</td>
<td>1.9 ± 0.8</td>
</tr>
<tr>
<td>AST (U/l)</td>
<td>19.1 ± 1.7</td>
<td>12.1 ± 0.7</td>
</tr>
<tr>
<td>Chylomicron-rich fraction</td>
<td></td>
<td></td>
</tr>
<tr>
<td>TAG (mmol/l of plasma)</td>
<td>0.07 ± 0.02</td>
<td>0.08 ± 0.01</td>
</tr>
<tr>
<td>apolipoprotein B content</td>
<td>2.3 ± 0.5</td>
<td>3.4 ± 0.5</td>
</tr>
</tbody>
</table>

HOMA-IR, homoeostatic model assessment of insulin resistance; ALT, alanine transaminase; AST, aspartate transaminase.

† Significance was determined by unpaired Student’s t test. *** P < 0.001, ** P < 0.01 compared with younger subjects.
subjects. The older group had a delayed median time to achieve peak plasma TAG concentrations relative to the younger group ($T_{\text{max}}$ 3 (IQR 3–4) v. 2 (IQR 1–3) h, respectively; $P<0.001$), reaching peak TAG concentration ($C_{\text{max}}$) after 3 h ($P<0.001$ v. baseline). The younger subjects reached maximum peak concentration at 2 h ($P=0.02$ v. baseline). The plasma TAG in the younger subjects also returned to baseline by 4 h; however, in the older group, plasma TAG remained significantly elevated at 5 h post-meal (Fig. 1(c)). As such, we observed a prolonged TAG response in older subjects. This pattern of response did not impact on the iAUC (0–5 h) response between the younger and the older groups (103±6 (SEM 21±6) v. 102±5 (SEM 19±1) mmol/l, respectively; $P=0.97$; data not shown). Marked suppression of circulating plasma NEFA was observed following the meal, with no difference between younger and older subjects (Fig. 1(d)). Postprandial total plasma cholesterol concentrations decreased in the older but not in the younger participants (Table 3; time–age interaction; $P<0.001$).

**Chylomycin-rich fraction dynamics**

CMRF particle size increased after the meal ($P=0.01$; main effect of time, Fig. 2(a)) and was smaller for the older compared with the younger participants ($P=0.011$, main effect of age). Owing to variation in time to peak size between individuals, median group maximum peak CMRF particle size and $T_{\text{max}}$ were compared. Overall median peak time was 3 h post-meal ($T_{\text{max}}$ 3 (IQR 2–3–5) v. 2 (IQR 1–4) h, younger v. older, respectively; $P=0.467$). Maximum CMRF particle size tended to be smaller in the older subjects ($C_{\text{max}}$ 321±21 (SEM 29) and 390±22 (SEM 22) nm, older v. younger, respectively; $P=0.072$). As CMRF particle size was not different at baseline between age groups, post hoc analysis was conducted to determine age-group postprandial changes in size, despite no time–age interaction. The older group’s CMRF particle size remained constant after the meal, whereas the younger group’s CMRF particle size increased, resulting in an age difference in size at 2–3 h ($P=0.022$ and $P=0.007$, respectively). Postprandial CMRF TAG concentrations decreased in the older but not in the younger participants ($P=0.01$ post hocs)

![Fig. 1. Postprandial plasma concentrations of glucose (a), insulin (b), TAG (c) and NEFA (d) in older (–––) and younger (---) subjects. Values are means (n 15/group), with their standard errors. There were no differences in glucose, insulin or NEFA responses between older and younger subjects. There were significant differences over time in the TAG response between older and younger subjects (age $\times$ time interaction; $P<0.001$).](https://www.cambridge.org/core/doi/10.1017/S000711451500505X/asset/155050DX/155050DX)

**Table 3. Postprandial total cholesterol in older and younger subjects**

<table>
<thead>
<tr>
<th>Time (h)</th>
<th>Younger subjects (n 15)</th>
<th>Older subjects (n 15)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean ± SEM</td>
<td>Mean ± SEM</td>
</tr>
<tr>
<td>Baseline</td>
<td>3.99 ± 0.18</td>
<td>5.04 ± 0.22</td>
</tr>
<tr>
<td>1 h</td>
<td>3.93 ± 0.17</td>
<td>4.92 ± 0.16</td>
</tr>
<tr>
<td>2 h</td>
<td>3.95 ± 0.18</td>
<td>4.84± 0.18</td>
</tr>
<tr>
<td>3 h</td>
<td>3.86 ± 0.18</td>
<td>4.75± 0.18</td>
</tr>
<tr>
<td>4 h</td>
<td>3.93 ± 0.20</td>
<td>4.75***± 0.16</td>
</tr>
<tr>
<td>5 h</td>
<td>3.97 ± 0.19</td>
<td>4.80**± 0.16</td>
</tr>
</tbody>
</table>

† Significance was determined by two-factor repeated-measure ANOVA with Sidak-corrected post hoc analysis; ** $P<0.01$, *** $P<0.001$ compared with baseline within age group; § $P<0.01$ compared with 1 h. All time point comparisons between younger and older subjects were significant to $P<0.01$. 

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concentration was not different between older and younger subjects (P=0.894; Fig. 2(b)).

Changes in CMRF particle numbers were assessed by apoB concentrations. Older subjects had higher apoB concentrations than younger subjects (mean postprandial concentration of 5.3 (SEM 0.4) v. 4.2 (SEM 0.4) × 10^{-2} g/l, respectively; P=0.035; Fig. 2(c)). Overall, apoB concentrations peaked at 2 h post-meal (P=0.001) returning to comparable baseline concentrations by 5 h.

**Chylomicron-rich fraction TAG and phospholipid fatty acid composition**

**Comparison of chylomicron-rich fraction TAG composition between ages.** Older subjects had greater proportions of MUFA in their CMRF TAG (5.6 % greater proportion in older v. younger subjects, P=0.014; Fig. 3). In addition, 4 h after the meal, older subjects had a lower proportion of PUFA in their CMRF TAG (9.4 % lower proportion in older v. younger subjects, P=0.004). Older participants’ CMRF TAG tended to have lower proportions of SFA at 2 h (7.0 % lower proportion in older v. younger subjects; P=0.079), a trend that was lost by 4 h.

These differences in MUFA and PUFA CMRF TAG composition were attributable to differences in the major fatty acids – oleic acid and linoleic acid. The older group’s CMRF TAG had relatively more oleic acid (18:1n-9; P=0.033; Fig. 4(a)). Although 18:2n-6 (linoleic acid) was found in equal proportions in both groups’ CMRF TAG at 2 h, by 4 h the older subjects had lower proportions (P=0.003). The older group had lower proportions of 18:0 (stearic acid) at 2 h compared with younger participants (P=0.009), whereas these age differences were not apparent after 4 h. Older subjects had lower proportions of 15:1 at 2 h compared with the younger group (P=0.037).

Pooling of the dairy-derived odd-chain fatty acids (15:0, 15:1, 17:0, 17:1) showed higher proportions in the CMRF TAG of older subjects (P<0.001). In both groups, CMRF TAG composition changed over the postprandial period. From 2 to 4 h after the meal, proportions of CMRF TAG 20:4n-6 (arachidonic acid) and 22:6n-3 (DHA) increased (P=0.002 and P=0.042, respectively).

**Comparison of chylomicron-rich fraction phospholipid composition between ages.** In both older and younger subjects, CMRF PL composition changed over the postprandial period. At 4 h, CMRF PL contained relatively more PUFA (P=0.034), less SFA (P=0.042) and tended to contain more MUFA (P=0.052) than at 2 h (Fig. 3). This difference in the later
Discussion

Following a high-fat meal, the older participants of the present study had smaller chylomicron particles with a greater proportion of oleic acid and a lower PUFA content over a 5-h postprandial period. This study highlights that there are subtle changes in the dynamics governing lipid absorption, which occur along with differences in relative absorption and/or clearance of specific fatty acids, predominately oleic acid. Furthermore, our data do not support either altered appearance or delayed chylomicron clearance as a major regulator of the exaggerated lipaemia, typically present in older individuals. Rather, this study shows that the major contributor to the heightened postprandial TAG response is in the non-chylomicron fraction and includes other TAG-rich particles including VLDL and chylomicron remnants. Consistent with previous reports, our data demonstrate prolonged postprandial triacylglycerolaemia in older adults. However, within this literature, a few studies have addressed the dynamics and role of chylomicron structure and composition in this hypertriacylglycerolaemic response. In the present study, the concentration of CMRF TAG was not different between older and younger participants. Previous studies reporting differences in chylomicron TAG have shown greater TAG content in large, rather than small, TRL in the elderly, plus a greater postprandial NEFA spillover from exogenous TRL. Neither mechanism was demonstrated in the present study. Rather, our data implicate alterations of small TRL (endogenous VLDL and small chylomicron remnants) production and clearance as a major determinant of the sustained postprandial hypertriacylglycerolaemia in the elderly. It must be noted that there were no age differences in BMI, HOMA-IR, fasting TAG and NEFA or indications of hepatic...
dysfunction in this present study. Previously, these factors have been shown to adversely affect postprandial chylomicron dynamics\(^{11}\). Furthermore, our findings are presented in the context of a mixed breakfast meal, rather than in response to the ingestion of isolated lipid loads.

In this study, the older participants had a maximum chylomicron particle size, on average 18% smaller and 27% more numerous compared with the younger group. Importantly, chylomicron sizes were measured directly through dynamic light scattering, a technique used previously for chylomicron and lipoprotein size analyses\(^{37,34}\), although rarely in the context of postprandial dynamics\(^{45}\). These smaller and more numerous TRL may be due to intestinal over-production of TAG in young subjects\(^{48}\), supporting a difference in post-meal TAG in young subjects\(^{48}\), suggesting an important impact of habitual diet\(^{47}\). In contrast, meal composition has been reported to significantly impact postprandial TAG in young subjects\(^{49}\), supporting a difference in post-meal fatty acid metabolism between old and young individuals. Furthermore, as our postprandial findings match longer-term evaluations of circulating fatty acids in elderly populations, the metabolic handling of fatty acids may contribute to age-related fatty acid differences in the elderly.

There are likely to be several mechanisms exerting influence on the differences in chylomicron fatty acid composition between the younger and the older subjects following the consumption of an identical meal, including possible differences in fatty acid absorption and clearance from TRL. Enterocyte fatty acid metabolism may be involved as chylomicron incorporation of fat-soluble vitamins has suggested age-related differences in postprandial absorption\(^{12,14}\). The greater proportions of MUFA observed in the older group may indicate enhanced intestinal absorption of oleic acid. This is supported by animal models indicating that uptake of oleic acid\(^{49}\), linoleic acid\(^{50}\) and SFA\(^{51}\) are increased in aged rats, fitting with our data. In addition, chylomicron TAG are known to contain proportions of fatty acids originating from enterocyte storage pools, containing fatty acids from the previous meal\(^{52,53}\) as well as early postprandial contributions from NEFA\(^{52}\). It has been suggested that in insulin-resistant states the intestines may store greater quantities of fat, contributing to aberrant apoB production\(^{54}\). It is unknown whether differences in intestinal enterocyte storage pools between age groups may have contributed to postprandial fatty acid compositional differences. However, it is probable that the predominant mediator is exchange with either circulating free or lipoprotein-derived PL and TAG or the presence of endogenous large TRL. Indeed, the postprandial reductions in total cholesterol we observed in the older group suggest increased inter-lipoprotein transfer; however, measures of this activity such as cholesterol ester transfer activity were not assessed. Furthermore, the origins of altered proportions of fatty acids in any endogenous lipoproteins, such as greater MUFA content, remain unknown.

The postprandial age-related differences in older participants’ chylomicron composition may have metabolic implications outside of transient effects on postprandial lipaemia. As older people incorporate ingested TAG at a greater rate into small TRL rather than chylomicrons\(^{11}\), any differences in fatty acid uptake will be amplified with the increased residence time of these remnants\(^{11,15}\). In addition, the increased surface area: volume ratio of chylomicrons in the older group implies proportionally more PL, magnifying the impact of these compositional differences. Specifically, enhanced uptake of oleic acid could have implications for uptake of other fatty acids, as long-chain fatty acid uptake into the enterocyte is competitive\(^{55}\), and could explain the lower proportions of MUFA found in older subjects’ chylomicron TAG. Furthermore, the fatty acid composition differences reported in these older adults have been identified as features in individuals with metabolic dysfunction; lower serum linoleic acid has been associated with an increased risk of developing impaired fasting glucose or type 2 diabetes mellitus in middle-aged men\(^{56}\). In addition, as we show that some age-related compositional differences are dependent on postprandial sampling time, differences in rates of uptake or clearance may contribute to these age-related differences, highlighting a downfall of compositional profile analysis from a single time point when temporal metabolic differences exist between populations.
Conclusions

The exaggerated postprandial lipaemia typical of older individuals is not characterised by elevated chylomicron TAG. This study demonstrated that in an older healthy cohort there is no impairment of large chylomicron clearance. Despite this preservation of a chylomicron response to a high-fat meal, the resultant chylomicrons are smaller and more numerous in the elderly, containing greater proportions of MUFA and lower proportions of PUFA. The implications of these fundamental differences in chylomicron dynamics and composition are unknown in the context of CVD risk and warrant further investigation.

Acknowledgements

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A. M. M. carried out laboratory analysis and data interpretation and drafted the manuscript. A. N. carried out lipidomic analysis. A. M. M., S. P. and C. A. P. were involved in the coordination, management and implementation of the clinical trial. J. F. M. provided laboratory supervision. J. F. M. and K. M. L. critically evaluated the manuscript. K. M. L. supervised lipidomic analysis. D. C.-S. formulated the research question, initiated and supervised all aspects of the study. All the authors approved the final version of the manuscript for submission.

The authors declare that there are no conflicts of interest.

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