

## **Association of oily fish intake, sex, age, BMI, and *APOE* genotype with plasma long chain n-3 fatty acid composition**<sup>1-3</sup>

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<sup>4</sup>Abbreviations:

LC n-3 PUFA, Long chain omega-3 polyunsaturated fatty acids; EPA, Eicosapentaenoic acid; DPA, Docosapentaenoic acid; DHA, Docosahexaenoic acid; APOE, Apolipoprotein E; BMI, body mass index; PC, Phosphatidylcholine; NEFAs, Non-esterified fatty acids; CEs, Cholesteryl esters; TGs, Triacylglycerols; FFQ, Food frequency questionnaire; FAMES, Fatty

acid methyl esters; GLM, General linear model; SEM, Standard error mean; LDL, Low-density lipoprotein; LDLRs, Low-density lipoprotein receptors; HDLs, High density lipoproteins; LDLC, LDL-cholesterol.

Running title: Determinants of fatty acid status

## 1 Abstract

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3 Omega-3 fatty acids are associated with better cardiovascular and cognitive health. However, the  
4 concentration of EPA, DPA and DHA in different plasma lipid pools differs and factors influencing  
5 this heterogeneity are poorly understood. Our aim was to evaluate the association of oily fish intake,  
6 sex, age, BMI and *APOE* genotype with concentrations of EPA, DPA and DHA in plasma PC,  
7 NEFAs, CEs and TGs. Healthy adults (148 male, 158 female, age 20-71 years) were recruited  
8 according to *APOE* genotype, sex and age. Fatty acid composition was determined by gas  
9 chromatography. Oily fish intake was positively associated with EPA in PC, CEs and TGs, DPA in  
10 TGs, and DHA in all fractions ( $P \leq 0.008$ ). There was a positive association between age and EPA  
11 in PC, CEs and TGs, DPA in NEFAs and CEs, and DHA in PC and CEs ( $P \leq 0.034$ ). DPA was  
12 higher in TGs in males than females ( $P < 0.001$ ). There was a positive association between BMI  
13 and DPA and DHA in TGs ( $P < 0.006$  and  $0.02$ , respectively). *APOE* genotype\*sex interactions  
14 were observed: the *APOE4* allele associated with higher EPA in males ( $P = 0.002$ ), and there was  
15 also evidence for higher DPA and DHA ( $P \leq 0.032$ ). In conclusion, EPA, DPA and DHA in plasma  
16 lipids are associated with oily fish intake, sex, age, BMI, and *APOE* genotype. Such insights may be  
17 used to better understand the link between plasma fatty acid profiles and dietary exposure and may  
18 influence intake recommendations across population subgroups.

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23 **Keywords:** apolipoprotein E (*APOE*) genotype; oily fish intake; omega 3 status; n-3 long chain  
24 polyunsaturated fatty acids; eicosapentaenoic acid (EPA); docosahexaenoic acid (DHA); fatty acid  
25 status; blood lipids.

## 26 Introduction

27 There is convincing evidence that higher intakes of the marine long chain n-3 PUFAs (LC n-  
 28 3 PUFAs) eicosapentaenoic acid (EPA) and docosahexaenoic acid (DHA), are beneficial to  
 29 cardiovascular and cognitive health, acting through a number of biological mechanisms, and that  
 30 the concentration of EPA and DHA present in blood and tissue lipids is correlated positively with  
 31 these effects <sup>1-5</sup>. Oily fish are a good source of EPA and DHA; therefore, national and international  
 32 authorities recommend regular consumption of oily fish such as salmon, mackerel, kippers, sardines,  
 33 herring, trout and fresh tuna, in order to provide approximately 500 mg EPA+DHA per day <sup>6</sup>, with  
 34 higher intakes of LC n-3 PUFAs recommended for those with diagnosed cardiovascular disease <sup>7</sup>.  
 35 However, the associations between intake and blood and tissue status, and therefore physiological  
 36 benefits, are highly variable <sup>8</sup>, and the factors influencing this heterogeneity are not well understood.  
 37 A greater knowledge of determinants of LC n-3 PUFA status could lead to the development of more  
 38 robust, and perhaps subgroup specific, recommendations for EPA and DHA intake.

39 In addition to intake of the specific LC n-3 PUFAs and their precursors, the heterogeneity in  
 40 habitual EPA, docosapentaenoic acid (DPA) and DHA concentrations may be influenced by  
 41 differences in fatty acid metabolism between sexes; females are reported to synthesise EPA, DPA  
 42 and DHA from shorter chain n-3 fatty acids more readily than males <sup>9-13</sup>. Lipid metabolism alters  
 43 with age and becomes dysregulated in obesity, and EPA and DHA concentrations have been  
 44 reported to be affected by increasing BMI <sup>12 14</sup> as well as with age <sup>10-12</sup>. Apolipoprotein E (*APOE*)  
 45 genotype is associated with altered lipid metabolism and transport, with differential responses in  
 46 *APOE4* carriers relative to non-carrier groups <sup>12 14</sup>. Recent reports highlight the importance of  
 47 *APOE* genotype in the response of EPA and DHA to supplementation and have indicated  
 48 interactions between genotype and BMI <sup>14</sup>. In addition, the concentrations of LC n-3 PUFAs in  
 49 individual lipid pools within blood (and in other tissues) differs <sup>15</sup>. However, despite these insights  
 50 from the published literature, the influence of oily fish intake, along with sex, age, BMI and *APOE*  
 51 genotype on EPA, DPA and DHA concentrations in different plasma pools has not been examined  
 52 systematically. Using samples from the FINGEN study <sup>4</sup>, where participants were prospectively  
 53 recruited based on a number of these variables (sex, age, and *APOE* genotype), we have conducted  
 54 such an analysis in a large number of participants to evaluate the independent and interactive impact  
 55 of a number of potential determinants (oily fish intake, sex, age, BMI and *APOE* genotype) on EPA,  
 56 DPA and DHA concentrations in the main plasma lipid fractions.

## 58 Participants and methods

59 The FINGEN study was a multi-centre trial conducted at the Universities of Glasgow,  
 60 Newcastle, Reading and Southampton in the United Kingdom. Three hundred and twelve  
 61 participants were recruited prospectively on the basis of *APOE* genotype (87 were *APOE2*  
 62 homozygotes or *APOE2/APOE3*, 111 were homozygous for *APOE3*, and 114 were *APOE4/APOE3*  
 63 or *APOE4* homozygotes), sex (149 male and 163 female) and age (20 to 71 years, with  
 64 approximately equal numbers in each of the 5 decades)<sup>4</sup>. Data from 306 participants were included  
 65 in the current analysis, with the numbers in each subgroup detailed in **Supplemental Table 1** and  
 66 **Supplemental Table 2**. Exclusion criteria included: diagnosed endocrine dysfunction including  
 67 diabetes or fasting glucose concentration > 6.5 mmol/L, myocardial infarction in the previous 2  
 68 years, the use of medication that may interfere with lipid metabolism, fasting total cholesterol of >  
 69 8.0 mmol/L or TG of > 3.0 mmol/L, a BMI of < 18.5 or > 36.0 kg/m<sup>2</sup>, or currently following a  
 70 weight loss diet. Individuals taking n-3 fatty acid supplements were also excluded. The study was  
 71 approved by the research ethics committee at each of the participating centres and written informed  
 72 consent was obtained from all subjects prior to participation.

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#### 74 **Study design**

75 The FINGEN study was a randomised double blind, placebo controlled, crossover study  
 76 testing two doses of fish oil compared with placebo<sup>4</sup>. Here we evaluate the association of oily fish  
 77 intake, sex, age, BMI and *APOE* genotype with fasting concentrations of EPA, DPA and DHA in  
 78 plasma phosphatidylcholine (PC), non-esterified fatty acids (NEFAs), cholesteryl esters (CEs) and  
 79 triacylglycerols (TGs) at baseline, prior to intervention. Habitual oily fish intake was estimated by  
 80 food frequency questionnaire (FFQ), using self-reported portions completed at baseline. Oily fish  
 81 was defined as salmon, herring, mackerel, fresh tuna, sardines, kippers, and trout.

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#### 83 **Fatty acid analysis**

84 The fatty acid composition of the plasma fractions was determined by gas chromatography.  
 85 Dipentadecanoyl PC, heneicosanoic acid, cholesteryl heptadecanoate and tripentadecanoin internal  
 86 standards were added to the plasma. Total plasma lipid was extracted using chloroform: methanol  
 87 (2:1, v/v) containing butylated hydroxytoluene (50 mg/L) as described by Folch et al<sup>16</sup>, and PC,  
 88 NEFA, CE and TG fractions were separated and isolated by solid phase extraction on aminopropyl  
 89 silica cartridges. CEs and TGs were eluted in a combined fraction with the addition of chloroform.  
 90 PC was then eluted from the cartridge with the addition of chloroform: methanol (60:40 v/v).  
 91 NEFAs were eluted from the cartridge with the addition of chloroform: methanol: glacial acetic acid  
 92 (100:2:2 v/v). CEs and TGs were separated on a hexane primed aminopropyl silica cartridge with  
 93 the addition of hexane to elute CEs, and the addition of hexane: methanol: ethyl acetate (100:5:5

v/v/v) to elute TGs. The fatty acids within the resulting lipid fractions were methylated by the addition of methanol in 2% (v/v) sulphuric acid at 50°C for 2 hours to produce fatty acid methyl esters (FAMES)<sup>17</sup>. FAMES were extracted into hexane and separated in a BPX-70 fused silica capillary column (30 m × 0.25 mm × 25 µm; SGE Analytical Science, United Kingdom) using an Agilent 6890 series gas chromatograph equipped with flame ionisation detection (Agilent Technologies, California, United States). The FAMES were identified by comparison with retention times of 37 FAME and menhaden oil standards run alongside the samples, and quantified with the use of the internal standards using ChemStation software (Agilent Technologies, California, United States) and Microsoft Excel (Microsoft Corporation, Washington, United States). Fatty acid composition data are expressed as absolute concentrations (µg/ml plasma) and as relative concentrations (g/100 g total fatty acid (%)).

## Statistics

Here we report baseline data obtained as part of the previous FINGEN trial<sup>4</sup>. Characteristics for participants included in the baseline analysis are detailed in **Supplemental Table 1** and **Supplemental Table 2**.

Results for the relative (%) and absolute concentrations (µg/ml) of fatty acids are reported for 303 to 306 and 292 to 306 participants in the four plasma lipid fractions. Data were checked for normality by plotting distributions of residuals obtained from general linear model (GLM) analysis of the data, and were analysed appropriately with a univariate GLM following log<sub>10</sub> transformation. All variables were included in the univariate model with individual associations analysed using ‘main effects’ and interaction between age and BMI, age and fish intake, and sex and *APOE* analysed using ‘interaction’ analysis options within the model. *P* values were corrected for multiple analyses using Bonferroni post hoc analysis resulting in a significance value of *P* = 0.006 for whole group analysis and *P* = 0.008 for analyses where males and females were analysed separately. All statistical analyses were conducted using SPSS software (version 21; SPSS Inc, Chicago, IL). Statistical significance was defined as *P* ≤ 0.05. Results are expressed as mean ± SEM or median (25<sup>th</sup>, 75<sup>th</sup> percentiles).

## Results

The group (n = 306) mean age and BMI was 45.1 ± 0.7 y and 25.2 ± 0.2 kg/m<sup>2</sup>, respectively.

Male and female participants were well matched for age, but males had a significantly higher average BMI (*P* < 0.001, **Supplemental Table 1** and **Supplemental Table 2**). There were no sex

differences in the proportion of total dietary energy consumed from fat, saturated fat (SFA), monounsaturated fat (MUFA) or polyunsaturated fat (PUFA) (data not shown). The average oily fish intake was 1.0 portion per week with no association of sex with oily fish intake.

For all three LC n-3 PUFAs, the greatest concentrations were evident in the PC fraction, with median absolute concentrations ( $\mu\text{g/ml}$ ) of 15.1, 11.9 and 44.1 for EPA, DPA and DHA, respectively. The median values for EPA, DPA and DHA for the whole group and  $P$  values for the association of oily fish intake, sex, age, BMI and *APOE* with the plasma concentrations of these fatty acids in the four lipid fractions are presented in **Table 1**. The data according to oily fish intake are shown in **Supplemental Figures 1-4**, while data according to age and BMI are shown in **Table 2 and Supplemental Tables 3-5**, and those according to *APOE* genotype\*sex in **Figures 1-3**.

#### Plasma EPA, DPA and DHA in the group as a whole

**EPA:** The concentration of EPA in plasma CEs and TGs was positively associated with oily fish intake ( $P \leq 0.004$ ), with evidence for positive association in plasma PC also ( $P = 0.018$ ) (**Table 1**). There was evidence for a positive association between EPA and age in plasma PC, CE's and TGs ( $P = 0.021$ ,  $0.019$ , and  $0.034$  respectively) and for the concentration of EPA in CEs to differ by sex ( $P = 0.055$ ), (**Table 2**). A higher concentration of EPA in CEs was observed in males (**Table 2**), and the concentration of EPA in TGs was associated with an *APOE*\*sex interaction ( $P = 0.044$ , data not shown).

**DPA:** The concentration of DPA was positively associated with oily fish intake in plasma TGs ( $P = 0.006$ ), with evidence for positive association in plasma PC also ( $P = 0.022$ ) (**Table 1**). DPA in TGs was positively associated with BMI ( $P = 0.006$ ) (**Table 1**), and there was evidence for the positive association of DPA in NEFAs and CEs with age ( $P = 0.031$  and  $0.007$  respectively, **Table 1**). The concentration of DPA significantly differed by sex with a higher concentration of DPA observed in plasma TGs in males ( $P < 0.001$ ), with a trend in PC also ( $P 0.031$ ) (**Table 1**). There was also a significant *APOE*\*sex interaction for the concentration of DPA in CEs ( $P \leq 0.005$ , data not shown). (**Table 1**),

**DHA:** The concentration of DHA in all plasma lipid fractions was positively associated with oily fish intake ( $P \leq 0.001$ ). There was evidence for a positive association of DHA in TGs with BMI ( $P = 0.020$ ) (**Table 1**) and with age in PC, -CEs and TGs ( $P = 0.037$ ,  $0.039$ , and  $0.050$  respectively, **Table 1**).

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Overall in PC, NEFAs, CEs, and TGs, the highest oily fish consumers (2+ portions of oily fish per week) had 55%, 42%, 52% and 119% higher EPA+DHA, respectively, compared with those reporting no oily fish intake (**Supplemental Figure 4**).

Due to the significant evidence of sex and *APOE*\*sex interactions, subgroup analysis was performed in males and females separately.

Subgroup analysis of plasma EPA, DPA and DHA according to sex

Significance data (*P*) are reported for EPA, DPA and DHA in **Table 2** and median data are reported for EPA, DPA and DHA in **Supplemental Tables 3, 4, and 5** respectively.

**EPA (Table 2, Supplemental Table 3):** The concentration of EPA in plasma TGs was positively associated with oily fish intake in both males and females ( $P \leq 0.008$ ), while the concentration of EPA in PC was positively associated with oily fish intake in females only ( $P \leq 0.004$ ). EPA concentration in TGs was positively associated with age and BMI in females ( $P = 0.006$ ), while EPA in TGs differed by *APOE* genotype in males ( $P = 0.002$ ), with evidence for this in CEs also ( $P = 0.019$ ), (**Figure 1**). A greater concentration of EPA in TGs was observed in male *APOE4* carriers ( $P = 0.002$ ) with evidence for this in PC and CEs also ( $P = 0.019$  and  $0.053$  respectively), (**Figure 1**).

**DPA (Table 2, Supplemental Table 4):** The concentration of DPA in plasma TGs was positively associated with oily fish intake in females ( $P = 0.008$ ). There was evidence for DPA concentration in PC to differ with *APOE* genotype in males ( $P \leq 0.053$ , **Figure 2**) with further analysis revealing evidence for higher concentrations of DPA in PC in *APOE4* allele carriers ( $P = 0.032$ , **Figure 2**).

**DHA (Table 2, Supplemental Table 5):** The concentration of DHA was positively associated with oily fish intake in plasma PC, NEFAs, and TGs in females ( $P \leq 0.002$ ) and plasma PC in males ( $P \leq 0.003$ ), (**Table 2**). There was evidence for DHA in plasma NEFAs to be associated with BMI in females ( $P = 0.010$ , **Table 2**), and for DHA in CEs to differ by *APOE* genotype in males. Further analysis revealed evidence for a higher concentration of DHA in CEs in *APOE4* carriers ( $P = 0.021$ , **Figure 3**).



## 197 Discussion

198 EPA and DHA have been widely reported for their beneficial effects on cardiovascular and  
 199 cognitive health<sup>1-4 18</sup> but a high level of variation in associations between intake and blood and  
 200 tissue status has been observed<sup>8</sup>. The current analysis aimed to identify factors associated with  
 201 concentrations of EPA, DPA and DHA in major lipid fractions in plasma from individuals  
 202 consuming their usual diet in order to identify sources of variation in these concentrations.  
 203 Identification of the contribution that oily fish intake, sex, age, BMI and *APOE* genotype make to  
 204 EPA, DPA and DHA status is important for two reasons. First it will highlight the sources of the  
 205 heterogeneity in status of these fatty acids, contributing to a better understanding of the use of fatty  
 206 acid profiles as a measure of dietary intake amongst different population subgroups. Secondly, it  
 207 may allow the development of sub-group specific recommendations for LC n-3 PUFA intake.

208 The current study reports associations for multiple confounding variables with the relative  
 209 and absolute concentrations of EPA, DPA and DHA in different plasma lipids. The relative  
 210 concentration allows investigation of LC n-3 PUFA concentrations in relation to all other fatty acids  
 211 within the plasma pool (% unit changes), while the absolute concentration allows investigation of  
 212 µg/ml unit changes in LC n-3 PUFAs independently of any other fatty acid within the plasma pool.  
 213 Both ways of expressing the data are useful and informative and both are used in the literature in the  
 214 field. The absolute concentration of a fatty acid within any plasma lipid fraction will be influenced  
 215 by the total concentration of that fraction. The absolute concentration of a particular fatty acid may  
 216 differ between individuals or between sub-groups while the relative concentration of that fatty acid  
 217 may not be different between those individuals or sub-groups. Conversely, the relative  
 218 concentration could be different but the absolute concentration may not be. Plasma lipids are  
 219 involved in transport of fatty acids between tissues where they have different actions depending  
 220 upon their structure. Hence, the absolute concentration of a fatty acid in a plasma lipid reflects the  
 221 exposure of tissues to that fatty acid and hence is likely to be a meaningful way of reporting the  
 222 fatty acid. Conversely, fatty acids often compete with one another for metabolism or for function  
 223 and hence the relative concentration of each fatty acid (i.e. %) is also likely to be meaningful.

224 Quantitatively, PC is the main plasma LC n-3 PUFA pool and the current study reports a  
 225 greater relative concentration of EPA+DHA in plasma PC (**Supplemental Figure 4**) in individuals  
 226 consuming 2+ portions of oily fish a week compared to those who reported not consuming oily fish,  
 227 as well as positive associations between EPA, DPA and DHA in other plasma lipid fractions and  
 228 oily fish intake. Positive associations for oily fish intake and EPA and DHA are reported for plasma  
 229 phospholipids<sup>19-21</sup> which are confirmed by data from the current analysis which shows 55% higher  
 230 EPA+DHA in plasma PC in those consuming two portions of oily fish (each 150 g) per week

231 compared with those reporting no oily fish consumption. Two portions of oily fish supply about 4-5  
 232 g of EPA+DHA per week, equivalent to 600-700 mg per day<sup>22 23</sup>. Previous studies report  
 233 comparable increases of 81% in plasma phospholipid EPA+DHA, and 8.8 µg/ml and 8.5 µg/ml in  
 234 total plasma EPA and DHA respectively following 16 week consumption of oily fish providing 485  
 235 mg EPA+DHA per day<sup>20</sup> and 6 week consumption of oily fish providing 927 mg EPA+DHA per  
 236 day, respectively<sup>21</sup>. Overall, the findings of the current analysis support existing reports that oily  
 237 fish intake is associated with, and at a population level is the main determinant of, LC n-3 PUFAs  
 238 in all major blood lipid pools, which may therefore be used as biomarkers of oily fish intake<sup>4 19 24 25</sup>.  
 239 Our analysis does not clearly indicate which plasma lipid fraction would best reflect dietary intake  
 240 of EPA and DHA, since, in general all four plasma lipid fractions showed dose-dependent increases  
 241 in EPA and DHA concentration (both absolute and relative) with increasing frequency of oily fish  
 242 consumption.

243         There is some evidence that age influences the concentration of EPA and DHA in various  
 244 plasma fatty acid fractions,<sup>10</sup> which has been attributed in part to higher habitual fish intake with  
 245 increasing age. Oily fish intake was controlled for in the current statistical analysis, allowing clearer  
 246 attribution of any observed associations of age with EPA, DPA and DHA concentrations to altered  
 247 metabolism and not to dietary differences in intakes of oily fish. Any influence of *APOE* group  
 248 distribution was also ruled out as, despite a greater number of individuals aged 50-59 yr being  
 249 included in the current analysis, there was no significant difference in the distribution of *APO E2*,  
 250 *E3* and *E4* genotypes between age groups (data not shown). A 28 d stable isotope tracer study in  
 251 young (mean age 27 y) vs older (mean age 77 y) adults reported a 1-2 fold greater enrichment of  
 252 <sup>13</sup>C-DHA in plasma phospholipids and CEs in the older age group, suggesting a medium term age-  
 253 related difference in DHA homeostasis associated with accumulation of DHA in the circulation in  
 254 older people<sup>26</sup>. The findings of the current analysis support reports of increased plasma DHA with  
 255 increasing age<sup>11 27 28</sup> and we further also report positive associations between age and EPA and  
 256 DPA, suggesting LC n-3 PUFAs accumulate in plasma pools during ageing. However, this may in  
 257 part be due to an increase in circulating cholesterol and CE with age (**Table 3**). Evidence of positive  
 258 associations of plasma total cholesterol with age dates back to the late 1970s<sup>29</sup>, and these have been  
 259 reported in both males and females<sup>30</sup>. Increased circulating LDL (**Table 3**) may be reflected in  
 260 higher absolute total PC and CE concentrations with age ( $P = 0.008$  and  $0.018$ , age 20-29 vs 60+ yr  
 261 for PC and CE respectively, data not shown) and we observed that total cholesterol (TC) and LDL-  
 262 cholesterol (LDLC) concentrations were significantly positively correlated with LC n-3 PUFA  
 263 concentrations in PC (TC,  $P = <0.001$ ,  $0.003$ ,  $0.027$ ; LDLC  $P = <0.001$ ,  $<0.001$ ,  $0.003$ , absolute  
 264 EPA, DPA and DHA respectively, data not shown), and that TC, LDLC and high density  
 265 lipoprotein cholesterol (HDL) concentrations were positively correlated with LC n-3 PUFA in

CEs (TC,  $P = <0.001$ , 0.002 absolute EPA and DHA respectively, LDL,  $P = <0.001$ , 0.046, 0.055 absolute EPA, relative DPA and DHA respectively, HDLC,  $P = 0.046$  relative DPA, data not shown). These data suggest CE levels may play a significant role in the association of age with LC n-3 PUFAs reported in this analysis.

Insulin has a role in the regulation of genes involved in whole body lipid homeostasis including in the removal of lipids from the circulation<sup>31</sup>; in cases of insulin resistance, such removal can be compromised. The occurrence of insulin resistance is reported to rise with increasing age and BMI and despite individuals with diabetes or a fasting glucose concentration  $> 6.5$  mmol/L being excluded from the current analysis, differences in fasting glucose were still evident between age and BMI groups (glucose positively correlated with age and BMI;  $P < 0.001$  both, data not shown). Thus, insulin resistance may contribute to the higher EPA and DPA concentrations in plasma lipid pools observed with increasing age and BMI.

Increasing body fatness and obesity influence many aspects of fatty acid and lipid metabolism and contribute to disease states such as hypertriglyceridemia, diabetes, and fatty liver disease<sup>12 32</sup>; loss of insulin sensitivity with increasing adiposity results in adipose tissue lipolysis and associated higher plasma NEFA concentrations<sup>32-34</sup>. In the current analysis, there was no correlation between total NEFA concentrations and BMI (data not shown); however, significant, but complex, associations between BMI and LC n-3 PUFAs were evident in plasma TGs, with an overall trend towards lower relative concentrations of EPA and DHA with increasing BMI, which is consistent with previous observations<sup>33 35</sup>. Increased  $\beta$ -oxidation of DHA associated with increased BMI may in part explain lower proportions of LC n-3 PUFAs in TGs<sup>36</sup> although altered TG synthesis and/or selective tissue uptake and partitioning in obesity may also be involved. We observed no association of BMI with absolute plasma concentrations of LC n-3 PUFAs and suggest the lower relative concentrations (i.e., %) of EPA and DHA are likely to be offset by increases in total TG concentrations with increasing BMI.

The proteins encoded by the *APOE* gene play a major role in the transport and metabolism of lipids via interaction with LDL receptors (LDLRs). Two common polymorphisms (rs7412 and rs429358) of the *APOE* gene in humans result in three protein isoforms, APOE2, E3 and E4. APOE2 and APOE3 are found in the circulation mainly on high density lipoproteins (HDLs) whereas APOE4 is found preferentially on very low density lipoproteins (VLDLs) with lower concentrations residing on HDLs<sup>37</sup>. The *APOE4* allele has been associated with reduced longevity<sup>38</sup>, and enhanced risk of cardiovascular disease<sup>39</sup> and Alzheimer's disease<sup>40</sup>. Although centrally involved in fatty acid transport and handling in plasma and tissues (and in particular within the brain where *APOE* is almost the only apolipoprotein present), the impact of *APOE* genotype on these processes, and the contribution of dysregulated EPA and DHA metabolism to disease risk is

301 unknown. However,  $^{13}\text{C}$ -DHA labelling studies provide evidence that DHA metabolism is  
 302 disturbed in those who are *APOE4* carriers <sup>41</sup>.

303 In the current analysis, *APOE4* carriers had significantly higher concentrations of TC and  
 304 HDLC, and lower concentrations of LDLC (**Table 3**); however, sex\**APOE* genotype interactions  
 305 were evident and in male *APOE4* carriers we observed to have significantly higher concentrations  
 306 of LDLC as well as of total CEs (data not shown). One advantage of investigating associations in  
 307 individual plasma lipid classes as opposed to total lipid is that possible effects of *APOE* and  
 308 lipoprotein transport and metabolism may be more easily identified. If the associations between  
 309 *APOE* and LC n-3 PUFAs are seen to occur in lipid pools which are predominantly related to LDL  
 310 and VLDL particles, they may reflect the dysregulation in lipoprotein handling in people with the  
 311 *E4* allele. However, if the associations between LC n-3 PUFA and *APOE* genotype are seen to  
 312 occur across all lipid pools, they may be indicative of alternative mechanisms. Further subgroup  
 313 analysis indicated higher EPA, DPA and DHA concentrations in CEs, EPA and DPA in PC, and  
 314 EPA in TGs in male *APOE4* carriers relative to the non-carrier groups. The higher EPA and DHA  
 315 may reflect higher overall CE and PC concentrations; however, the lack of association between  
 316 *APOE* genotype and fatty acid concentrations in females is suggestive of a sex specific association  
 317 independent of CE and PC metabolism.

318 Interestingly, we have previously reported *APOE* genotype mediated differences in the  
 319 response of plasma EPA and DHA to a fish oil supplement given over eight weeks in males, with  
 320 lower enrichment in total lipid and phospholipid EPA and DHA in *APOE4* carriers relative to the  
 321 wild-type *APOE3/E3* genotype, but only in overweight participants <sup>14</sup>. The aetiology of these  
 322 associations with LC n-3 PUFA metabolism is currently unknown. As with the association with age,  
 323 higher plasma LC n-3 PUFAs in *APOE4* carriers may reflect reduced tissue uptake and DHA  
 324 accumulating in the circulation. Although lower overall concentrations of *APOE* were observed in  
 325 *APOE4* carriers (data not shown) no difference in plasma *APOE* concentrations were evident  
 326 between sexes, which potentially could have contributed to the differential associations of *APOE*  
 327 genotype with EPA, DPA and DHA concentrations. The preferential binding of VLDL by *APOE4*  
 328 and possible associations of *APOE* genotype with PC and CE synthesis and cellular uptake of EPA  
 329 and DHA via the LDLR family, LDLR concentrations and specific LC PUFA transporters such as  
 330 the MFSD2A transporter in the brain <sup>42</sup> may be involved, and are worthy of future investigations.  
 331 Associations between sex and the activity of these transporters and receptors would also be of  
 332 interest, along with sex and *APOE* associations with FADS and ELOVL genes which encode  
 333 desaturation and elongation enzymes required for the synthesis of LC n-3 PUFAs. Differential  
 334 synthesis of EPA and DHA has been reported between sexes; Pawlosky *et al* report greater ability  
 335 of females to convert ALA to DHA through increased conversion of DPA to DHA compared to

336 males when consuming a beef based diet. These results were not observed when consuming a fish  
 337 based diet in which the capacity to convert DPA to DHA was equal between males and females.  
 338 These findings suggest LC n-3 PUFA metabolism in females may be more sensitive to dietary  
 339 alterations or may be affected by hormonal regulation<sup>43</sup>. Indeed there is evidence for up-regulation  
 340 of the desaturase-elongase pathway via oestrogenic actions resulting in increased conversion of  
 341 ALA to EPA<sup>19 44 45</sup> and to DHA<sup>11 13 46</sup> indicating significant effects of female sex hormones on the  
 342 metabolism of LC n-3 PUFAs. Consistent with these observations, there is evidence for an increase  
 343 in DHA in relation to EPA and DPA at baseline and in response to EPA+DHA intake in females  
 344 compared to males<sup>47 48</sup>. The current analysis further reports lower concentrations of both DPA (-  
 345 36% lower absolute concentration in TGs) and EPA (20% lower absolute concentration in TGs) in  
 346 females but does not report higher concentrations of DHA in females or find a significant effect of  
 347 sex on the ratio of DPA: DHA ( $P > 0.50$ , data not shown). However, these results are also in  
 348 contrast to other reports describing increased concentrations of EPA and DHA in females<sup>19 44 45</sup>.  
 349 These data from the current analysis suggest investigation into associations between sex, *APOE*,  
 350 and fatty acid synthesis enzymes and transporters would be of worthwhile to further understand the  
 351 mechanisms by which these associations occur.

352

353 In conclusion, we report concentrations of EPA, DPA and DHA to vary across *APOE*  
 354 genotype and that sex is an important factor to consider when evaluating LC n-3 PUFA  
 355 concentrations in these genotypic subgroups. Our results also confirm that concentrations of EPA,  
 356 DPA and DHA in plasma pools are suitable population markers of oily fish consumption and show  
 357 that age and sex are important contributors to the variation in EPA, DPA and DHA concentrations  
 358 in plasma lipids independent of *APOE* genotype. These variables should be considered when  
 359 interpreting LC n-3 PUFA concentrations as a marker of dietary intake and when suggesting dietary  
 360 LC n-3 PUFA recommendations to ensure benefits are achieved across population subgroups.  
 361 Investigation into the handling of supplemental EPA and DHA in these subgroups is to be  
 362 addressed in a further publication and could provide the basis for more detailed advice. However,  
 363 the aetiology and physiological significance of the interaction between sex and *APOE* genotype and  
 364 its association with EPA, DPA and DHA status still requires further investigation.

365

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371

**372 Authors' responsibilities**

373 The authors' responsibilities were as follows: GL, CKA, JCM, CJP, PCC and AMM (the study  
374 management group) were responsible for designing the original FINGEN study and supervising all  
375 aspects of the reported work; EAM, BMK, PJC and CKA recruited and screened volunteers, carried  
376 out the intervention, collected the blood samples and collected the anthropometric, questionnaire  
377 and compliance data; HLF conducted the laboratory analysis reported herein; HLF and MI  
378 conducted statistical analysis; HLF wrote the draft of the manuscript; all authors contributed to the  
379 final version of the manuscript.

380

**381 Conflicts of interest**

382 PCC is an advisor to Pronova BioPharma, Aker Biomarine, Smartfish, Sancilio, Solutex, Dutch  
383 State Mines, Cargill and Danone/Nutricia. None of the other authors has any conflict to declare.

384

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503



	%	%	%	%	µg/ml	µg/ml	µg/ml	µg/ml
Median	2.86	1.1	0.46	0.61	44.13	2.07	9.01	4.28
(25th, 75th percentile)	(2.08, 3.93)	(0.80, 1.54)	(0.32, 0.61)	(0.39, 0.98)	(29.94, 57.68)	(1.43, 3.18)	(5.88, 12.59)	(2.38, 7.19)
Oily Fish Intake <sup>2</sup>	<0.001	<0.001	0.001	<0.001	<0.001	0.002	0.045	<0.001
Sex	-	-	-	-	-	-	-	-
Age <sup>3</sup>	0.037	-	-	-	0.043	-	0.039	0.050
BMI <sup>4</sup>	-	-	-	0.02	-	-	-	-

EPA, eicosapentaenoic acid; DPA, docosapentaenoic acid; DHA, docosahexaenoic acid; PC, phosphatidylcholine; NEFAs, non-esterified fatty acids; CEs, cholesteryl esters; TGs, triacylglycerol.

<sup>1</sup> *P* values obtained using log<sub>10</sub> data in univariate general linear model analysis. Individual associations were investigated for by the addition of all other variables as covariates, controlling for any associations between confounding variables that may influence the dependant variable. The resulting *P* values are therefore reflective of the sole association between the variable of interest and the dependant variable.

<sup>2</sup> Oily fish intake: 0 portions/week, 0.1-0.99/week, 1.0-1.99/week, and 2+/week. Oily fish defined as: salmon, herring, mackerel, fresh tuna, sardines, kippers, and trout.

<sup>3</sup> Age: 20-29y, 30-39y, 40-49y, 50-59y, 60+y.

<sup>4</sup> BMI: Normal weight = 18-25 (kg/m<sup>2</sup>), Overweight = 25.1-30 (kg/m<sup>2</sup>) and Obese = 30.1-46 (kg/m<sup>2</sup>).

**Table 2**

Statistical significance (*P*) of the associations between oily fish intake, sex, age, BMI and LC n-3 PUFAs in males and females<sup>1</sup>

		<b>MALES</b>							
		PC <i>P</i> <sup>1</sup>	NEFAs <i>P</i> <sup>1</sup>	CEs <i>P</i> <sup>1</sup>	TGs <i>P</i> <sup>1</sup>	PC <i>P</i> <sup>1</sup>	NEFAs <i>P</i> <sup>1</sup>	CEs <i>P</i> <sup>1</sup>	TGs <i>P</i> <sup>1</sup>
		%	%	%	%	µg/ml	µg/ml	µg/ml	µg/ml
EPA	Oily fish intake <sup>2</sup>	-	-	-	0.028	0.062	0.061	-	0.008
	Age <sup>3</sup>	-	-	-	-	-	-	0.058	0.019
	BMI <sup>4</sup>	-	-	-	0.014	-	-	-	-
DPA	Oily fish intake <sup>2</sup>	-	-	NS	-	0.066	0.026	-	-
	Age <sup>3</sup>	-	-	0.068	-	-	-	0.012	-
	BMI <sup>4</sup>	-	-	-	-	-	-	-	-
DHA	Oily fish intake <sup>2</sup>	0.003	0.023	0.016	-	0.002	0.014	-	-
	Age <sup>3</sup>	-	-	0.011	-	-	0.024	0.005	-
	BMI <sup>4</sup>	-	-	-	-	-	-	-	-
		<b>FEMALES</b>							
		PC <i>P</i> <sup>1</sup>	NEFAs <i>P</i> <sup>1</sup>	CEs <i>P</i> <sup>1</sup>	TGs <i>P</i> <sup>1</sup>	PC <i>P</i> <sup>1</sup>	NEFAs <i>P</i> <sup>1</sup>	CEs <i>P</i> <sup>1</sup>	TGs <i>P</i> <sup>1</sup>
		%	%	%	%	µg/ml	µg/ml	µg/ml	µg/ml
EPA	Oily fish intake <sup>2</sup>	0.004	-	0.009	<0.001	0.003	-	-	<0.001
	Age <sup>3</sup>	0.04	-	-	-	0.039	-	-	0.006
	BMI <sup>4</sup>	-	-	-	-	-	0.052	-	0.006
DPA	Oily fish intake <sup>2</sup>	-	-	-	0.008	-	-	-	0.067
	Age <sup>3</sup>	-	-	-	-	-	-	-	-
	BMI <sup>4</sup>	-	-	-	-	-	-	-	-
DHA	Oily fish intake <sup>2</sup>	0.001	0.001	0.003	<0.001	<0.001	0.048	-	0.001

Age <sup>3</sup>	0.035	-	-	-	0.047	-	-	0.032
BMI <sup>4</sup>	-	-	-	-	-	0.010	-	-

EPA, eicosapentaenoic acid; DPA, docosapentaenoic acid; DHA, docosahexaenoic acid; PC, phosphatidylcholine; NEFAs, non-esterified fatty acids; CEs, cholesteryl esters; TGs, triacylglycerol.

<sup>1</sup> *P* values obtained using log<sub>10</sub> data in univariate general linear model analysis. Individual associations were investigated for by the addition of all other variables as covariates, controlling for any associations between confounding variables that may influence the dependant variable. The resulting *P* values are therefore reflective of the sole association between the variable of interest and the dependant variable.

<sup>2</sup> Oily fish intake: 0 portions/week, 0.1-0.99/week, 1.0-1.99/week, and 2+ /week. Oily fish defined as: salmon, herring, mackerel, fresh tuna, sardines, kippers, and trout.

<sup>3</sup> Age: 20-29y, 30-39y, 40-49y, 50-59y, 60+y.

<sup>4</sup> BMI: Normal weight = 18-25 (kg/m<sup>2</sup>), Overweight = 25.1-30 (kg/m<sup>2</sup>) and Obese = 30.1-46 (kg/m<sup>2</sup>).

**TABLE 3**

Blood cholesterol (mmol/l) concentration according to sex, age, BMI, *APOE* genotype and oily fish intake

	TC		HDLc		LDLC	
	Mean	SEM	Mean	SEM	Mean	SEM
Male	5.16	0.08	1.26	0.02	3.34	0.07
Female	5.16	0.08	1.61	0.03	3.17	0.07
<sup>1</sup> <i>P</i>	NS		<0.001		NS	
Age group						
20-29y	4.39	0.14	1.46	0.05	2.56	0.13
30-39y	4.68	0.1	1.34	0.04	2.93	0.1
40-49y	5.34	0.11	1.47	0.04	3.41	0.09
50-59y	5.57	0.1	1.45	0.05	3.62	0.09
60+y	5.59	0.13	1.49	0.06	3.52	0.1
<sup>2</sup> <i>P</i>	<0.001		NS		<0.001	
<sup>3</sup> BMI group						
Normal weight	4.91	0.08	1.57	0.03	2.98	0.07
Overweight	5.33	0.08	1.34	0.03	3.43	0.07
Obese	5.63	0.18	1.20	0.05	3.77	0.17
<sup>2</sup> <i>P</i>	<0.001		<0.001		<0.001	
<i>APOE</i> genotype <sup>4</sup>						
<i>E2</i>	4.71	0.09	1.54	0.04	2.76	0.08
<i>E3</i>	5.19	0.1	1.43	0.04	3.31	0.08
<i>E4</i>	5.46	0.08	1.37	0.03	3.57	0.07
<sup>1</sup> <i>P</i>	<0.001		0.006		<0.001	
Oily fish intake <sup>5</sup>						
0/wk	4.9	0.12	1.41	0.04	3.11	0.1
0.1-0.99/wk	5.21	0.09	1.44	0.03	3.25	0.07
1-1.99/wk	5.3	0.12	1.48	0.05	3.38	0.1
2+/wk	5.16	0.15	1.41	0.07	3.28	0.15
<sup>2</sup> <i>P</i>	NS		NS		NS	NS

TC, Total cholesterol; HDLC, high density lipoprotein cholesterol; LDLC, low density lipoprotein cholesterol.

<sup>1</sup>*P* values obtained from one-way ANOVA model.

<sup>2</sup>*P* values obtained from Pearson's correlation model.

<sup>3</sup> BMI: Normal weight = 18-25 (kg/m<sup>2</sup>), Overweight = 25.1-30 (kg/m<sup>2</sup>) and Obese = 30.1-46 (kg/m<sup>2</sup>).

<sup>4</sup> *APOE* genotype: *E2* (*E2/E2* and *E2/E3*), *E3* (*E3/E3*), and *E4* (*E3/E4* and *E4/E4*) .

<sup>5</sup> Oily fish defined as: salmon, herring, mackerel, fresh tuna, sardines, kippers, and trout.

**FIGURE 1** Absolute concentrations ( $\mu\text{g/ml}$ ) of eicosapentaenoic acid (EPA) in phosphatidylcholine (PC), non-esterified fatty acids (NEFAs), cholesteryl esters (CEs) and triacylglycerols (TGs) lipid fractions in male and female subjects according to *APOE* genotype. Distribution of participants in each *APOE* allele group are as follows; PC: Males: E2/E2 = 1, E2/E3 = 32, E3/E3 = 40, E3/E4 = 51, and E4/E4 = 2. Total: 126. PC: Females: E2/E2 = 3, E2/E3 = 39, E3/E3 = 44, E3/E4 = 43, and E4/E4 = 10. Total: 139. NEFAs: Males: E2/E2 = 2, E2/E3 = 29, E3/E3 = 45, E3/E4 = 50, and E4/E4 = 2. Total: 128. NEFAs: Females: E2/E2 = 3, E2/E3 = 42, E3/E3 = 44, E3/E4 = 45, and E4/E4 = 10. Total: 144. CEs: Males: E2/E2 = 2, E2/E3 = 33, E3/E3 = 50, E3/E4 = 52, and E4/E4 = 2. Total: 139. CEs: Females: E2/E2 = 3, E2/E3 = 44, E3/E3 = 48, E3/E4 = 49, and E4/E4 = 10. Total: 154. TGs: Males: E2/E2 = 2, E2/E3 = 32, E3/E3 = 50, E3/E4 = 53, and E4/E4 = 2. Total: 139. TGs: Females: E2/E2 = 3, E2/E3 = 45, E3/E3 = 49, E3/E4 = 48, and E4/E4 = 10. Total: 155. \*  $P < 0.050$ , and \*\*  $P > 0.050$  but  $< 0.060$ .  $P$  values were obtained using log-10 data in univariate general linear model (GLM) analysis controlling for covariates (age, BMI, and oily fish intake). Where there was a significant association with *APOE* genotype, significance between specific *APOE* alleles was assessed using parameter estimates obtained from the GLM results.



**FIGURE 2** Absolute concentrations ( $\mu\text{g/ml}$ ) of docosapentaenoic acid (DPA) in phosphatidylcholine (PC), non-esterified fatty acids (NEFAs), cholesteryl esters (CEs) and triacylglycerols (TGs) lipid fractions in male and female subjects according to *APOE* genotype. Distribution of participants in each *APOE* allele group are as follows; PC: Males: E2/E2 = 1, E2/E3 = 32, E3/E3 = 40, E3/E4 = 51, and E4/E4 = 2. Total: 126. PC: Females: E2/E2 = 3, E2/E3 = 39, E3/E3 = 44, E3/E4 = 43, and E4/E4 = 10. Total: 139. NEFAs: Males: E2/E2 = 2, E2/E3 = 29, E3/E3 = 45, E3/E4 = 50, and E4/E4 = 2. Total: 128. NEFAs: Females: E2/E2 = 3, E2/E3 = 42, E3/E3 = 44, E3/E4 = 45, and E4/E4 = 10. Total: 144. CEs: Males: E2/E2 = 2, E2/E3 = 33, E3/E3 = 50, E3/E4 = 52, and E4/E4 = 2. Total: 139. CEs: Females: E2/E2 = 3, E2/E3 = 44, E3/E3 = 48, E3/E4 = 49, and E4/E4 = 10. Total: 154. TGs: Males: E2/E2 = 2, E2/E3 = 32, E3/E3 = 50, E3/E4 = 53, and E4/E4 = 2. Total: 139. TGs: Females: E2/E2 = 3, E2/E3 = 45, E3/E3 = 49, E3/E4 = 48, and E4/E4 = 10. Total: 155. \*  $P < 0.050$ . \*\*  $P > 0.050$  but  $< 0.070$ .  $P$  values were obtained using log-10 data in univariate general linear model (GLM) analysis controlling for covariates (age, BMI, and oily fish intake). Where there was a significant association with *APOE* genotype, significance between specific *APOE* alleles was assessed using parameter estimates obtained from the GLM results.

**FIGURE 3** Absolute concentrations ( $\mu\text{g/ml}$ ) of docosahexaenoic acid (DHA) in phosphatidylcholine (PC), non-esterified fatty acids (NEFAs), cholesteryl esters (CEs) and triacylglycerols (TGs) lipid fractions in male and female subjects according to *APOE* genotype. Distribution of participants in each *APOE* allele group are as follows; PC: Males: E2/E2 = 1, E2/E3 = 32, E3/E3 = 40, E3/E4 = 51, and E4/E4 = 2. Total: 126. PC: Females: E2/E2 = 3, E2/E3 = 39, E3/E3 = 44, E3/E4 = 43, and E4/E4 = 10. Total: 139. NEFAs: Males: E2/E2 = 2, E2/E3 = 29, E3/E3 = 45, E3/E4 = 50, and E4/E4 = 2. Total: 128. NEFAs: Females: E2/E2 = 3, E2/E3 = 42, E3/E3 = 44, E3/E4 = 45, and E4/E4 = 10. Total: 144. CEs: Males: E2/E2 = 2, E2/E3 = 33, E3/E3 = 50, E3/E4 = 52, and E4/E4 = 2. Total: 139. CEs: Females: E2/E2 = 3, E2/E3 = 44, E3/E3 = 48, E3/E4 = 49, and E4/E4 = 10. Total: 154. TGs: Males: E2/E2 = 2, E2/E3 = 32, E3/E3 = 50, E3/E4 = 53, and E4/E4 = 2. Total: 139. TGs: Females: E2/E2 = 3, E2/E3 = 45, E3/E3 = 49, E3/E4 = 48, and E4/E4 = 10. Total: 155. \*  $P = 0.021$ .  $P$  values were obtained using log-10 data in univariate general linear model (GLM) analysis controlling for covariates (age, BMI, and oily fish intake). Where there was a significant association with *APOE* genotype, significance between specific *APOE* alleles was assessed using parameter estimates obtained from the GLM results.