Supplementation with high-dose docosahexaenoic acid increases the Omega-3 Index more than high-dose eicosapentaenoic acid

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ABSTRACT

Background: Recent studies suggest that eicosapentaenoic (EPA) and docosahexaenoic (DHA) acids have distinct effects on cardiometabolic risk factors. The Omega-3 Index (O3I), which is calculated as the proportion of EPA and DHA in red blood cell (RBC) membranes, has been inversely associated with the risk of coronary heart diseases and coronary mortality. The objective of this study was to compare the effects of EPA and DHA supplementation on the O3I in men and women with abdominal obesity and subclinical inflammation.

Methods: In a double-blind controlled crossover study, 48 men and 106 women with abdominal obesity and subclinical inflammation were randomized to a sequence of three treatment phases: 1) - 2.7 g/d of EPA, 2) - 2.7 g/d of DHA, and 3) - 3 g/d of corn oil (0 g of EPA+DHA). All supplements were provided as 3×1 g capsules for a total of 3 g/d. The 10-week treatment phases were separated by nine-week washouts. RBC membrane fatty acid composition and O3I were assessed at baseline and the end of each phase. Differences in O3I between treatments were assessed using mixed models for repeated measures.

Results: The increase in the O3I after supplementation with DHA (+5.6% compared with control, P < 0.0001) was significantly greater than after EPA (+3.3% compared with control, P < 0.0001; DHA vs. EPA, P < 0.0001). Compared to control, DHA supplementation decreased (−0.8%, P < 0.0001) while EPA increased (+2.5%, P < 0.0001) proportion of docosapentaenoic acid (DPA) in RBCs (DHA vs. EPA, P < 0.0001). The baseline O3I was higher in women than in men (6.3% vs. 5.8%, P=0.011). The difference between DHA and EPA in increasing the O3I tended to be higher in men than in women (+2.6% vs. +2.2% respectively, P for the treatment by sex interaction=0.0537).

Conclusions: The increase in the O3I is greater with high dose DHA supplementation than with high dose EPA, which is consistent with the greater potency of DHA to modulate cardiometabolic risk factors. The extent to which such differences between EPA and DHA in increasing the O3I relates to long-term cardiovascular risk needs to be investigated in the future.

1. Introduction

Considerable research has been conducted to determine the association between long-chain polyunsaturated omega-3 fatty acids (LCn3-PUFAs) consumption and cardiovascular risk. LCn3-PUFAs modulate a variety of cardiometabolic risk factors such as blood lipids, blood pressure, thrombosis and inflammation [1]. Fatty fish and supplements, often combining eicosapentaenoic and docosahexaenoic acids (EPA and DHA), are the main dietary sources of LCn3-PUFAs. There is emerging evidence suggesting that EPA and DHA exert different effects on blood lipids, blood pressure, and other cardiovascular risk factors. EPA and DHA have different mechanisms of action, with EPA having a more pronounced effect on platelet aggregation and thrombosis compared to DHA. DHA has a more pronounced effect on brain and eye development and function.

Abbreviations: BMI, body mass index; CHD, coronary heart disease; CRP, C-reactive protein; CVD, cardiovascular diseases; C, cholesterol; DHA, docosahexaenoic acid; DPA, docosapentaenoic acid; ELOVL2, ELOVL fatty acid elongase 2; ELOVL5, ELOVL fatty acid elongase 5; EPA, eicosapentaenoic acid; FADS1, fatty acids desaturase 2 or delta-5 fatty acids desaturase; FADS2, fatty acids desaturase 2 or delta-6 fatty acids desaturase; GAPDH, glyceraldehyde 3-phosphate dehydrogenase; HDL, high-density lipoprotein; INAF, Institute of nutrition and functional foods; LCn3-PUFA, long-chain omega-3 polyunsaturated fatty acid; LDL, low-density lipoprotein; MetS, metabolic syndrome; MUFA, monounsaturated fatty acid; O3I, Omega-3 Index; PUFAs, polyunsaturated fatty acids; RBC, red blood cell; SFA, saturated fatty acid; TG, triglyceride.

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lipids and inflammation markers [2,3]. However, such evidence is limited, most studies to date having assessed these effects using a mixture of EPA and DHA in different forms and proportions. Little is known with regard to the specific effects of EPA and DHA on metabolic pathways and biological processes underlying cardiometabolic health in humans.

The fatty acid composition of cell membranes influences their physico-chemical properties and, ultimately, organ functions [1,4]. The Omega-3 Index (O3I), which is calculated as the relative content of red blood cell (RBC) membranes as EPA plus DHA, reflects the phospholipid LCn3-PUFA composition of major organs [4], including cardiac tissue [4,5]. A high O3I (8–12%) has been associated with a lower risk of coronary heart disease (CHD) and coronary mortality in epidemiological studies [6,7]. Supplementation with EPA + DHA is recommended by various health agencies including the American Heart Association for secondary CHD prevention or management of plasma triglycerides (TG) [8]. Yet, whether EPA and DHA have distinct effects on the O3I is currently unknown. Considering that the O3I is modifiable by diet [5], studies are required to compare the effects of different LCn3-PUFAs on this promising clinical tool for the management of diet-related CHD risk.

The objective of this study was to compare the effects of high doses of re-esterified EPA and DHA on the O3I using a randomized double-blind controlled crossover study design, in men and women with abdominal obesity and subclinical inflammation. We hypothesized that the O3I increases more with DHA than with EPA. Based on previous studies of fatty acid metabolism in men and women [9], we also hypothesized that the increase in the O3I with both EPA and DHA is greater among women than among men.

2. Patients and methods

2.1. Study design

This analysis is based on data from a double-blind randomized, controlled crossover study with three treatment phases (1- EPA, 2- DHA and 3- corn oil as control), for which the primary outcome was the change in C-reactive concentrations (CRP) concentrations. Details of the study design and results of primary analyses have been published previously [10]. Briefly, each treatment phase had a median duration of 10 weeks and were separated by a nine-week washout. Randomization of participants to one of six treatment sequences was performed using an in-house computer program and was stratified by sex. Participants were supplemented with three identical 1 g capsules of > 90% purified LCn3-PUFA per day providing either 2.7 g/d EPA or 2.7 g/d DHA. Corn oil was used as a control (0 g/d EPA + DHA). LCn3-PUFA supplements were formulated as re-esterified TG and provided by Douglas Laboratories. Participants were instructed to maintain a constant body weight during the course of the study. They were also counseled on how to exclude fatty fish (including salmon, tuna, mackerel, and herring), other LCn3-PUFA supplements, flax products, walnuts, and LCn3-PUFA-enriched products during the three study phases.

2.2. Study population

Primary eligibility criteria were to have abdominal obesity based on the International Diabetes Federation sex specific cut-offs (≥ 80 cm for women, ≥ 94 cm for men) [11] in combination with a screening plasma CRP concentration ≥ 1 mg/L but < 10 mg/L. Subjects had to be otherwise healthy. Adult subjects (18 and 70 years of age) were recruited at the Institute of Nutrition and Functional Foods (INAF). Body weight had to be stable for at least three months prior to randomization. Exclusion criteria were plasma CRP > 10 mg/L at screening, extreme dyslipidemias such as familial hypercholesterolemia, having a personal history of cardiovascular diseases (CVD) (CHD, cerebrovascular disease or peripheral arterial disease), taking medications or substances known to affect inflammation (e.g. steroids, binging alcohol), and use of LCn3-PUFA supplements within two months of study onset. All participants signed an informed consent document approved by local Ethics Committees at the beginning of the study and the study protocol was registered March 4, 2013 at ClinicalTrials.gov (NCT01810003).

2.3. Anthropometry

Anthropometric measures including waist and hip circumferences were obtained according to standardized procedures [12]. Body composition was measured by Dual-energy X-ray absorptiometry (GE Healthcare, Madison, WI).

2.4. Dietary habits

Food intakes during each phase was monitored using a validated quantitative web-based, self-administered food frequency questionnaire at the end of each treatment phase [13].

2.5. Risk factor assessment

Serum total cholesterol (C), TG and high-density lipoprotein (HDL)-C were assessed on a Roche/Hitachi Modular (Roche Diagnostics, Mannheim, Germany) according to the manufacturer’s specifications and using proprietary reagents. Plasma low-density lipoprotein (LDL)-C concentrations were calculated using the Friedewald Equation. Total C, LDL-C, HDL-C and TG were measured twice on consecutive days at the end of each treatment. The mean of the two measurements were used in the analyses. Treatment-specific baseline values were measured once. All personnel involved in the measurements of the study outcomes were blinded to treatments. Metabolic syndrome (MetS) was defined using the International Diabetes Federation criteria [11].

2.6. RBC membrane fatty acids measurements

RBC membrane fatty acid composition was measured at baseline and at the end of each treatment phase. RBC membrane was analyzed by OmegaQuant Analytics, LLC, (Sioux Falls, South Dakota, United States) according to the Omega-3 Index® methodology as modified from Harris et al. [14]. Fatty acid methyl esters were generated from erythrocytes by transesterification with boron trifluoride and analyzed by gas chromatography. Fatty acids were identified by comparison with a standard mixture of fatty acids characteristics of RBCs. Each fatty acid is expressed as a weight percent of total identified fatty acids after a response factor correction (based on calibration curves) was applied to each fatty acid. The O3I represents the sum of EPA and DHA expressed as a percent of total RBC fatty acids [14]. The RBC composition at baseline and after each phase is presented in Supplemental Table 1. Baseline RBC composition for men and women is presented in Supplemental Table 2.

2.7. Gene expression of polyunsaturated fatty acid metabolism

Fasting fresh blood was collected in PAXgene Blood RNA tubes (Becton Dickinson, Canada) after each treatment in a subsample of 44 randomly selected participants for gene expression analyses. RNA was isolated using a PAXgene RNA-kit according to manufacturer’s instructions (Qiagen, Canada). Quantity of total RNA was measured using a NanoDrop ND-1000 Spectrophotometer (NanoDrop Technologies, Wilmington, DE, USA) and total RNA quality was assayed on an Agilent BioAnalyzer 2100 (Agilent Technologies, Santa Clara, CA, USA). Reverse transcription was performed on 1.5–2 µg total RNA. cDNA corresponding to 20 ng of total RNA was used to perform fluorescent-based Realtime PCR quantification using the LightCycler 480 (Roche Diagnostics, Mannheim, DE). The genes targeted were ELOVL fatty acid elongase 2 and 5 (ELOVL2 and ELOVL5) and fatty
acids desaturase 2, delta-5 and delta-6 (FADS1 and FADS2). Sequence
primers and gene description are available in the Supplemental Table 3.
Values were normalized to expression of the housekeeping gene
glyceraldehyde 3-phosphate dehydrogenase (GAPDH).

2.8. Statistical analyses

Differences in RBC membrane fatty acid composition between
treatments were assessed using the MIXED procedure for repeated
measures in SAS (v9.3, Cary, NC), with treatment as fixed effect and a
compound symmetry or autoregressive covariance matrix to account for
within-subject correlations. The change vs. the control treatment (post
treatment EPA minus control and post treatment DHA minus control)
was used as the dependent variable in all analyses [15]. To be included
in the analyses, subjects had to have completed the control phase plus
at least one of the two treatment phases. Using this approach, the main
treatment effect in the models reflected the direct comparison of EPA
and DHA. Adjustment for multiple comparisons was not necessary as
the main treatment effect had only two levels. In the same model, the
change vs. control for each treatment was tested against the null
hypothesis by the LSMEANS statement in the MIXED procedure.
Potential confounders of the outcome measure response to treatment,
mainly body mass index, age, sex, energy intake were added to models
and results from analyses based on the most parsimonious models (i.e.
retaining only the variables that contributed significantly to variations
in any given study outcome) are shown. We considered sex, sequence of
treatments and baseline values interaction by adding interaction terms
with the main treatment effect into the model statement in the MIXED
procedure. The skewness in the distribution of model residuals was
considered and data were log-transformed when required. Correlations
were tested using Pearson’s correlations coefficients.

3. Results

3.1. Subjects’ characteristics

Of the 173 eligible men and women, 154 were randomized to
treatment sequences. Characteristics of the 154 randomized partici-
pants at screening are shown by sex in Table 1. A total of 123
participants have completed both the EPA and the control
phases. 121 participants have completed both the EPA and the control
phases.

3.2. Fatty acid composition of RBC membranes

Fig. 1 shows the individual O3I response to EPA and DHA
supplementation. The O3I increased with EPA and DHA in all but one
participant. Self-reported compliance for this participant was high
(93%). Both EPA (+3.3%, P < 0.0001) and DHA supplementation
( +5.6%, P < 0.0001) significantly increased the O3I in RBC mem-
branes compared with control (Table 2). DHA led to a greater increase
in the O3I than EPA in all participants (Fig. 1). Thus, the increase in
the O3I with DHA was significantly greater than with EPA (P < 0.0001).
The correlation between the change in O3I with EPA and with DHA (vs.
control) was r = 0.65 (P < 0.0001). DPA levels in RBC increased after
EPA (+2.5%, P < 0.0001) but decreased after DHA (−0.8%, P < 0.0001)
compared with control (EPA vs. DHA, P < 0.0001, Table 2).

The increase in O3I was not statistically different between men and
women after EPA (+3.1% and +3.4% respectively) and after DHA
(+5.7% and +5.6% respectively) (Fig. 2). However, the difference
between DHA and EPA in increasing the O3I tended to be higher in men
than in women (+2.6% vs. +2.2% respectively, P for the treatment-by-
sex interaction=0.054). This apparent difference between men and
women was independent of baseline EPA (P = 0.11) and DHA (P = 0.19)
levels as well as of the baseline O3I (P = 0.12, not shown).

The change in the O3I was inversely correlated with the correspond-
ing change in TG after both EPA and DHA (r = −0.21, P = 0.017 and
r = −0.24, P = 0.0076 respectively) and correlated positively with the
change in LDL-C after DHA only (r = 0.19, P = 0.039 for DHA and
r = −0.02, P = 0.86 for EPA). Changes in the O3I with EPA and DHA did
not correlate significantly with changes in other biomarkers, including
inflammation markers. Finally, there was no carryover effect of any
of the baseline O3I (not shown).

3.3. Gene expression of polyunsaturated fatty acid metabolism

There was no significant difference in the expression of ELOVL2 and
ELOVL5 and FADS1 and FADS2 in whole blood cells between EPA and
DHA (Table 3). EPA supplementation increased the expression of
ELOVL2 by 24% compared with control (P = 0.0084).

4. Discussion

A high O3I, which reflects a relatively high content of EPA and DHA
in the membranes of RBCs, has been associated with a lower risk of
CHD and mortality in observational studies [6,7]. While there is
emerging evidence suggesting that DHA may be more potent than
EPA in modifying cardiometabolic risk [2,3], their respective impacts
on the O3I have not been thoroughly examined. To the best of our
knowledge, this is the first randomized double-blind controlled cross-
over trial to show that the increase in O3I is significantly greater after
supplementation with high dose DHA (2.7 g/d) than with a comparable
dose of EPA.

Grimsgaard et al. [16] found in a parallel study that the increase in
serum phospholipid EPA (+4.7%) was greater after supplementation with 3.8 g/d of ethyl ester EPA for 7 weeks than the corresponding increase of serum phospholipid DHA (+3.2%) after a supplementation of with 3.6 g/d of ethyl ester DHA. More recently, Tsunoda et al. [17] have shown that the increase in EPA (+1.9%) in peripheral blood mononuclear cells after supplementation with 1.8 g/d of EPA compared to an olive oil control was similar to the corresponding increase in DHA (+2.0%) after 1.8 g/d of DHA supplementation. Authors of these two studies did not report the changes in O3I with EPA and DHA supplementation. Our data further indicated that the relative content of DPA in RBC was increased after EPA and reduced after DHA compared with control. This is consistent with results from Tsunoda et al. [17] This increase in EPA levels after supplementation with EPA was also consistent with the observed increase in the expression of ELOVL2, which catalyzes the elongation of EPA to DPA. Since DPA is not accounted for in the calculation of the O3I, “disappearance” of EPA in RBC through in vivo elongation to DPA does in large part explain the smaller increase in O3I with EPA compared with DHA in our study. In fact, an index based on the sum of EPA, DPA and DHA in RBCs would have increased more after high dose supplementation with EPA than after supplementation with a corresponding dose of DHA (+5.8% vs. +4.8% respectively, P < 0.001, not shown).

Levels of DPA in RBCs have been inversely associated with TG and CRP concentrations in healthy adults following supplementation with LCn3-PUFAs (EPA + DPA + DHA) [18], suggesting that DPA may also be partly responsible for some of the biological and cardioprotective effects attributed to LCn3-PUFA [19]. Lower levels of DPA in serum or plasma have been associated with greater risk of acute myocardial infarction [20], total mortality [21], cardiovascular mortality [21] and nonfatal myocardial infarction [22]. DPA concentrations in different tissues (total plasma lipids, plasma or RBC phospholipids and adipose tissue) have also been inversely associated with fatal coronary heart disease in a recent meta-analysis of 19 epidemiological studies (total N = 45,637) [23]. On the other hand, data from the present study indicate that the correlation between the O3I based on EPA and DHA and the sum of EPA + DPA + DHA in RBC is 0.97 after DHA supplementation, 0.93 after EPA supplementation and 0.92 after the control (not shown, P < 0.0001 for all). Such strong correlations suggest little added contribution of DPA to the predictive value of the O3I. Consistent with this, addition of DPA did not significantly improve the mortality risk prediction of the O3I in the Women’s Health Initiative Memory Study [24]. Although a growing body of evidence suggests that DPA may be a highly bioactive compound, studies are needed to dissect out its contribution to CHD risk from that of EPA and DHA, which are also strongly correlated with CHD risk [25].

ELOVL and FADS are the enzymes responsible of the elongation of EPA to DPA and DPA to DHA. Grimsgaard et al. [16] estimated the change in the activities of the FADS1 and FADS2 after seven weeks of DHA (3.6 g/d), EPA (3.8 g/d) or corn oil (4 g/d) supplementation, using ratios of phospholipid fatty acids proportions in serum. Estimated FADS2 activity was increased while estimated FADS1 activity was decreased after DHA supplementation compared with baseline. Supplementation with EPA increased estimated FADS1 activity but had no

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**Table 2**

<table>
<thead>
<tr>
<th>Table 2</th>
<th>Changes in proportions of fatty acids in red blood cell after EPA and DHA supplementation, vs. control.</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>Change with EPA vs. control</td>
</tr>
<tr>
<td>% of total FA</td>
<td></td>
</tr>
<tr>
<td>Total SFA</td>
<td>40.13 ± 0.07</td>
</tr>
<tr>
<td>Myristic acid, 14:0</td>
<td>0.28 ± 0.01</td>
</tr>
<tr>
<td>Palmitic acid, 16:0</td>
<td>2.17 ± 0.10</td>
</tr>
<tr>
<td>Stearic acid, 18:0</td>
<td>17.04 ± 0.08</td>
</tr>
<tr>
<td>Total cis-MUFA</td>
<td>16.27 ± 0.09</td>
</tr>
<tr>
<td>Oleic acid, 18:1, n9</td>
<td>14.92 ± 0.08</td>
</tr>
<tr>
<td>Total trans-MUFA</td>
<td>0.69 ± 0.01</td>
</tr>
<tr>
<td>Total PUFA n6</td>
<td>33.56 ± 0.13</td>
</tr>
<tr>
<td>Linoleic acid, 18:2 n6</td>
<td>11.79 ± 0.13</td>
</tr>
<tr>
<td>Arachidonic acid, 20:4 n6</td>
<td>15.53 ± 0.11</td>
</tr>
<tr>
<td>Total PUFA n3</td>
<td>9.36 ± 0.11</td>
</tr>
<tr>
<td>EPA, 20:5 n3</td>
<td>0.18 ± 0.00</td>
</tr>
<tr>
<td>DPA, 22:5 n3</td>
<td>2.94 ± 0.04</td>
</tr>
<tr>
<td>DHA, 22:6 n3</td>
<td>5.35 ± 0.09</td>
</tr>
<tr>
<td>Omega-3 Index</td>
<td>6.23 ± 0.10</td>
</tr>
</tbody>
</table>

Data are presented as means ± SEM. Total % does not add up to 100% because not all individual fatty acids are shown in this table.

Covariates (sex, age, self-reported energy intake and baseline fatty acid proportions in red blood cells) was included in the mixed models only when they were found to be significant at P < 0.05 in the mixed models. 

DHA: docosahexaenoic acid; DPA: docosapentaenoic acid; EPA: eicosapentaenoic acid; MUFA: monounsaturated fatty acid; PUFA: polyunsaturated fatty acid; SFA: saturated fatty acid.

1 P values for the EPA and DHA change vs. the control treatment, as determined by the LSMEANS statement and tested against the null hypothesis in the mixed models. * P < 0.05, ** P < 0.01, *** P < 0.001. N=123 for DHA, 121 for EPA, 125 for control.

2 Main treatment P values for the comparison between EPA and DHA change vs. control, as determined by the main treatment effect in the mixed models. The mixed model for the main effect comparing ΔEPA and ΔDHA is based on n = 123 observations, with ΔEPA data excluded for 2 participants, due to low compliance.

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**Fig. 2** Difference in change of Omega-3 Index (O3I) after EPA and DHA supplementation, vs. control in men and women (%). Values are means ± SEM. * P < 0.0001 for the treatment-specific effect vs. control for ΔEPA and ΔDHA. The increase in O3I was higher after DHA than after EPA supplementation for both men and women. We found no difference between men and women response to EPA and DHA supplementation.
impact on FADS2 activity compared with baseline [16]. In our study, supplementation with high dose EPA or DHA had no significant impact on either FADS1 or FADS2 gene expression in whole blood cells compared with the control corn oil treatment. It is not entirely surprising that estimated FADS1/2 activities based on fatty acids ratios and analysis of gene expression of these enzymes in whole blood cells yielded inconsistent results among existing studies, considering the complex relationship with actual activities of the corresponding enzymes.

EPA levels in RBC increased significantly after high dose supplementation with DHA. This is consistent with data having demonstrated a dose-dependent relationship between EPA levels in blood and the dose of supplemented DHA, which most likely attributable to retro-conversion of DHA into EPA [26]. The fatty acid content of RBC are expressed as a relative weight of all identified fatty acids. It is therefore difficult to assess if the reduction in the proportion of DPA after DHA supplementation is due to further retro-conversion to EPA, or to an increase in the proportion of other fatty acids.

In vivo conversion rate of ALA to EPA and to DHA is estimated at 8–20% and 0.5–9% respectively [27], with higher rates among women of reproductive age than among men [9]. The regulatory effect of estrogens combined with a lower muscle mass may result in a smaller proportion of ALA being channelled towards beta-oxidation in women, and thus a greater proportion of ALA being converted into EPA and DHA [9]. This suggests potential sex-dependent difference in the response to dietary LnC3-PUFAs. While the effects EPA and DHA supplementation on DPA and DHA levels were similar among men and women in the present study (not shown), women compared with men accumulated more EPA in their RBC after supplementation with EPA (+4.0% vs. +3.5%, respectively, P=0.0002, not shown). Our study further showed that the increase in the O3I with EPA and DHA tended to be higher in men than in women. Self-reported fish consumption and compliance were similar between men and women (not shown) and thus differences in the O3I response to EPA and DHA are unlikely to be explained by these factors. This apparent sex-dependent difference in the response to EPA and DHA supplementation and consequences in terms of cardiovascular risk in men and women needs further investigation.

Albert et al. [7] have shown that men in the top quartile of the whole blood EPA + DPA + DHA distribution in the Physicians’ Health Study (corresponding to an average O3I of 6.9%) [6] had a 90% lower risk of sudden cardiac death than men in the lowest quartile (corresponding to an average O3I of 3.8%) [6]. In the present study, the O3I after DHA was 2.3% greater than after EPA supplementation. Data from epidemiological studies would therefore suggest a greater benefit of DHA compared with EPA on risk of CVD [10,28,29]. However, the greater increase in LDL-C concentrations with high dose DHA compared to EPA also needs to be factored in when assessing the impact of LnC3-PUFAs on CVD risk. We have shown that the reduction in serum TG, interleukin-18 and the total cholesterol/HDL-C ratio, as well as the increase in serum HDL-C, adiponectin and LDL-C were significantly more important after high dose supplementation with DHA than after EPA supplementation [10]. The extent to which the greater improvements in inflammation markers and in HDL-C with DHA compared with EPA counterbalances the potential risk associated with higher LDL-C is uncertain [10]. While changes in the O3I predicted concurrent changes in TG after both DHA and EPA, this was not the case for the change in LDL-C, which correlated with changes in the O3I only after DHA. Correlation analyses also revealed that the change in the O3I was a poor predictor of the response of other cardiometabolic risk factors to EPA and DHA. Very few studies thus far have assessed and compared the individual contributions of EPA and DHA to the association between the O3I and CVD risk, as well as to its overall predictive value. Additional intervention studies are therefore needed to determine potential benefits of a long term DHA vs. EPA supplementation on the O3I as it relates to cardiovascular outcomes.

Our use of a crossover study design had many advantages including attenuation of residual confounding and increase in statistical power. The large sample size of this carefully controlled randomized trial also provided power to detect small changes in RBC composition. The composition of RBCs tracked well with each supplementation phase, reflecting high compliance to treatments and high quality data. One risk of crossover design studies pertains to potential carryover effects of treatments. However, we found no evidence of carryover effects on changes in the O3I after EPA and DHA supplementation, indicating that washout plus intervention periods were of sufficient duration to eliminate the effect of preceding treatments (data not shown). The effects of EPA and DHA on the O3I were compared to 3 g of corn oil as control, which slightly decreased EPA and DPA relative content in RBC (−0.07% and −0.11% respectively, P < 0.05, not shown), but had no effect on DHA content (+0.15%, P=0.16, not shown) and on the O3I (+0.01%, P=0.66, not shown) compared with baseline. Most previous controlled studies on LnC3-PUFAs have also used low doses of vegetable oils as a control and to that extent, our study design is similar to previous ones on this topic [3].

### 5. Conclusions

Data from this randomized double-blind crossover study shows that the increase in the O3I is numerically greater after supplementation for 10 weeks with 2.7 g/d of DHA than after 2.7 g/d of EPA. This difference may simply reflect the fact that a proportion of EPA is elongated into DPA, which is not included in the O3I calculation. Data also suggested that the O3I may increase slightly more in men than in women after

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**Table 3** Gene expression after EPA and DHA supplementation, vs. control (N=44).

<table>
<thead>
<tr>
<th>Gene</th>
<th>Control</th>
<th>ΔEPA vs. control</th>
<th>P-value EPA vs. control</th>
<th>ΔDHA vs. control</th>
<th>P-value EPA vs. control</th>
<th>P-value EPA vs. DHA</th>
</tr>
</thead>
<tbody>
<tr>
<td>FADS1</td>
<td>1970.5 ± 314.0</td>
<td>+251.8 ± 171.5</td>
<td>0.0026</td>
<td>+16.5 ± 17.8</td>
<td>0.49</td>
<td>0.92</td>
</tr>
<tr>
<td>FADS2</td>
<td>1153.8 ± 314.0</td>
<td>−24.4 ± 17.8</td>
<td>0.11</td>
<td>−5.9 ± 17.4</td>
<td>0.21</td>
<td>0.76</td>
</tr>
<tr>
<td>ELOVL2</td>
<td>12.3 ± 3.1</td>
<td>+3.1 ± 1.7</td>
<td>0.0008</td>
<td>+2.4 ± 1.8</td>
<td>0.21</td>
<td>0.23</td>
</tr>
<tr>
<td>ELOVL5</td>
<td>970.15 ± 281.3</td>
<td>+491.4 ± 357.4</td>
<td>0.11</td>
<td>+423.6 ± 297.4</td>
<td>0.17</td>
<td>0.85</td>
</tr>
</tbody>
</table>

For each gene, expression is presented as no. of copies of mRNA normalized for the expression of the housekeeping gene glyceraldehyde 3-phosphate dehydrogenase. Values are expressed as means ± SEM. Bold indicates P < 0.01.

Adjustment for potential covariates (sex, age, energy intake and baseline fatty acid proportions) was considered only when these covariates were found to be significant at P < 0.05 in the mixed models.

1 P values for the EPA and DHA change vs. the control in the outcome, as determined by the LSMEANS statement and tested against the null hypothesis in the mixed models.

2 Main treatment P values for the comparison between EPA and DHA change vs. the control in the outcome, as determined by the main treatment effect in the mixed models.

3 Log-transformed data were used in these analyses due to skewness of the distributions of post-treatment values.
DHA supplementation. Additional studies are needed to better understand how such differences in the net change in OSI after supplementation with high dose DHA compared with EPA relate to the risk of cardiovascular outcomes.

Clinical trial registry number and website

Clinical Trial Registry number and website: http://www.clinicaltrials.gov (NCT01810003).

Disclosures

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Appendix A. Supporting information

Supplementary data associated with this article can be found in the online version at doi:10.1016/j.jplef.2017.03.008.

References


The authors’ responsibilities were as follows: BL, AT and PC designed research and obtained funding from CIHR. WSH, AC, CV, JM and KHJ conducted the research and performed laboratory analyses. JA performed statistical analyses and wrote the manuscript. BL had primary responsibility for final content. All authors critically revised the manuscript and contributed intellectually to its development, provided final approval of the submitted manuscript, had full access to all of the data in the study, take responsibility for the integrity of the data and the accuracy of the data in the analysis, affirm that the article is an honest, accurate, and transparent account of the study being reported and that no important aspects of the study have been omitted.

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