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Corresponding Author: Antoni Caimari

Additional Authors: Josep Maria del Bas, Maria Isabel Rodriguez-Naranjo, Caroline E Childs, Carolina Paras Chavez, Annette L West, Elizabeth A Miles, LLuís Arola, Philip C Calder

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Impairment of lysophospholipid metabolism in obesity: altered plasma profile and desensitization to the modulatory properties of n-3 PUFAs in a randomized controlled trial.<sup>1-3</sup>

*Josep M. del Bas, Antoni Caimari, Maria Isabel Rodriguez-Naranjo, Caroline E. Childs, Carolina Paras Chavez, Annette L. West, Elizabeth A. Miles, Lluis Arola, and Philip C. Calder*

<sup>1</sup>From Nutrition and Health Research Group, Technological Center for Nutrition and Health, Tecnio, Campus de Excelencia Internacional Cataluña Sur (CEICS), Reus, Spain (JMdB, AC, MIRN, LA); Human Development and Health Academic Unit, Faculty of Medicine, University of Southampton, Southampton, United Kingdom (CEC, CPC, ALW, EAM, PCC); National Institute for Health Research Southampton Biomedical Research Centre, University Hospital Southampton National Health Service Foundation Trust and University of Southampton, Southampton, United Kingdom (PCC); and Nutrigenomics Research Group, Department of Biochemistry and Biotechnology, Universitat Rovira i Virgili, Tarragona, Spain (LA).

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<sup>3</sup>Address correspondence to Dr Antoni Caimari, Nutrition and Health Research Group, Technological Center for Nutrition and Health, Avinguda Universitat 1, Reus 43204, Spain. Email: [antonи.caimari@ctns.cat](mailto:antonи.caimari@ctns.cat). Telephone 0034977752965.

**Authors last names:** del Bas, Caimari, Rodriguez-Naranjo, Childs, Paras Chavez, West, Miles, Arola, Calder.

**Short running head:** Lysophospholipid metabolism in obesity.

**<sup>4</sup>Abbreviations used:** ANCOVA, analysis of covariance; BMI, body mass index; CVD , cardiovascular disease; DHA, docosahexaenoic acid ; ETA, eicosatetraenoic acid; HFM, high fat meal; HOMA, homeostasis model assessment; IR, insulin resistance; LCAT, lecithin cholesterol acyl transferase; LP, lysophospholipid; LPC, lysophosphatidylcholine; LPE, lysophosphatidylethanolamine; LPI, lysophosphatidylinositol; MetS, metabolic syndrome; MTT, thiazolyl blue tetrazolium bromide ; NAFLD, non-alcoholic fatty liver disease; NASH, non-alcoholic steatohepatitis; ORO, Oil red O; PC, principal component; PCA, principal component analysis; PLS-DA, partial least square discriminant analysis; PPAR, peroxisome proliferator activated receptor; PUFA, polyunsaturated fatty acid; ROC, receiver operating characteristics.

Clinical Trial registration number: ISRCTN96712688 ([www.controlled-trials.com](http://www.controlled-trials.com))

1    ABSTRACT

2    **Background:** Plasma lysophospholipids (LPs) have emerged as signalling molecules  
3    with important effects in inflammation, insulin resistance and fatty liver disease, each  
4    of which is closely linked to obesity. Dietary n-3 PUFAs may be able to improve  
5    these conditions.

6    **Objective:** The objective was to assess the response of the plasma LPs to obesity, n-3  
7    PUFA consumption and a high fat meal challenge in order to better understand the  
8    role of LP metabolism in the progression of obesity related disorders.

9    **Study design:** We determined the concentrations of 8 lysophosphatidylcholines  
10   (LPCs), 11 lysophosphatidylethanolamines and 7 lysophosphatidylinositols in plasma  
11   of 34 normoweight and 38 obese subjects randomized to corn oil (as control) or n-3  
12   PUFA-rich fish oil (3 g/day, n=15-19 per group) for 90 days. Blood samples were  
13   collected on the last day of the study under fasting conditions and 6 h after a high fat  
14   meal (1135 kcal, 86 g fat) challenge. The profile of secreted LPs was studied in  
15   HepG2 cells under palmitate induced steatosis.

16   **Results:** Obese and normoweight subjects had different profiles of plasma LPs.  
17   Multivariate combination of the 26 LPs could discriminate between normoweight and  
18   obese subjects with an accuracy of 98%. The high fat meal challenge altered the  
19   concentration of plasma LPCs in an oil treatment-dependent manner in normoweight,  
20   but not in obese, subjects, suggesting that obesity impairs the sensitivity of LP  
21   metabolism to n-3 PUFAs. Non-cytotoxic steatosis in HepG2 cells affected the  
22   secretion pattern of LPs partially resembling the changes observed in plasma of obese  
23   subjects.

24     **Conclusions:** Obesity has a significant impact on LP metabolism, altering plasma LP  
25     profile and abolishing its sensitivity to dietary n-3 PUFAs. These effects could  
26     contribute to the onset or progression of alterations associated with obesity such as  
27     inflammation, insulin resistance and fatty liver disease. This trial was registered at  
28     [www.controlled-trials.com](http://www.controlled-trials.com) as ISRCTN96712688.

29

30     **Keywords:** lysophospholipid metabolism, lysophosphatidylcholine,  
31     lysophosphatidylethanolamine, obesity, insulin resistance, inflammation, fatty liver  
32     disease, polyunsaturated fatty acids, omega-3, fish oil

33

**34 INTRODUCTION**

35       Obesity is linked to chronic low-grade inflammation, insulin resistance (IR),  
36   non-alcoholic fatty liver disease (NAFLD), metabolic syndrome (MetS) and  
37   cardiovascular disease (CVD) among other conditions (1,2). Recent advances in the  
38   field of metabolomics have allowed untargeted exploration of the metabolic changes  
39   induced by obesity or obesity-related complications. Using this approach different  
40   lysophospholipid (LP) species, mainly lysophosphatidylcholines (LPCs), have been  
41   identified as being differentially changed in the plasma of subjects with obesity (3–6),  
42   non-alcoholic steatohepatitis (NASH) (7–9) or NAFLD (10), and to accompany the  
43   amelioration of different features associated to obesity and MetS (11,12). These  
44   findings suggest that obesity and obesity-related conditions are linked to altered  
45   plasma LPs. LPs act as signalling molecules, modulating processes such as  
46   inflammation, insulin production and insulin sensitivity through their interaction with  
47   G-protein coupled receptors (13). Thus, LPs may be important molecules in obesity  
48   and its related disorders. Nevertheless, the effects of obesity on LP metabolism and  
49   the changes that the obese phenotype can induce in the plasma LP profile remain  
50   unexplored.

51       There is increasing evidence that increased intake of n-3 polyunsaturated fatty  
52   acids (n-3 PUFAs) can partly ameliorate some obesity-associated conditions. Thus,  
53   different meta-analyses show that n-3 PUFAs have beneficial effects on obesity and  
54   insulin resistance (14–17) and exert anti-inflammatory actions at the local and  
55   systemic level (14,18,19). These effects are proposed to take place through different  
56   mechanisms such as modulation of the activity of the peroxisome proliferator  
57   activated receptor (PPAR) family of nuclear receptors or of the production of  
58   eicosanoids and other lipid mediators (16). Little is known regarding the actions of

59 n-3 PUFAs on the metabolism of LPs, although it has been shown that LPC  
60 containing n-3 polyunsaturated acyl chains (i.e. docosahexaenoic acid (DHA) and  
61 eicosapentaenoic acid (EPA)) exert anti-inflammatory actions *in vitro* and in mice by  
62 affecting prostaglandin formation (20,21). Therefore, incorporation into  
63 glycerophospholipids represents a mechanism by which n-3 PUFAs can interplay  
64 with LP metabolism.

65 Because fatty acid handling and hepatic lipid metabolism are dysregulated in  
66 obesity, we hypothesized that LP metabolism, which is highly interconnected with  
67 these processes, is also altered, resulting in measurable changes in the concentrations  
68 of plasma LP species that have been related with the progression of alterations linked  
69 to obesity. Therefore, our objective was to assess the effects of obesity on LP  
70 metabolism and how different factors that either ameliorate this condition, such as n-3  
71 PUFA consumption, or induce metabolic stress, such as a high fat meal (HFM)  
72 challenge, can modulate the plasma LP profile. Due to the emerging role of LPs as  
73 metabolic signals, our aim is to provide new evidence to support the role of LP  
74 metabolism as a key factor in the onset and progression of obesity-related diseases.

75     **SUBJECTS, MATERIALS AND METHODS**

76     **Subjects and intervention**

77                 All procedures involving human subjects were approved by the NRES South  
78                 Central–Berkshire Research Ethics Committee (submission number 11/SC/0384).  
79                 Normoweight ( $n = 50$ ) and obese ( $n = 50$ ) male and female subjects were recruited  
80                 from February 2012 to October 2013 at the University of Southampton. The primary  
81                 aim of the study was to assess the effect of n-3 PUFA consumption on blood  
82                 inflammatory markers in normoweight and obese subjects in the fasting state and in  
83                 response to a standard HFM challenge. Secondary outcomes include the examination  
84                 of the effects of obesity, n-3 PUFAs and the HFM on blood lipids and related  
85                 metabolites, which was the focus of the present study. Subjects were eligible for  
86                 enrolment into the study if they were men or women aged 18–65 y, had a BMI 18.5–  
87                  $25 \text{ kg/m}^2$  (normoweight) or BMI 30–40  $\text{kg/m}^2$  with a waist circumference > 94 cm for  
88                 men or > 80 cm for women (obese), did not eat more than one oily fish meal per  
89                 week, and if they provided written informed consent. Subjects were excluded if they  
90                 met any of the following criteria: diagnosed with diabetes; use of prescribed medicine  
91                 to control inflammation, hypertension or dyslipidaemia; use of fish oil or other oil  
92                 supplements; having chronic gastrointestinal problems; being pregnant or planning to  
93                 become pregnant within the study period; participation within another clinical trial.  
94                 The trial was registered at [www.controlled-trials.com](http://www.controlled-trials.com) as ISRCTN96712688.

95                 The trial was designed in two separated phases (**supplementary Figure 1**).

96                 The first one was a crossover study designed to determine the effects of including n-3  
97                 PUFAs or corn oil (as control) with a single HFM on post-prandial changes in  
98                 metabolites and inflammatory markers in both normoweight and obese subjects

99 (supplementary Figure 1). The HFM consisted of two croissants served with 28 g of  
100 butter and 18 g of jam, 3 x 1 g capsules containing corn oil (or n-3 PUFAs) and a  
101 milkshake made with 250 mL of semi-skimmed milk, 32 g of Nesquik™ powder and  
102 75 g of double cream. The HFM provided 85.8 g fat, 76.9 g carbohydrate, 18.6 g  
103 protein and 1134 kcal. An abdominal adipose tissue biopsy was collected during the  
104 first phase of the trial. The current work is based on the second phase of the trial,  
105 which was focused on the effects of daily intake of n-3 PUFAs or corn oil (as control)  
106 for 90 days on blood lipids and other metabolites and inflammatory markers  
107 (supplementary Figure 1). Sample size for the trial was calculated based upon  
108 anticipated findings from the second phase of the trial: a 20% reduction in plasma  
109 concentration of interleukin-6 with 90 days n-3 PUFA treatment was anticipated.  
110 Based upon mean and SD for plasma interleukin-6 concentration from previous  
111 studies it was calculated that 20 subjects per group (i.e. corn oil; n-3 PUFA) would be  
112 needed to detect a 20% difference with 80% power at the 5% significance level ( $p <$   
113 0.05). Thus, to allow for a 20% drop out rate, 50 normoweight and 50 obese subjects  
114 needed to be recruited. 50 normoweight and 50 obese subjects were initially recruited  
115 and participated in the first phase of the trial (supplementary Figure 1). Before  
116 starting this second phase, 16 normoweight and 12 obese subjects withdrew from the  
117 study; this was mainly because of the requirement to provide a second abdominal  
118 adipose tissue biopsy at the end of the second phase. The remaining cohort (34  
119 normoweight and 38 obese subjects) was randomly allocated, in a double blinded  
120 fashion, to 3 g/day corn oil as control or to 3 g/day EPAX6000 TG (EPAX, Oslo,  
121 Norway), a source of eicosapentaenoic acid (EPA; 1.1 g/day) and DHA (0.8 g/day)  
122 for 90 days (supplementary Figure 1). EPAX6000 TG is referred to as “fish oil”  
123 throughout the rest of the manuscript. Random allocation of subject study code to

124 treatment (blinded as A or B) was performed using an online random number  
125 generator. Corn oil and fish oil were provided as one g gelatine coated capsules.  
126 Capsules were provided to subjects in sealed containers. The appearance of the  
127 capsules and containers and the labelling on containers were identical for both capsule  
128 types. Subjects and all researchers were blinded to allocation until after statistical  
129 analysis was complete. After 90 days subjects attended the National Institute for  
130 Health Research Wellcome Trust Clinical Research Facility at Southampton General  
131 Hospital in the morning after an overnight fast (> 10 hours without food or drink  
132 except water). A blood sample was collected into heparin as anticoagulant and **then**  
133 **subjects consumed the same HFM described in phase one with placebo capsules.** All  
134 subjects consumed the same meal and placebo capsules. Blood was collected into  
135 heparin as anticoagulant 1, 2, 3, 4 and 6 h after finishing the meal. Plasma was  
136 prepared by centrifugation and stored at -80°C until analysis. Here data for the fasting  
137 and 6 h plasma samples are presented. There were no subject withdrawals in the  
138 second phase of the trial and so data are available for all subjects.

139 **Determination of plasma lipid, glucose and insulin concentrations**

140 Plasma triglyceride, cholesterol, HDL-cholesterol, non-esterified fatty acid  
141 (NEFA) and glucose concentrations were measured using an iLAB 600 clinical  
142 chemistry analyser and software (Instrumentation Laboratories, Warrington, UK) and  
143 enzyme-based kits provided by Wako and Instrumentation laboratories. LDL-  
144 cholesterol concentrations were estimated using the Friedwald equation. Plasma  
145 insulin concentrations were measured using a quantitative sandwich ELISA kit from  
146 Dako. Homeostasis model assessment of insulin resistance (HOMA-IR) was  
147 calculated as:

148 [Glucose in mmol/L] x [(Insulin in  $\mu$ U/L)/22.5]

149 **Sample preparation for determination of LPs**

150 For the extraction of metabolites, 100  $\mu$ L of human plasma were added to 900  
151  $\mu$ L of methanol:water (8:1 vol/vol) containing 0.5 mg/L of LPC (13:0) and 0.1 mg/L  
152 deuterated taurocholic acid-D5 as internal standard. The mixture was homogenized by  
153 ultrasonication (30 s) and vortexing (20 s). After that, samples were incubated on ice  
154 for 10 min, and then centrifuged (16000  $\times$  g, 5 min). The supernatant was dried under  
155 nitrogen flow to eliminate the solvent. Finally, it was re-dissolved in 200  $\mu$ L of  
156 methanol:water (1:1 vol/vol) to obtain a 2-fold dilution of the initial plasma  
157 concentration.

158 **Separation of LPs by reverse phase liquid chromatography**

159 LPs were separated by reverse phase (RP) liquid chromatography performed  
160 using an Agilent ZORBAX C18 SB-Aq (Agilent Technologies, Santa Clara, CA,  
161 USA) 2.1 mm  $\times$  50 mm, 1.8  $\mu$ m particle analytical column. An Agilent ZORBAX C-8  
162 (Agilent Technologies, Santa Clara, CA, USA), 2.1 mm  $\times$  30 mm, 3.5- $\mu$ m particle  
163 guard column was placed in series in front of the analytical column. An Agilent 1290  
164 Infinity HPLC system (Agilent Technologies, Santa Clara, CA, USA) with a binary  
165 pump and degasser, thermostated well plate autosampler and column compartment  
166 were used. The autosampler temperature was 4 °C, the injection volume was 2  $\mu$ L,  
167 column temperature was 60 °C, and the flow rate was 0.6 mL/min. A 2%–98% linear  
168 gradient of solvent A (0.2% acetic acid in water) to B (0.2% acetic acid in methanol  
169 (Honeywell, Morristown, NJ, USA)) was employed over 16 min followed by a  
170 solvent B hold of 2 min and a 5 min post-time for both positive and negative ion  
171 polarity analysis.

**172 Identification of LP species by mass spectrometry**

173 LP species were identified by mass spectrometry. An Agilent 6550 Accurate-  
174 Mass Quadrupole-Time of Flight (Q-TOF) mass spectrometer (MS) (Agilent  
175 Technologies, Santa Clara, CA, USA) was operated in ESI+ and ESI- modes.  
176 Dynamic mass axis calibration was achieved by continuous infusion of a reference  
177 mass solution (121.050873 and 922.009798 for positive polarity, and 119.03632 and  
178 980.016375 for negative polarity). Scanning conditions were as follows: drying gas  
179 temperature of 325 °C and flow rate of 10 L/min; vaporizer temperature of 350 °C;  
180 nebulizer pressure of 45 psi; capillary voltage 4,000 V. MS data acquired in positive  
181 mode were used to quantitative analysis of LPC species and negative mode MS data  
182 were used to quantify LPE and LPI species. Quantitative analysis was performed as  
183 described previously (22). Calibration curves were constructed using 1 to 1500 µg/L  
184 LPC (16:0), LPC (18:0), LPC (20:0), LPE (18:1), and LPI (18:1) as standards. LPC  
185 (16:0) calibration curve was used to quantify LPC (14:0), LPC (16:0) and LPC (16:1);  
186 LPC (18:0) calibration curve was used to quantify LPC (18:0), LPC (18:1), LPC  
187 (18:2), LPC (18:3) and LPC (18:4); and LPC (20:0) was used to quantify LPC (20:0),  
188 LPC (20:1), LPC (20:2), LPC (20:3), LPC (20:4), and LPC (22:5). All LPE and LPI  
189 species were quantified using the calibration curves of LPE (18:1) and LPI (18:1),  
190 respectively. The limit of detection was 0.04 µM.

**191 Cell culture experiments**

192 HepG2 cells were maintained in complete cell culture medium obtained by  
193 supplementing DMEM (Lonza Ibérica, Barcelona, Spain) with 1% L-Glutamine  
194 (Lonza Ibérica, Barcelona, Spain), 1% penicillin/streptomycin-EDTA (Sigma,  
195 Madrid, Spain), 1% non-essential amino acids (Sigma, Madrid, Spain) and 10% fetal

196 bovine serum (FBS, Sigma, Madrid, Spain). Cells were seeded either in 12 or 48 well  
197 plates at a concentration of 120,000 cells/mL in complete cell culture medium.  
198 Twelve hours after seeding, media were replaced with complete serum free media  
199 containing 1% of bovine serum albumin (BSA, Sigma, Madrid, Spain) bound to  
200 sodium palmitate (Sigma, Madrid, Spain) at 0.2, 0.5 and 0.75 mmol/L or without  
201 sodium palmitate as control for 24 h. Treatment media preparation has been described  
202 previously (23). After 24 h treatment, cell media were collected, centrifuged at 1000 g  
203 for 10 min at 4 °C to discard cell debris and stored at -80 °C until extraction and  
204 quantification of LPs. For neutral lipid staining, cells were extensively washed with  
205 PBS, fixed with 4% paraformaldehyde and subsequently incubated with oil red O  
206 (ORO; Sigma, Madrid, Spain). After microscopic analysis and micrography (Nikon  
207 Eclipse Ti-S, Izasa, Barcelona, Spain), ORO was eluted with isopropyl alcohol  
208 (Sigma, Madrid, Spain) and the absorbance read at 510 nm. For cytotoxicity assays,  
209 after 24 h of treatment, media were replaced with complete serum free media  
210 containing a 1% BSA and thiazolyl blue tetrazolium bromide (MTT; Sigma, Madrid,  
211 Spain) as previously described (23). After 4.5 h incubation, cells were washed with  
212 PBS, micrographed and intracellular MTT was extracted with DMSO (Sigma,  
213 Madrid, Spain) and quantified at 570 and 660 nm. Cell viability was quantified as  
214 Viability (%) = [A570 – A660 (palmitate)/A570-A660 (control)]×100.

215 **Statistical analyses**

216 Data were available for 15 normoweight subjects in the control group, 19  
217 obese subjects in the control group, 19 normoweight subjects in the n-3 PUFA group  
218 and 19 obese subjects in the n-3 PUFA group. Data are expressed as means ± SEM.  
219 Differences in baseline characteristics between subject groups were analysed by  
220 analysis of covariance (ANCOVA), in which treatment group assignment (G) was

221 used as a fixed factor and the variables age and sex were included as covariates. An  
222 ANCOVA model was also used to analyse the data at the end point, in both fasting  
223 and postprandial conditions. The circulating levels of LPs or their delta values  
224 (changes between the pre- and post-high fat challenge) and the delta values of  
225 anthropometric and biochemical parameters (changes from baseline) were used as  
226 dependent variables, obesity (O) and n-3 PUFA treatment (T) were included as fixed  
227 factors and the variables age and sex were included as covariates. When the  
228 interaction between obesity and treatment was statistically significant under the  
229 ANCOVA, a Bonferroni post hoc test was used to compute pairwise comparisons  
230 between groups (i.e. the effect of treatment within obesity groups and the effect of  
231 obesity within treatment groups). Grubbs' test was used to detect outliers, which were  
232 discarded for subsequent analyses. All statistical analyses were performed with SPSS  
233 Statistics 18 (SPSS, Inc., Chicago, IL), setting the level of statistical significance at  
234 bilateral 5%. Principal component analyses (PCA), partial least squares discriminant  
235 analyses (PLS-DA), multivariate biomarker validation using receiver operating  
236 characteristics (ROC) curves and hierarchical clustering analyses were performed  
237 after data normalization and autoscaling using the software Metaboanalyst 3.0 (24).

238 **RESULTS**239 **Subject characteristics**

240 Data were collected between February 2012 and October 2013 from 34  
241 normoweight subjects treated daily with 3 g of either corn oil (N-CO group, n=15) or  
242 fish oil (N-FO group, n=19) for 90 days and from 38 obese subjects who received the  
243 same treatments for the same duration (O-CO group, n=19; O-FO, n=19). At baseline,  
244 obese subjects were significantly older than the normoweight subjects and had higher  
245 body weight, BMI, waist circumference and body fat mass (**Table 1**). Obese subjects  
246 had higher plasma concentrations of total cholesterol, LDL-cholesterol, triglycerides  
247 and non-esterified fatty acids than normoweight subjects (Table 1), although the mean  
248 levels of these parameters in the obese subjects were typically within the acceptable  
249 concentration range, according to the American Association of Clinical  
250 Endocrinologist's Medical (AACE) Guidelines for management of dyslipidaemia and  
251 prevention of atherosclerosis (25). Obese subjects also had higher plasma glucose and  
252 insulin concentrations (Table 1) and the mean HOMA-IR value above 2.5 indicates  
253 insulin resistance (26).

254 There was a significant effect of treatment on fasting HDL-cholesterol and  
255 triglyceride concentrations (Table 1). N-3 PUFAs lowered triglycerides and elevated  
256 HDL-cholesterol concentrations compared with corn oil, the control (Table 1). No  
257 significant changes were found in other anthropometric and biochemical parameters  
258 analysed (Table 1).

259 **Obesity and n-3 PUFAs induce changes in plasma LP concentrations in the**  
260 **fasted state**

261           Obesity was associated with significant differences in plasma LP  
262 concentrations in the fasting state compared with those seen in normoweight  
263 individuals (**Table 2**). Obese subjects independent of treatment group, had lower  
264 concentrations of most of the identified LPCs, with the exception of LPC 20:2, which  
265 was higher in the obese subjects (Table 2). Obese subjects had lower concentrations  
266 of 5 out of 11 LPEs (LPE 14:1, LPE 18:1, LPE 18:2, LPE 20:0 and LPE 20:2) than  
267 seen in normoweight subjects (Table 2). The interaction between obesity and  
268 treatment was significant for the plasma concentration of LPE 18:0, which was lower  
269 in the obese subjects that received the n-3 PUFAs compared to the normoweight  
270 participants submitted to the same treatment (**Table 2**). Treatment with n-3 PUFAs  
271 affected the concentration of 8 out of 26 LPs. Thus, n-3 PUFA groups had lower  
272 concentrations of LPC 18:2, LPC 20:3, LPE 18:1, LPE 18:2 and LPE 20:4 and higher  
273 concentrations of LPE 20:5, LPE 22:6 and LPI 18:0.

274           PCA analysis including all the LPs revealed a clear phenotype-dependent  
275 clustering of the subjects when the scores of the 3 first principal components,  
276 explaining ~56 % of the variance, were represented (**Figure 1A**). Consistent with the  
277 ANCOVA and post-hoc analyses, when the individual scores were colored depending  
278 on both the phenotype and the oil treatment no clear treatment-dependent clusters  
279 were identified (**Figure 1B**).

280           The 26 LPs were used to set up a PLS-DA predictive model for discriminating  
281 between normoweight and obese subjects (**Figure 1C**). The quality parameters  
282 associated to the model were excellent. The degree of fit of the model to the data,  
283 represented by R<sup>2</sup> was 0.86, the quality assessment statistic (Q<sub>2</sub>), which reports the  
284 result of crossvalidation of the model was 0.78, while a threshold of > 0.4 has been  
285 proposed as an acceptable value for a biological model (27), and the prediction

286 accuracy of the crossvalidation process was 97.7%. The cross-validation revealed  
287 that the accuracy of the model was maximal (97.7%) when all 26 LPs were used,  
288 presenting an area under the ROC curve of 0.996 with a narrow confidence interval  
289 ranging from 0.982 to 1 (**Figure 1D**). These results suggest that the profile of LP  
290 abundance clearly differs between normoweight and obese subjects, and that the more  
291 LP species used to characterize the subject the better the discrimination between the  
292 two phenotypes.

293 **A high fat meal challenge magnifies the differences in plasma LPs between**  
294 **normoweight and obese subjects and modifies the long-term effects of n-3 PUFA**  
295 **treatment**

296 The abundance of the 26 LPs was determined in the same subjects 6 h after the  
297 intake of a HFM (**Table 3**). The HFM challenge accentuated the differences between  
298 normoweight and obese groups, producing the same differences in the plasma  
299 concentrations of LPCs and LPEs reported in the fasted state and revealing additional  
300 differences in the concentrations of LPEs (18:0, 20:4, 22:5 and 22:6) and LPIs (18:1  
301 and 22:6) (Table 3). These results suggest that the obesity related differences in LPC  
302 profile are robust regardless of the absorptive state of the subject while the profile of  
303 LPE and LPI species is sensitive to short-term high fat loads. **By contrast, the HFM**  
304 **challenge produced, in general, different changes in the circulating levels of LPs to**  
305 **those seen in the fasting state in the subjects chronically treated with n-3 PUFAs.**  
306 **Thus, significant interactions between obesity and treatment were found after the**  
307 **HFM intake for the plasma concentrations of LPCs (18:0, 20:1 and 20:3), which were**  
308 **significantly lower in the normoweight subjects that consumed the fish oil (N-FO**  
309 **group) compared to the normoweight participants that were treated with the corn oil**  
310 **(N-CO group) (Table 3).** In addition, although the treatment effect was maintained for

311 LPEs (18:2, 20:4 and 22:6), it was abolished for LPCs (18:2 and 20:3), LPEs (18:1  
312 and 20:5) and LPI 18:0 (Table 3).

313 The post-HFM challenge concentrations of the 26 LPs were used for  
314 unsupervised classification using the PLS-DA model obtained in the fasted state. As a  
315 result (**Figure 2A**) 67 out of 72 subjects were correctly classified (93% accuracy),  
316 obtaining an area under the ROC curve of 0.991 for the post-HFM challenge values  
317 (**Figure 2B**). Since the plasma LP profile 6 h after the HFM challenge can be used to  
318 discriminate between normoweight and obese subjects by applying a model that was  
319 obtained with the plasma LP profile in fasting conditions, these results reinforce the  
320 conclusion that the changes on plasma LP profile induced by obesity are robust and  
321 maintained even after refeeding.

322 **The response of LPC metabolism to a high fat meal challenge is modulated by n-  
323 3 PUFAAs in normoweight but not in obese subjects**

324 Different studies have shown that the obese state is associated with a  
325 decreased sensitivity to fasting and refeeding conditions in terms of regulation of  
326 genes and proteins involved in key metabolic processes (28,29). To further  
327 characterize the effects of long term n-3 PUFA intake on the postprandial response to  
328 a HFM challenge we tested whether the plasma LP profile is also sensitive to  
329 refeeding conditions. When the differences between the pre- and post-high fat  
330 challenge plasma concentrations of LPs were analysed, an interaction between the oil  
331 treatment and the obese phenotype was found for all the LPC species, revealing that  
332 n-3 PUFA treatment affected the response of LPCs to the HFM challenge only in  
333 normoweight subjects (**Figure 3** and **supplementary Table 1**). Thus, in normoweight  
334 subjects the concentrations of LPC 16:0, LPC 18:0 and LPC 20:1 decreased in

335 response to the HFM in normoweight subjects who had consumed n-3 PUFAs (N-FO  
336 group) compared to the normoweight subjects who had consumed corn oil (N-CO  
337 group). In addition, N-CO subjects showed increased circulating concentrations of  
338 LPC 18:1, LPC 18:2, LPC 20:2, LPC 20:3 and LPC 20:4 after the HFM challenge  
339 compared to the N-FO group (Figure 3 and supplementary Table 1). These treatment-  
340 dependent responses found in normoweight subjects were not observed in obese  
341 subjects (Figure 3 and supplementary Table 1). Treatment- and phenotype-dependent  
342 changes were not observed in LPE or LPI species, with the exception of LPE 20:4,  
343 which significantly increased in response to the HFM in the N-FO group but not in  
344 the O-FO group (supplementary Table 1). These results indicate an insensitivity to the  
345 effects of high fat feeding on LPC species in obese subjects.

346 **An in vitro model of steatosis partially reproduces the plasma LP profile of obese  
347 subjects**

348 Due to the key contribution of the liver to the pool of plasma LPs (30), we  
349 hypothesized that the changes in the profile of LPs associated with obesity could be,  
350 at least in part, a response of hepatocytes to the increasing load of fatty acids that  
351 occurs in obesity (1). To test this hypothesis we established a model of mild steatosis  
352 in HepG2 cells. Cells cultured in serum depleted media were exposed to one of three  
353 concentrations of palmitate. All three concentrations resulted in statistically  
354 significant intracellular neutral lipid accumulation (**Figures 4A and 4B**).

355 Nevertheless, only 0.2 mmol/L palmitate maintained full viability of the cells, while  
356 0.5 mmol/L and 0.75 mmol/L palmitate significantly decreased cell viability by 50%  
357 and 60% respectively (**Figures 4C and 4D**). The concentrations of all the LPs that  
358 could be detected (compiled in **Table 4**) were determined in the preconditioned media  
359 of HepG2 cells. PCA (80.9% of the variance explained by PC1, PC2 and PC3, **Figure**

360 **4E)** and hierarchical clustering analysis (**Figure 4F**) using all the detected LPs  
361 revealed that the three doses of palmitate resulted in very different LP profiles in the  
362 media. Thus, the cluster of cells treated with 0.2 mmol/L palmitate was clearly  
363 differentiated from the cluster of cells cultured without palmitate and from cells  
364 treated with 0.5 or 0.75 mmol/L palmitate. In turn, the clusters formed by the cells  
365 treated with 0.5 and 0.75 mmol/L palmitate overlapped and were clearly differentiated  
366 from the other two treatments. These results suggest that both steatosis and cell  
367 viability highly affect the profile of LPs produced by HepG2 cells.

368 Analysis of individual LPs (Table 4) revealed exacerbated changes for cells treated  
369 with 0.5 or 0.75 mmol/L palmitate, likely due to the release of intracellular contents  
370 into the medium upon apoptosis of the cells. Nevertheless, comparison of the cells  
371 treated with the vehicle versus cells treated with the 0.2 mmol/L palmitate revealed  
372 differences similar to those found in human plasma. Thus, 6 out of 9 LPCs were  
373 changed. Although most of the LPC species detected in HepG2 media were different  
374 from those detected in human plasma, two of the common species (LPC18:1 and  
375 LPC20:2) changed in the same direction in HepG2 cell media and human plasma. The  
376 exception to this was LPC16:0, which contains palmitic acid, the fatty acid used to  
377 establish the model. LPEs and LPIs followed a similar pattern in HepG2 media and  
378 human plasma, with 6 out of 7 common forms unchanged and only one, LPE 18:1,  
379 differing in both models. These results suggest that, despite the evident differences  
380 between the HepG2 and human models, induction of non-cytotoxic steatosis in  
381 HepG2 cells partially reproduces the changes of LP profiles found in human plasma  
382 when obese subjects are compared to normoweight subjects.

383 **DISCUSSION**

384 Our results agree with previous research showing decreased levels of different  
385 species of LPCs in obesity or insulin resistance in humans and mice (3,4,10,31) that  
386 persist even after weight loss (3). In contrast, increased levels of LPC species  
387 associated to the onset of type 1 diabetes, IR and obesity have been described as well  
388 (5,6,32–35), suggesting that the onset of these alterations might be associated to  
389 transient increases of LPC. Beside LPC species, it has been shown that plasma LPEs  
390 negatively correlate with BMI (36) and increase as obesity is ameliorated by diet (11).  
391 In agreement with these results, our analyses performed on plasma collected in fasting  
392 conditions revealed lower circulating concentrations of some LPEs in obese subjects  
393 compared to normoweight subjects. Beyond the changes found in individual LP  
394 species, it is remarkable that, with our data, the performance of the PLS-DA  
395 predictive model was maximal when all the analysed LP species were used. These  
396 results highlight that the impact of obesity on plasma LPs is not limited to a discrete  
397 number of species but to the whole profile of plasma LPs, including LPCs, LPEs or  
398 LPIs, suggesting that obesity widely affects the metabolism of LPs.

399 We have found that the metabolism of plasma LPCs is sensitive to long-term  
400 intake of n-3 PUFAs. Thus, after the HFM challenge, the concentrations of saturated  
401 LPCs were sharply decreased and those of unsaturated LPCs remained unchanged in  
402 normoweight subjects treated with n-3 PUFAs. Circulating LPCs have been related to  
403 inflammation, although whether they exert pro- or anti-inflammatory actions is still  
404 under debate. In fact the role of LPCs in inflammation might depend on their fatty  
405 acyl chain (13,37). Thus, unsaturated LPCs (20:4 and 22:6) counteract the pro-  
406 inflammatory actions of saturated (16:0) LPCs (20,21). Therefore, it could be  
407 suggested that the changes exerted by the n-3 PUFA-rich oil are positive, reflecting a

408 lower inflammatory state after the HFM. This interpretation is consistent with the  
409 anti-inflammatory properties attributed to n-3 PUFAs (14,15,38,39). In contrast, long-  
410 term intake of corn oil, rich in the n-6 PUFA linoleic acid, had no effects on the  
411 response of saturated LPCs to the HFM challenge although the concentration of LPC  
412 20:4 was increased. It is believed that the postprandial state is a period of transient  
413 acute inflammation that contributes to increasing CVD risk (40). Therefore, it could  
414 be hypothesized that, together with other mechanisms (19,38), n-3 PUFAs might  
415 attenuate postprandial inflammation by modulating the metabolism of LPs.  
416 Furthermore, in contrast to the findings in normoweight subjects, obesity impaired the  
417 sensitivity of LPC metabolism to the HFM. Recently Kardinaal *et al.* have shown that  
418 the response of plasma inflammatory markers, such as interleukin-6 and different pro-  
419 inflammatory lipids, to a high fat challenge is blunted in obese subjects with MetS  
420 compared to healthy subjects (41). These differences in the response to a high fat  
421 challenge were attributed to the lack of phenotypic flexibility that might lead to MetS.  
422 Our results support this explanation. Nevertheless, it has to be considered that most of  
423 the LPCs, including the pro-inflammatory LPC16:0, were lower in obese subjects  
424 than normoweight subjects in the fasting state. These findings might seem paradoxical  
425 at first, since obesity is associated to chronic low-grade inflammation (42). However,  
426 it is possible that the impaired sensitivity of LPC metabolism to pro-inflammatory  
427 cues in obese subjects is responsible for this finding. Other possible causes for these  
428 differences, such as age differences between groups, need further research.

429         Although our results highlight the clear effects of obesity and, to a lesser  
430 extent, n-3 PUFAs on LPs metabolism, a potential limitation of the present study is  
431 that the lack of adjustment for multiple testing might have increased the number of  
432 false positives. Therefore, results concerning individual LPs species should be

433 interpreted with caution, although these represent a starting point to assess the effects  
434 of obesity and n-3 PUFA consumption on individual LPs.

435 The changes observed in the profile of LPs could be related to other processes  
436 that are also affected by obesity. For example, it has been shown that LPCs might  
437 have an important role in insulin sensitivity. Thus, LPC12:0, 14:0 and 16:0 but not  
438 LPCs of 18 and 20 carbons induce the uptake of glucose by cultured adipocytes (43),  
439 LPC18:1 induces pancreatic insulin release (44) and circulating levels of LPC16:0 are  
440 lower in insulin resistant than in insulin sensitive subjects with NAFLD (10). In fact,  
441 lysophosphatidic acid, a product of LPC hydrolysis, has been suggested as a  
442 promising agent to treat insulin resistance (13). Therefore, the lower plasma levels of  
443 these metabolites observed in obese subjects could be related to their lower insulin  
444 sensitivity compared with normoweight subjects, as suggested by the differences  
445 found in HOMA-IR indices. This possibility is reinforced by the greater decrease of  
446 LPC16:0 observed in normoweight subjects treated with n-3 PUFAs than in the other  
447 groups after the HFM challenge, which might indicate transient adverse effects on  
448 insulin sensitivity. Indeed, a recent systematic review of meta-analyses has revealed  
449 that n-3 PUFAs have unfavourable effects on type 2 diabetes in Caucasians (14).  
450 Therefore, the interactions of obesity with LP metabolism could also contribute to or  
451 result from the onset of obesity-associated IR.

452 The differences observed in plasma LPs between normoweight and obese  
453 subjects prompted us to further explore the origin of these changes. Plasma LPs might  
454 be formed in blood by the actions of LCAT and secretory phospholipases, but a direct  
455 hepatic origin has been also demonstrated in rats and proposed as a quantitatively  
456 relevant source of LPCs (30). In fact, it has been shown that hepatic alterations such  
457 as NAFLD and NASH alter the metabolism of phospholipids and LPs in mice and

458 humans (7–10). Overall, the effects of NAFLD on circulating levels of LPCs are very  
459 similar to those described by us and others in obese subjects. In view of this evidence,  
460 it is plausible to hypothesize that these common changes have a common origin. It is  
461 well known that increasing BMI is associated with increased risk of NAFLD (45) and  
462 adipocyte cell size is associated to liver injury (46). The main underlying mechanism  
463 relates to the increased flux of fatty acids from the adipose tissue to the liver, together  
464 with altered hepatic metabolism of fatty acids and lipoproteins (1,45). Therefore, we  
465 hypothesized that accumulation of lipids in hepatocytes might alter the profile of  
466 secreted LPs. Our experiment in the *in vitro* model of steatosis obtained by  
467 challenging HepG2 with palmitate supports this hypothesis. What is more, despite the  
468 obvious differences between the *in vitro* model and human subjects, the changes  
469 induced in the profile of LPs in HepG2 media by non-cytotoxic steatosis and in  
470 plasma by obesity are remarkably similar. Therefore, it could be hypothesized that the  
471 changes induced by obesity in the LP profile are partially due to the metabolic stress  
472 that obesity induces in the liver, and place these molecules as promising biomarkers  
473 for studying alterations of liver homeostasis. More research is still needed in order to  
474 confirm this hypothesis.

475 Overall, our results suggest that obesity has a profound impact in the  
476 metabolism of LPs, modifying the profile of plasma LPs in the long term and  
477 affecting the sensitivity of these metabolites to dietary fatty acids. These effects could  
478 be mediated, at least partially, by the influence of obesity in the metabolism of  
479 hepatocytes. Due to the role of LPs as signalling molecules in processes that are  
480 usually altered in obesity, these findings provide more evidence to understand the  
481 mechanisms that favour the progression of alterations such as insulin resistance,  
482 inflammation, NAFLD or MetS. More research is needed in order to better understand

483 the exact role of plasma LPs in disease progression and, therefore, to assess whether  
484 LP metabolism represents a promising target for the prevention and treatment of  
485 obesity associated diseases as well as being a source of biomarkers for the early and  
486 non-invasive detection of metabolic alterations.

487

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492

493 **Conflict of interests**

494 Authors have no potential conflicts of interest.

495

496 **Authors' contributions**

497 JdB, AC, LA, EAM, PCC designed research; JdB, MIRN, CEC, CPC, ALW  
498 conducted research; JdB, AC, MIRN analyzed data; JdB, AC, PCC drafted the  
499 manuscript; JdB, AC, PCC had primary responsibility for final content. All authors  
500 read and approved the final manuscript.

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**Table 1. Anthropometric and biochemical characteristics of normoweight and obese subjects at baseline and changes from baseline after the treatment with 3 g/day corn oil or n-3 PUFA rich oil for 90 days.**

	Before treatment (absolute values)				After treatment ( $\Delta$ values)			
	N-FO	O-FO	N-CO	O-CO	N-FO	O-FO	N-CO	O-CO
N	19	19	15	19				
Males, N (%)	8 (42.1)	5 (26.3)	2 (13.3)	5 (26.3)				
Age (y)	30.8 $\pm$ 3.5	47.6 $\pm$ 2.7	31.9 $\pm$ 3.5	40.5 $\pm$ 2.9	O			
Weight (kg)	63.2 $\pm$ 2.1	94.9 $\pm$ 2.7	61.3 $\pm$ 2.4	101.7 $\pm$ 3.2	O	0.38 $\pm$ 0.37	0.86 $\pm$ 0.68	0.25 $\pm$ 0.46
BMI (kg/m <sup>2</sup> )	22.2 $\pm$ 0.4	34.4 $\pm$ 0.5	22.5 $\pm$ 0.5	35.5 $\pm$ 0.7	O	0.14 $\pm$ 0.13	0.34 $\pm$ 0.26	0.09 $\pm$ 0.16
Waist (cm)	75.6 $\pm$ 1.2	106.8 $\pm$ 2.5	75.3 $\pm$ 1.7	108.9 $\pm$ 2.9	O	0.04 $\pm$ 0.70	-0.01 $\pm$ 1.35	-0.92 $\pm$ 0.44
Fat mass (kg)	12.4 $\pm$ 1.1	38.0 $\pm$ 1.4	14.4 $\pm$ 1.1	41.6 $\pm$ 1.7	O	0.20 $\pm$ 0.30	0.47 $\pm$ 0.38	0.14 $\pm$ 0.35
TC (mmol/L)	4.38 $\pm$ 0.21	5.56 $\pm$ 0.27	4.37 $\pm$ 0.22	4.87 $\pm$ 0.14	O	0.01 $\pm$ 0.11	-0.07 $\pm$ 0.15	0.03 $\pm$ 0.11
HDL-C (mmol/L)	1.50 $\pm$ 0.08	1.51 $\pm$ 0.07	1.58 $\pm$ 0.08	1.41 $\pm$ 0.07		0.12 $\pm$ 0.06	0.05 $\pm$ 0.05	-0.06 $\pm$ 0.04
LDL-C (mmol/L)	2.64 $\pm$ 0.18	3.69 $\pm$ 0.22	2.63 $\pm$ 0.21	3.23 $\pm$ 0.12	O	-0.09 $\pm$ 0.09	-0.15 $\pm$ 0.12	0.09 $\pm$ 0.09
TG (mmol/L)	0.85 $\pm$ 0.08	1.43 $\pm$ 0.19	0.79 $\pm$ 0.06	1.15 $\pm$ 0.12	O	-0.16 $\pm$ 0.08	-0.10 $\pm$ 0.08	-0.03 $\pm$ 0.04
NEFAs ( $\mu$ mol/L)	423 $\pm$ 45	630 $\pm$ 37	486 $\pm$ 56	575 $\pm$ 46	O	46.4 $\pm$ 57.2	-95.3 $\pm$ 50.5	37.6 $\pm$ 67.1
								-26.9 $\pm$ 32.3

Glucose (mmol/L)	$4.79 \pm 0.09$	$5.49 \pm 0.18$	$4.73 \pm 0.07$	$5.48 \pm 0.07$	<i>O</i>	$0.00 \pm 0.10$	$0.16 \pm 0.16$	$0.14 \pm 0.11$	$-0.04 \pm 0.10$
Insulin ( $\mu$ IU/mL)	$5.19 \pm 0.61$	$12.7 \pm 1.6$	$6.00 \pm 0.56$	$16.0 \pm 1.9$	<i>O</i>	$-0.63 \pm 0.80$	$0.68 \pm 1.21$	$0.44 \pm 0.59$	$-0.41 \pm 1.26$
HOMA-IR	$1.11 \pm 0.12$	$3.33 \pm 0.48$	$1.28 \pm 0.12$	$3.74 \pm 0.47$	<i>O</i>	$-0.12 \pm 0.18$	$0.54 \pm 0.32$	$0.12 \pm 0.14$	$-0.01 \pm 0.31$

Blood was collected after overnight fasting at study entry and on the last day of the 90 day intervention. The data are given as the mean  $\pm$  SEM (n = 15-19). Baseline and post-treatment delta values (changes from baseline) are shown. Differences in anthropometric and biochemical characteristics (dependent variables) among groups were evaluated by ANCOVA, in which the variables age and sex were included as covariates. At baseline, treatment group assignment (G) was used as a fixed factor, whereas at the endpoint obesity (O) and n-3 PUFA treatment (T) were included as fixed factors. O: significant effect of obesity, T: significant effect of treatment (ANCOVA, p<0.05). N-FO: normoweight subjects treated with fish oil; O-FO: obese subjects treated with fish oil; N-CO: normoweight subjects treated with corn oil; O-CO: obese subjects treated with corn oil. TC: total cholesterol; LDL-C: low density lipoprotein cholesterol; HDL-C: high density lipoprotein cholesterol; TG: triglycerides; NEFAs: non-esterified fatty acids; HOMA-IR: homeostatic model assessment of insulin resistance.

**Table 2. Lysophospholipid concentrations in plasma of normoweight and obese subjects treated with 3 g/day corn oil or and n-3 PUFA-rich oil for 90 days and then submitted to overnight fasting**

Metabolite ( $\mu\text{mol/L}$ )	N-FO	O-FO	N-CO	O-CO	ANCOVA
<b>LPC</b>					
LPC 16:0	266.5 $\pm$ 23.6	125.7 $\pm$ 10.2	253.3 $\pm$ 23.9	131.3 $\pm$ 16.6	<i>O</i>
LPC 18:0	84.0 $\pm$ 12.0	22.2 $\pm$ 3.1	85.0 $\pm$ 10.1	24.8 $\pm$ 3.8	<i>O</i>
LPC 18:1	45.6 $\pm$ 5.2	11.9 $\pm$ 1.3	49.7 $\pm$ 6.9	14.4 $\pm$ 1.9	<i>O</i>
LPC 18:2	3.25 $\pm$ 0.39	0.771 $\pm$ 0.09	3.73 $\pm$ 0.56	1.06 $\pm$ 0.11	<i>O, T</i>
LPC 20:1	0.409 $\pm$ 0.039	0.177 $\pm$ 0.010	0.409 $\pm$ 0.038	0.169 $\pm$ 0.015	<i>O</i>
LPC 20:2	0.297 $\pm$ 0.021	0.469 $\pm$ 0.036	0.299 $\pm$ 0.027	0.524 $\pm$ 0.038	<i>O</i>
LPC 20:3	2.15 $\pm$ 0.23	0.952 $\pm$ 0.07	2.95 $\pm$ 0.28	1.24 $\pm$ 0.15	<i>O, T</i>
LPC 20:4	6.32 $\pm$ 0.62	2.87 $\pm$ 0.26	7.01 $\pm$ 0.82	3.40 $\pm$ 0.41	<i>O</i>
<b>LPE</b>					
LPE 14:1	0.854 $\pm$ 0.028	0.791 $\pm$ 0.013	0.858 $\pm$ 0.023	0.786 $\pm$ 0.014	<i>O</i>
LPE 16:0	0.096 $\pm$ 0.004	0.093 $\pm$ 0.005	0.096 $\pm$ 0.006	0.088 $\pm$ 0.003	
LPE 18:0	0.374 $\pm$ 0.025	0.310 $\pm$ 0.021*	0.316 $\pm$ 0.014	0.322 $\pm$ 0.021	<i>O, OxT</i>
LPE 18:1	0.241 $\pm$ 0.014	0.212 $\pm$ 0.014	0.292 $\pm$ 0.029	0.256 $\pm$ 0.017	<i>O, T</i>
LPE 18:2	0.496 $\pm$ 0.038	0.394 $\pm$ 0.035	0.585 $\pm$ 0.051	0.516 $\pm$ 0.046	<i>O, T</i>
LPE 20:0	0.468 $\pm$ 0.034	0.433 $\pm$ 0.021	0.443 $\pm$ 0.026	0.423 $\pm$ 0.031	<i>O</i>
LPE 20:2	1.014 $\pm$ 0.060	0.766 $\pm$ 0.042	1.118 $\pm$ 0.117	0.853 $\pm$ 0.047	<i>O</i>
LPE 20:4	0.298 $\pm$ 0.016	0.293 $\pm$ 0.023	0.393 $\pm$ 0.028	0.356 $\pm$ 0.027	<i>T</i>
LPE 20:5	0.097 $\pm$ 0.019	0.109 $\pm$ 0.009	0.065 $\pm$ 0.005	0.086 $\pm$ 0.011	<i>T</i>
LPE 22:5	0.085 $\pm$ 0.004	0.087 $\pm$ 0.004	0.080 $\pm$ 0.005	0.090 $\pm$ 0.006	
LPE 22:6	0.410 $\pm$ 0.028	0.381 $\pm$ 0.023	0.305 $\pm$ 0.016	0.275 $\pm$ 0.017	<i>T</i>
<b>LPI</b>					
LPI 16:1	0.431 $\pm$ 0.014	0.405 $\pm$ 0.005	0.428 $\pm$ 0.013	0.403 $\pm$ 0.006	<i>O</i>
LPI 18:0	0.097 $\pm$ 0.005	0.105 $\pm$ 0.005	0.085 $\pm$ 0.005	0.094 $\pm$ 0.004	<i>T</i>
LPI 18:1	0.076 $\pm$ 0.004	0.081 $\pm$ 0.004	0.077 $\pm$ 0.003	0.078 $\pm$ 0.004	
LPI 18:2	0.091 $\pm$ 0.006	0.091 $\pm$ 0.004	0.091 $\pm$ 0.006	0.094 $\pm$ 0.005	
LPI 20:3	0.062 $\pm$ 0.002	0.064 $\pm$ 0.002	0.063 $\pm$ 0.002	0.065 $\pm$ 0.002	
LPI 20:4	0.149 $\pm$ 0.008	0.157 $\pm$ 0.006	0.150 $\pm$ 0.012	0.159 $\pm$ 0.010	
LPI 22:6	0.081 $\pm$ 0.004	0.086 $\pm$ 0.004	0.082 $\pm$ 0.004	0.086 $\pm$ 0.005	

Blood was collected after overnight fasting on the last day of the 90 day intervention.

Data are mean  $\pm$  SEM (n = 15-19). The effects of treatment, obesity and their

interaction were evaluated by ANCOVA. Lysophospholipid concentrations in plasma

were used as dependent variables, obesity (O) and n-3 PUFA treatment (T) were included as fixed factors and the variables age and sex were included as covariates. O: significant effect of obesity; T: significant effect of treatment;  $O \times T$ , significant interaction between obesity and treatment (ANCOVA,  $p < 0.05$ ). \* Effect of obesity within FO groups (Bonferroni *post hoc* comparisons,  $p < 0.05$ ). N-FO: normoweight subjects treated with fish oil; O-FO: obese subjects treated with fish oil. N-CO: normoweight subjects treated with corn oil; O-CO: obese subjects treated with corn oil.

**Table 3. Lysophospholipid concentrations in plasma of normoweight and obese subjects treated with 3 g/day of corn oil or n-3 PUFA-rich oil for 90 days and then submitted to a high-fat meal challenge**

Metabolite ( $\mu\text{mol/L}$ )	N-FO	O-FO	N-CO	O-CO	ANCOVA
<b>LPC</b>					
LPC 16:0	199.2 $\pm$ 17.2	131.9 $\pm$ 17.9	244.2 $\pm$ 15.9	113.3 $\pm$ 15.5	<i>O</i>
LPC 18:0	60.2 $\pm$ 8.1\$	25.5 $\pm$ 4.4*	88.9 $\pm$ 9.8	21.3 $\pm$ 4.3#	<i>O, T, OxT</i>
LPC 18:1	51.7 $\pm$ 5.7	19.9 $\pm$ 2.8	72.3 $\pm$ 7.8	18.0 $\pm$ 2.7	<i>O</i>
LPC 18:2	4.28 $\pm$ 0.47	1.79 $\pm$ 0.21	5.89 $\pm$ 0.54	1.76 $\pm$ 0.21	<i>O</i>
LPC 20:1	0.319 $\pm$ 0.028\$	0.186 $\pm$ 0.019*	0.411 $\pm$ 0.031	0.166 $\pm$ 0.015#	<i>O, OxT</i>
LPC 20:2	0.315 $\pm$ 0.025	0.524 $\pm$ 0.048	0.437 $\pm$ 0.036	0.550 $\pm$ 0.045	<i>O, T</i>
LPC 20:3	2.66 $\pm$ 0.28\$	1.50 $\pm$ 0.16*	4.61 $\pm$ 0.30	1.76 $\pm$ 0.20#	<i>O, T, OxT</i>
LPC 20:4	6.55 $\pm$ 0.62	4.07 $\pm$ 0.46	10.51 $\pm$ 1.14	4.19 $\pm$ 0.52	<i>O, T</i>
<b>LPE</b>					
LPE 14:1	0.854 $\pm$ 0.005	0.787 $\pm$ 0.012	0.899 $\pm$ 0.026	0.807 $\pm$ 0.009	<i>O</i>
LPE 16:0	0.103 $\pm$ 0.005	0.089 $\pm$ 0.004	0.098 $\pm$ 0.005	0.098 $\pm$ 0.004	
LPE 18:0	0.393 $\pm$ 0.030	0.342 $\pm$ 0.024	0.433 $\pm$ 0.039	0.308 $\pm$ 0.018	<i>O</i>
LPE 18:1	0.634 $\pm$ 0.057	0.421 $\pm$ 0.027	0.723 $\pm$ 0.064	0.402 $\pm$ 0.029	<i>O</i>
LPE 18:2	1.277 $\pm$ 0.147	0.802 $\pm$ 0.054	1.528 $\pm$ 0.139	0.866 $\pm$ 0.065	<i>O, T</i>
LPE 20:0	0.454 $\pm$ 0.041	0.434 $\pm$ 0.029	0.547 $\pm$ 0.053	0.413 $\pm$ 0.028	<i>O</i>
LPE 20:2	1.64 $\pm$ 0.15	1.13 $\pm$ 0.07	1.83 $\pm$ 0.15	1.12 $\pm$ 0.05	<i>O</i>
LPE 20:4	0.517 $\pm$ 0.033	0.421 $\pm$ 0.022	0.740 $\pm$ 0.069	0.467 $\pm$ 0.037	<i>O, T</i>
LPE 20:5	0.205 $\pm$ 0.037	0.135 $\pm$ 0.015	0.164 $\pm$ 0.032	0.134 $\pm$ 0.019	
LPE 22:5	0.108 $\pm$ 0.004	0.097 $\pm$ 0.005	0.119 $\pm$ 0.011	0.099 $\pm$ 0.006	<i>O</i>
LPE 22:6	0.455 $\pm$ 0.034	0.379 $\pm$ 0.019	0.408 $\pm$ 0.031	0.292 $\pm$ 0.013	<i>O, T</i>
<b>LPI</b>					
LPI 16:1	0.442 $\pm$ 0.018	0.399 $\pm$ 0.006	0.437 $\pm$ 0.013	0.410 $\pm$ 0.008	<i>O</i>
LPI 18:0	0.105 $\pm$ 0.005	0.109 $\pm$ 0.003	0.104 $\pm$ 0.005	0.103 $\pm$ 0.005	
LPI 18:1	0.133 $\pm$ 0.010	0.109 $\pm$ 0.004	0.133 $\pm$ 0.012	0.109 $\pm$ 0.005	<i>O</i>
LPI 18:2	0.142 $\pm$ 0.013	0.129 $\pm$ 0.005	0.138 $\pm$ 0.012	0.123 $\pm$ 0.008	
LPI 20:3	0.068 $\pm$ 0.003	0.068 $\pm$ 0.002	0.072 $\pm$ 0.004	0.068 $\pm$ 0.002	
LPI 20:4	0.234 $\pm$ 0.020	0.224 $\pm$ 0.010	0.243 $\pm$ 0.024	0.220 $\pm$ 0.013	
LPI 22:6	0.141 $\pm$ 0.011	0.116 $\pm$ 0.004	0.141 $\pm$ 0.012	0.116 $\pm$ 0.005	<i>O</i>

Blood was collected 6 h after the administration of a high-fat meal (1135 kcal, 86 g

fat, of which 59 g saturated fat). Data are mean  $\pm$  SEM ( $n = 15-19$ ). The effects of

treatment, obesity and their interaction were evaluated by ANCOVA.

Lysophospholipid concentrations in plasma were used as dependent variables, obesity (O) and n-3 PUFA treatment (T) were included as fixed factors and the variables age and sex were included as covariates. O: significant effect of obesity; T: significant effect of treatment;  $O \times T$ , significant interaction between obesity and treatment (ANCOVA,  $p < 0.05$ ). \* Effect of obesity within FO groups; # effect of obesity within CO groups; \$ effect of treatment within normoweight groups (Bonferroni *post hoc* comparisons,  $p < 0.05$ ). N-FO: normoweight subjects treated with fish oil; O-FO: obese subjects treated with fish oil. N-CO: normoweight subjects treated with corn oil; O-CO: obese subjects treated with corn oil.

**Table 4. Concentrations of lysophospholipids in media of HepG2 cells treated with vehicle or with different concentrations of palmitate during 24 hours to induce steatosis.**

Metabolite ( $\mu\text{mol/L}$ )	vehicle	Change in			0.75 mmol/L	ANOVA
		0.2 mmol/L	Obese humans <sup>1</sup>	0.5 mmol/L		
<i>LPC</i>						
LPC 14:0	9 $\pm$ 1 <sup>a</sup>	0.3 $\pm$ 0.2 <sup>b,*</sup>	N.D.	15 $\pm$ 2 <sup>a</sup>	26 $\pm$ 4 <sup>c</sup>	**
LPC 16:0	172 $\pm$ 7 <sup>a</sup>	173 $\pm$ 10 <sup>a</sup>	Decreased	1017 $\pm$ 127 <sup>b</sup>	1455 $\pm$ 159 <sup>c</sup>	**
LPC 16:1	1876 $\pm$ 30	1841 $\pm$ 34	N.D.	1862 $\pm$ 20	1 976 $\pm$ 48	
LPC 18:1	185 $\pm$ 8 <sup>a</sup>	146 $\pm$ 8 <sup>b,*</sup>	Decreased	245 $\pm$ 17 <sup>c</sup>	330 $\pm$ 31 <sup>d</sup>	**
LPC 18:3	246 $\pm$ 8 <sup>a</sup>	250 $\pm$ 12 <sup>a</sup>	N.D.	1220 $\pm$ 304 <sup>b</sup>	2042 $\pm$ 336 <sup>c</sup>	**
LPC 18:4	54 $\pm$ 3 <sup>a</sup>	71 $\pm$ 6 <sup>b,*</sup>	N.D.	137 $\pm$ 7 <sup>c</sup>	186 $\pm$ 16 <sup>d</sup>	**
LPC 20:0	6 $\pm$ 1 <sup>a</sup>	10 $\pm$ 2 <sup>ab,*</sup>	N.D.	21 $\pm$ 6 <sup>b</sup>	26 $\pm$ 3 <sup>b</sup>	**
LPC 20:2	130 $\pm$ 5 <sup>a</sup>	170 $\pm$ 8 <sup>b,*</sup>	Increased	210 $\pm$ 4 <sup>c</sup>	200 $\pm$ 14 <sup>c</sup>	**
LPC 22:5	306 $\pm$ 9	266 $\pm$ 11 <sup>*</sup>	N.D.	328 $\pm$ 46	269 $\pm$ 61	
<i>LPE</i>						
			Unchanged in			
LPE 18:0	11 $\pm$ 1 <sup>a</sup>	12 $\pm$ 1 <sup>a</sup>	CO group	29 $\pm$ 3 <sup>b</sup>	36 $\pm$ 3 <sup>b</sup>	**
LPE 18:1	61 $\pm$ 3 <sup>a</sup>	58 $\pm$ 6 <sup>a</sup>	Decreased	98 $\pm$ 6 <sup>b</sup>	133 $\pm$ 9 <sup>c</sup>	**
LPE 18:4	1.0 $\pm$ 0.1 <sup>a</sup>	3.2 $\pm$ 0.1 <sup>b,*</sup>	N.D.	14 $\pm$ 0.4 <sup>c</sup>	15 $\pm$ 1 <sup>c</sup>	**
LPE 20:4	9 $\pm$ 1 <sup>a</sup>	9 $\pm$ 1 <sup>a</sup>	Unchanged	22 $\pm$ 2 <sup>b</sup>	33 $\pm$ 3 <sup>c</sup>	**
<i>LPI</i>						
LPI 18:1	43 $\pm$ 2	40 $\pm$ 5	Unchanged	60 $\pm$ 3	84 $\pm$ 5	
LPI 18:2	12 $\pm$ 1	13 $\pm$ 2	Unchanged	12 $\pm$ 1	16 $\pm$ 1	
LPI 20:3	32 $\pm$ 2 <sup>a</sup>	27 $\pm$ 2 <sup>bc</sup>	Unchanged	24 $\pm$ 1 <sup>b</sup>	30 $\pm$ 1 <sup>ac</sup>	**
LPI 20:4	48 $\pm$ 3	44 $\pm$ 4	Unchanged	41 $\pm$ 1	54 $\pm$ 4	

Cells were maintained for 24 hours with vehicle (media supplemented with BSA) or increasing concentrations of BSA-bound palmitate (0.2, 0.5 and 0.75 mmol/L). Mean  $\pm$  SEM of culture medium LPCs, LPEs and LPIs of two different experiments run in duplicate.<sup>1</sup> Observed change in overweight compared with normoweight subjects as reported in table 2. N.D. denotes that the metabolite was not detected in human plasma. \*significantly different from vehicle ( $p<0.05$ ; Student's t-test). \*\*denotes  $p$

<0.05 for the one-way ANOVA. Superscript letters denote significantly different groups as identified by Fisher's LSD post-hoc test.

## Figure Legends

### Figure 1.

Multivariate analyses of 26 lysophospholipids detected in plasma of 34 normoweight and 38 obese subjects. (A) Scores plot of principal components 1, 2 and 3 after a principal component analysis. Subjects are colored as normoweight or obese. (B) Same plot shown on panel A with subjects colored as normoweight (N) or obese (O) treated with corn oil (CO) or n-3 PUFA rich oil (FO) for 90 days. (C) Scores plot of components 1, 2 and 3 after a partial least squares discriminant analysis (PLS-DA) using phenotype, i.e. normoweight and obese, as discriminant factor. (D) Receiver Operating Characteristics (ROC) curve analysis of the PLS-DA model using the 2, 3, 5, 10, 20 and 26 lysophospholipids with the highest variable importance in the projection score. The area under the ROC curve and the corresponding 95% confidence interval are reported in the inset.

### Figure 2.

Projection in the PLS-DA model shown in figure 1 of lysophospholipid profiles measured 6 hours after a high fat meal challenge in normoweight and obese subjects. (A) Classification of subjects by their lysophospholipid profile as normoweight (**open circles and squares**) or obese (**filled circles and squares**) during the cross-validation of the PLS-DA model set up (training group) and classification of subjects by their lysophospholipid profile measured 6 hours after the challenge (validation group). (B) Receiver Operating Characteristics (ROC) curves calculated during the cross-validation of the PLS-DA model (**darker line** and shadowed 95% confidence interval of 0.982 to 1) and ROC curve obtained with the lysophospholipid profile determined 6 hours after the challenge as a holdout group validation (**lighter line**).

The areas under the ROC curve for the cross-validation (CV) and the holdout groups are reported in the inset.

Figure 3.

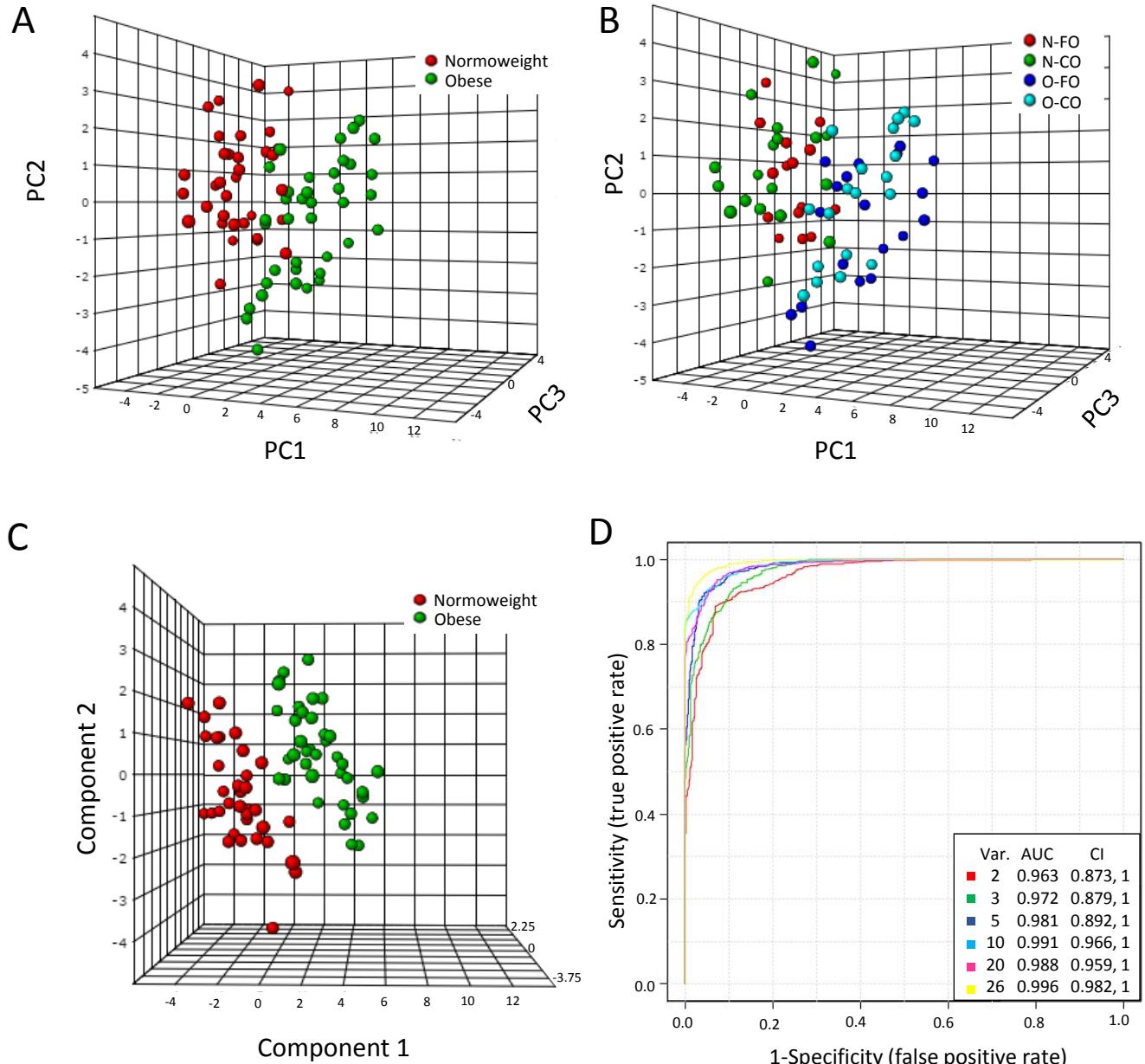
Differences in lysophosphatidylcholines concentrations between fasting conditions and 6 hours after a high fat meal challenge in normoweight (N) and obese (O) subjects treated with corn oil (CO) or n-3 PUFA rich oil (FO) for 90 days (n=15-19)..

Data are mean  $\pm$  SEM. The effects of treatment, obesity and their interaction were evaluated by ANCOVA. Delta values (changes between the pre- and post-high fat challenge) of lysophospholipid concentrations in plasma were used as dependent variables, obesity (O) and n-3 PUFA treatment (T) were included as fixed factors and the variables age and sex were included as covariates. *O*: significant effect of obesity; *T*: significant effect of treatment; *OxT*, significant interaction between obesity and treatment (ANCOVA,  $p<0.05$ ). \* Effect of obesity within FO groups; # effect of obesity within CO groups; \$ effect of treatment within normoweight groups (Bonferroni *post hoc* comparison,  $p<0.05$ ).

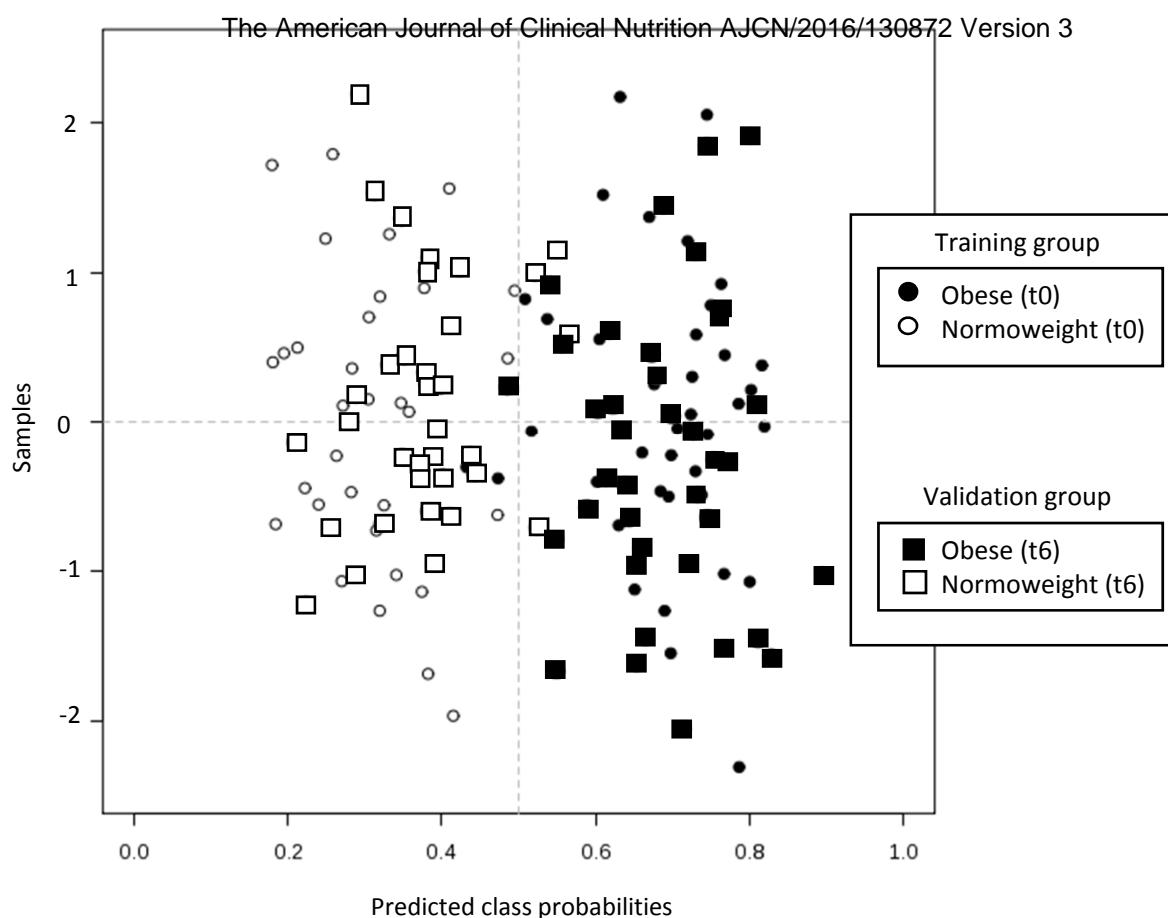
Figure 4.

Changes in the profile of secreted lysophospholipids induced by steatosis in HepG2 cells. Cells were cultured with vehicle or the indicated doses of palmitate during 24 hours. (A) Intracellular accumulation of neutral lipids was evaluated by Oil Red O (ORO) staining. (B) Spectrophotometric quantification after ORO elution from cells. (C) Cell viability was determined by incubating cells with methyl-thiazolyl-tetrazolium (MTT) after 24 hours of treatment with vehicle or palmitate at different doses. (D) Reduced MTT was eluted, quantified spectrophotometrically and cell viability was determined as % of the vehicle signal. (E) The concentrations of 17

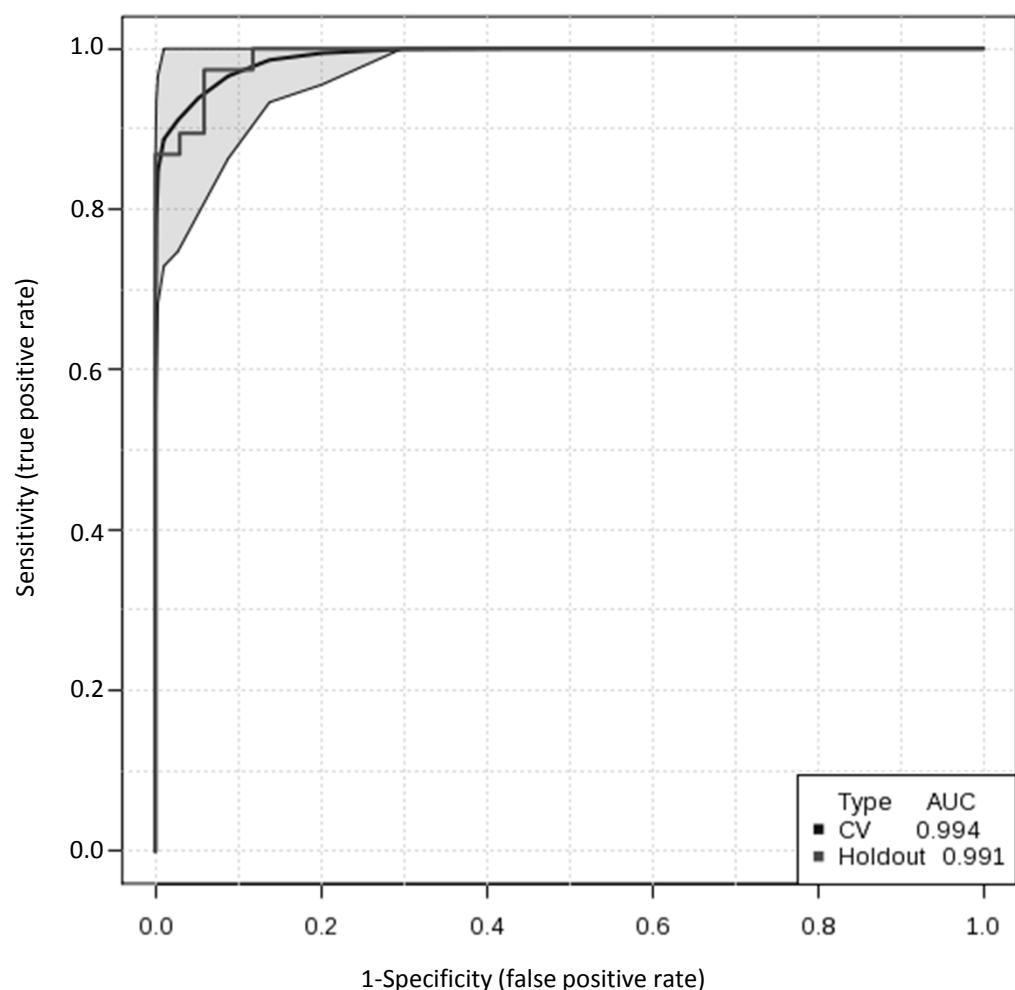
lysophospholipids detected in media of vehicle or palmitate treated HepG2 cells were subjected to a principal component analysis and scores of principal components 1,2 and 3 are represented for each replicate. (F) Clustering trends of the different experimental groups were further confirmed by hierarchical clustering analysis. Data in columns are mean  $\pm$  SEM of two independent experiments run in triplicate. \*\* p<0.01 respect the vehicle group by Student's T test.



A

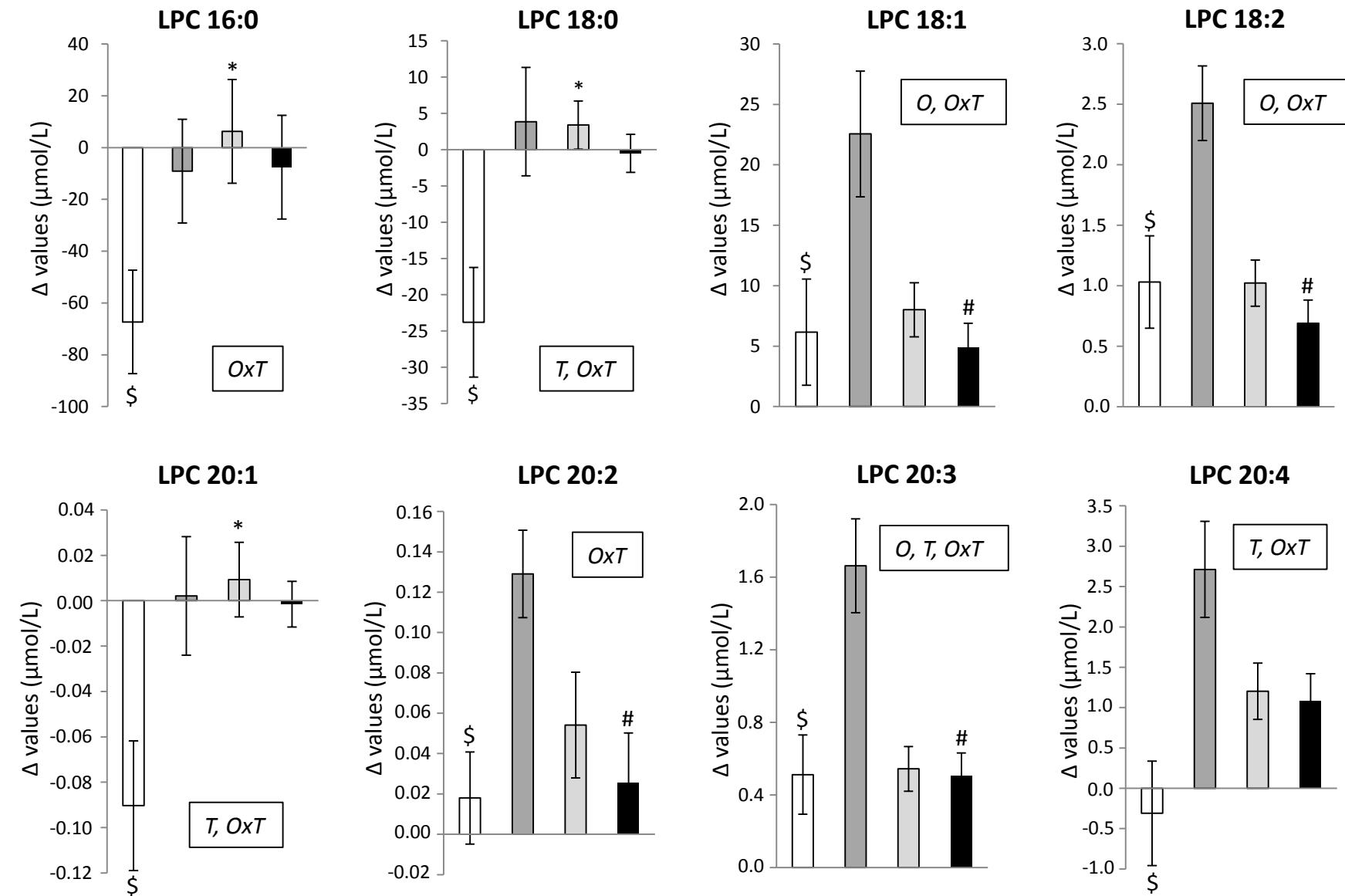


B

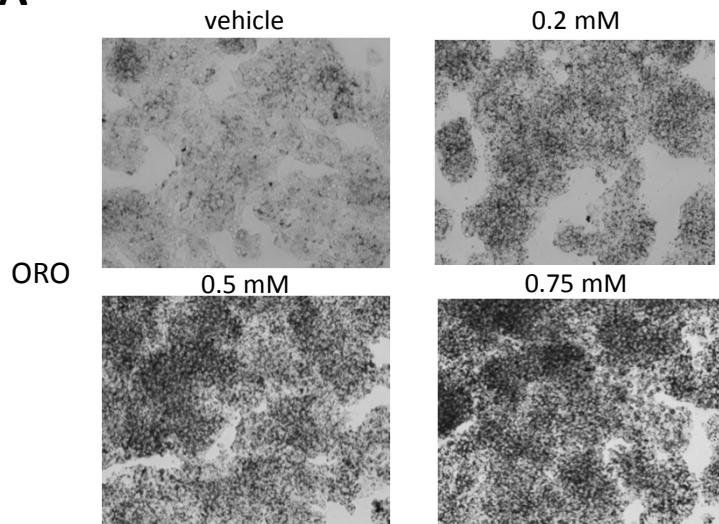


**Figure 3**

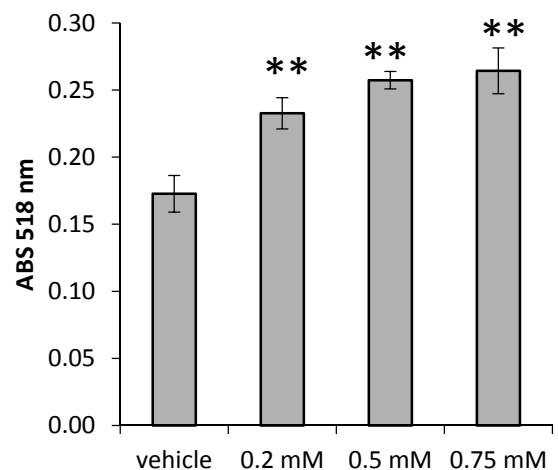
Legend:  
N-FO (white box)  
N-CO (grey box)  
O-FO (light grey box)  
O-CO (black box)



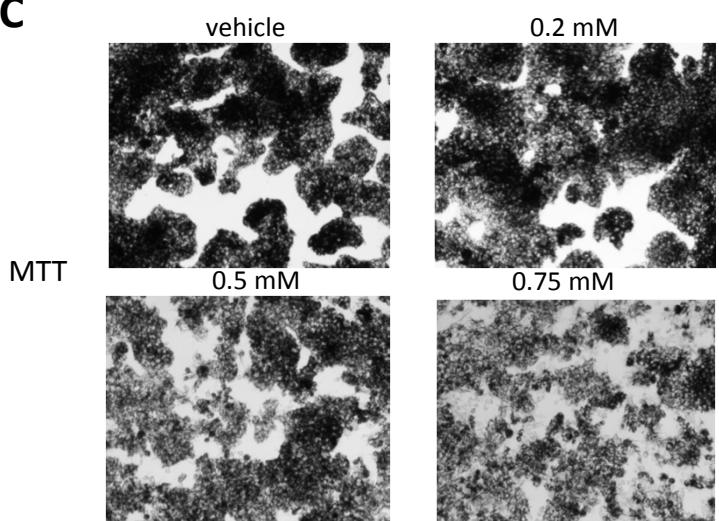
**A**



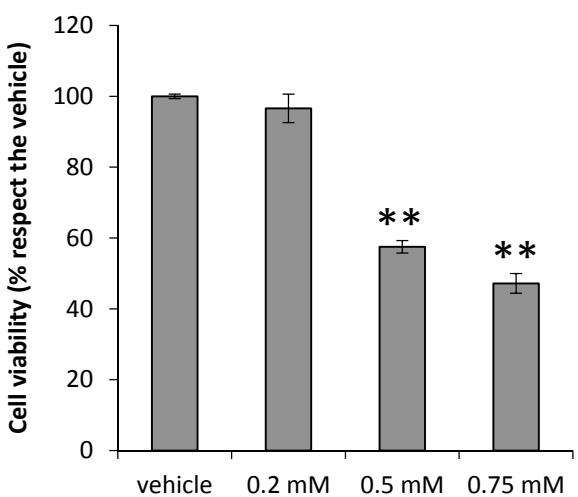
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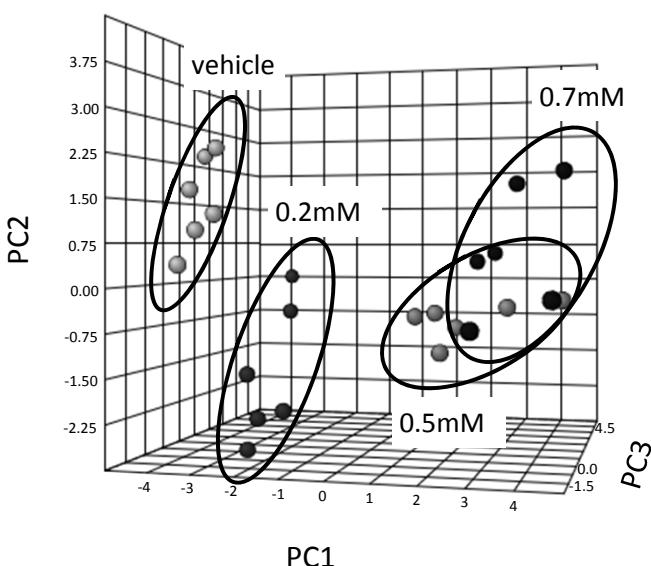
**C**



**D**



**E**



**F**

