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Change in brain plasmalogen composition by exposure to prenatal
undernutrition leads to behavioral impairment of rats

Abbreviated title: Ethanolamine plasmalogen and behavior

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44

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51

52 Abstract

53 Epidemiological studies suggest that poor nutrition during pregnancy influences
54 offspring predisposition to experience developmental and psychiatric disorders. Animal
55 studies have shown that maternal undernutrition leads to behavioral impairment, which
56 is linked to alterations in monoaminergic systems and inflammation in the brain. In this
57 study, we focused on the ethanolamine plasmalogen of the brain as a possible
58 contributor to behavioral disturbances observed in offspring exposed to maternal
59 undernutrition. Maternal food or protein restriction between gestational day (GD) 5.5
60 and GD 10.5 resulted in hyperactivity of rat male adult offspring. Genes related to the
61 phospholipid biosynthesis were found to be activated in the prefrontal cortex (PFC), but
62 not in the nucleus accumbens or striatum, in the offspring exposed to prenatal
63 undernutrition. Corresponding to these gene activations, increased ethanolamine
64 plasmalogen (18:0p-22:6) was observed in the PFC using mass spectrometry imaging. A
65 high number of crossings and the long time spent in the center area was observed in the
66 offspring exposed to prenatal undernutrition and was mimicked in adult rats via the
67 intravenous injection of ethanolamine plasmalogen (18:0p-22:6) incorporated into the
68 liposome. Additionally, plasmalogen (18:0p-22:6) increased only in the PFC, and not in
69 the nucleus accumbens or striatum. These results suggest that brain plasmalogen is one
70 of the key molecules to control behavior and its injection using liposome is a potential
71 therapeutic approach for cognitive impairment.

72

73 Keywords: Hyperactivity; Maternal undernutrition; Plasmalogen
74 phosphatidylethanolamine; Prefrontal cortex

75

76 Significance Statement

77 Maternal undernutrition correlates to developmental and psychiatric disorders. Here, we
78 found that maternal undernutrition in early pregnancy led to hyperactivity in rat male
79 offspring and induced gene activation of phospholipid-synthesizing enzyme and
80 elevation of ethanolamine plasmalogen (18:0p-22:6) level in the prefrontal cortex (PFC).
81 Intravenous injection of ethanolamine plasmalogen (18:0p-22:6) incorporated into the
82 liposome maintained crossing activity and was circumscribed to the center area for a
83 long time period, in prenatally undernourished offspring with aberrant behavior.
84 Furthermore, the amount of ethanolamine plasmalogen (18:0p-22:6) increased in the
85 PFC of the rat after injection. Our result suggests that brain plasmalogen is one of the
86 key molecules to control behavior and that its injection using liposome is a potential
87 therapeutic approach for cognitive impairment.

88 Introduction

89 Epidemiological studies have linked maternal stress during pregnancy, including
90 malnutrition, infection, daily life stress, and traumatic events, to the presence of
91 psychological and developmental disorders in offspring (Hoek et al., 1998; Khashan et
92 al., 2008; Kinney et al., 2008; Marques et al., 2015; Fineberg et al., 2016; Kundakovic
93 and Jaric, 2017). A previous study suggested that brain development is disrupted by
94 prenatal exposure to stress, which alters fetal programming by affecting the epigenome,
95 such as via changes to DNA methylation and histone modification, and induces
96 behavioral disturbances (Kundakovic and Jaric, 2017). A postmortem study of the brain
97 of a patient with schizophrenia suggested decreased *Reelin* and *GAD67* expression due
98 to the hypermethylation of their promoter regions, which was led by the upregulation of
99 DNA methyltransferase 1 (*DNMT1*) genes and may be involved in the etiology of
100 schizophrenia (Kundakovic, 2014). Meanwhile, a Dutch famine study reported the
101 relationship between prenatal undernutrition during the first trimester and the increased
102 incidence of schizophrenia (Brown and Susser, 2008). These reports suggest that
103 maternal stress may alter brain function through disturbances in the neurotransmission
104 of certain systems, such as the GABAergic system. In fact, restraint stress on mice
105 during pregnancy leads to the overexpression of DNMT1 and DNMT3a mRNA, which
106 is accompanied by the downregulation of Reelin and GAD67 protein levels, as well as
107 glutamine receptor protein from the hypermethylation of their promoter regions in the
108 frontal cortex. This cascade of events induced a schizophrenia-like phenotype
109 observable in behavioral tests performed on the male offspring after birth (Matrisciano
110 et al., 2013). Prenatal stress is thus associated with a predisposition toward
111 neurobehavioral disorders. Not only restrained stress, but prenatal caloric restriction has

112 also been shown to affect the dopamine system and neuronal excitability, resulting in a
113 decrease in anxiety-like behavior, while protein restriction results in deficits in pre-pulse
114 inhibition and locomotor activity (Markham and Koenig, 2011; Amaral et al., 2015).
115 Based on these findings, various neurotransmitter systems, including monoamine,
116 GABAergic, and glutaminergic systems, appear to be viable therapeutic targets for
117 treating behavioral disturbances induced by prenatal stress; however, membrane lipids
118 also seemed to differ between the brains of patients with psychiatric disorders and those
119 without such disorders, and little is known regarding the underlying mechanisms of this
120 process (Ghosh et al., 2017). Therefore, we hypothesized that behavioral disturbance
121 due to early prenatal undernutrition is led by aberrant brain phospholipid metabolism
122 via fetal programming. In this study, we focused on alterations in the composition of
123 brain phospholipids that are induced by prenatal undernutrition, and we have identified
124 a candidate phospholipid to control behavior.
125

126 Materials and methods

127 *Animals*

128 All animal procedures were approved by the Institutional Review Board of the Shiga
 129 University of Medical Science Animal Care and Use Committee (2011-8-1, 2014-3-7,
 130 2015-12-1, and 2019-4-2). For experiments investigating maternal undernutrition,
 131 9-week-old male (body weight [BW], 250–280 g) and 8-week-old female (BW, 160–
 132 190 g) Wistar rats were obtained from CLEA Japan, Inc. (Tokyo, Japan). Six-week-old
 133 male rats were obtained for experiments involving phosphatidylethanolamine (PE)
 134 injection. All rats were housed under a 12-h light:dark cycle (lights were turned on at
 135 08:00) and were allowed to acclimate for greater than 1 week.

136 *Diet*

137 Female rats were acclimated to a standard diet for pregnant rats (AIN-93G), containing
 138 20% casein for 2 days prior to mating, for which each female was housed with one male
 139 overnight. We defined gestational day (GD)0 as the day when a vaginal plug was
 140 observed. Pregnant rats were randomly assigned to the *ad libitum* (AL) group, the
 141 food-restriction (F) group, or the isocaloric, low-protein-diet (LPD) group, and
 142 subjected to undernutrition from GD5.5 to GD10.5 or from the day of blastocyst
 143 implantation to the day just before the closure of the neural tube (Fig. 1A) (Erb, 2006).
 144 In humans, this period is comparable to the days from E6.5 to approximately E30,
 145 which is nearing the first half of the first trimester (Bystron et al., 2008; Schoenwolf et
 146 al., 2015). Neural stem cells, and not neurons, exist in the telencephalon because this
 147 period is prior to neurogenesis (Götz and Huttner, 2005; Bystron et al., 2008). The F
 148 group was fed 50% (50F, four dams) or 40% (40F, six dams) of the daily food intake of

the AL group (eight dams). The rationale is that daily rations fell to no more than 800 kcal during the Dutch famine that occurred between December 1944 and April 1945, representing 40% of rations (>2000 calories) after June 1945 (Roseboom et al., 2001). The LPD group (five dams) was fed a diet containing 9% casein. After delivery, pups were culled to produce litters of eight offspring (four males and four females) per a dam on postnatal day (P)4. During lactation, the dams were fed CE-2, a standard pellet chow for rearing and breeding. Subsequently, the offspring were weaned on P28, and afterward, they were fed CE-2 *ad libitum*. Male offspring were used in this study because male humans show a higher risk of neuropsychiatric or neurobehavioral disorders (e.g., schizophrenia, attention deficit hyperactivity disorder [ADHD], and autism spectrum disorder [ASD]), than females (Aleman et al., 2003; Werling and Geschwind, 2013; Arnett et al., 2015). Furthermore, sex differences in ADHD and ASD may be, in part, genetically mediated (Werling and Geschwind, 2013; Arnett et al., 2015). The male offspring experienced handling once a week after weaning, and body weight of the offspring were measured at 9 and 12 weeks of age.

164 *Preparation of liposomes*

Large unilamellar liposomes composed of egg phosphatidylcholine (PC) and C18:0-22:6 plasmalogen PE (PlsEtn) or C16:0-18:1 diacyl phosphatidylethanolamine (POPE) were prepared by the extrusion method (Morita et al., 2008). Briefly, a thin film was obtained by evaporating the lipid chloroform solution and was subsequently hydrated with saline so that the concentrations of egg PC and PE (18:0p-22:6) (Avanti, AL) were 8 mg/mL and 2 mg/mL, respectively, for PE liposomes (PELs). Similarly, mixed solutions of PE (16:0-18:1) and egg PC were prepared at the same concentration as for the POPE liposomes (POPELs). To produce control liposomes (CLs), PC (10

173 mg/mL) with no PE was prepared. After five rounds of freezing and thawing, the lipid
 174 suspension was extruded through a polycarbonate filter with 100-nm pore size.

175 *Behavioral test*

176 *Effects of prenatal undernutrition on behavior*

177 From June to October, the locomotor activity of the male offspring was evaluated by the
 178 open-field test at 8 weeks of age for the AL (n = 32 from 8 dams), LPD (n = 18 from 5
 179 dams), 50F (n = 13 from 4 dams), and 40F (n = 17 from 6 dams) groups and at 12
 180 weeks for the AL (n = 18 from 5 dams), LPD (n = 19 from 5 dams), and 50F (n = 14
 181 from 5 dams) groups, to examine the impact of maternal undernutrition on behavior.
 182 Behavioral data that included device errors in tracing animals were excluded. The
 183 apparatus, measuring 90 cm in diameter and 45 cm in height, was used to monitor the
 184 behavior of rats, and the behavior was recorded for 10 min under 9 lux of light. Data
 185 were analyzed using the Limelight video tracking system (Actimetrics, IL, USA). The
 186 distance traveled and the time spent in the center was measured under the following
 187 analysis conditions: the open field was divided into the center and peripheral regions so
 188 that 1) the center region was bordered by a concentric circle passing through the
 189 midpoint of the radius of the open field (Condition 1 and 2) the area of the center region
 190 (A1) was the same as that of the peripheral region (A2) (Condition 2). The former
 191 condition was selected to allow crossing behavior to be analyzed.

192 *Effects of plasmalogen (18:0p-22:6) on behavior*

193 To examine the effect of PE (18:0p-22:6), the locomotor activity of male rats was
 194 evaluated before and after PE (18:0p-22:6) injection (for 8- and 14-week-old offspring,
 195 respectively). Rats were assigned to two different groups based on the results of
 196 crossing from the open-field test so that rats with similar locomotor activities were

197 evenly divided among the groups. A PEL or CL suspension (1 mL/kg BW) was injected
 198 into the tail vein at 14 weeks of age, and the second injection of liposome suspension
 199 was performed 2 days later. The rats in the PEL (n = 6) and CL (n = 7) groups were
 200 subjected to the open-field test or the elevated plus maze test (PEL: n = 6, CL: n = 6) 1
 201 day or 4 days after the second injection, respectively. Behavioral data of one CL rat
 202 acquired by using the elevated plus maze test was excluded because it included device
 203 errors in tracing animals. In the elevated plus maze test, rats were placed in the central
 204 square platform facing the closed arms, and their behavior was recorded for 250
 205 seconds under 8, 10, and 4 lux of light at the central square platform, facing the open
 206 arms and closed arms, respectively (Hino et al., 2019). Time spent in the open and
 207 closed arms was measured in this test. To verify the specific effect of PlsEtn
 208 (18:0p-22:6) on behavior, the alteration of locomotor activity was examined before and
 209 after POPEL, CL, or saline injection. Male rats were assigned to three different groups
 210 based on the results of the crossing analysis, and behavior was evaluated for POPEL (n
 211 = 5), CL (n = 5), and saline injection (n = 4) groups. This experiment was conducted
 212 separately from the PEL injection study. The experimental groups allocated in this study
 213 are listed in Table 2.

214 *Metabolic profiling of plasma and cerebrospinal fluid (CSF)*

215 Blood samples were collected from male offspring (AL: four dams and seven litters,
 216 40F: four dams and six litters) and CSF (AL: four dams and seven litters, 40F: four
 217 dams and five litters) at 9 weeks of age. CSF samples that got mixed with blood were
 218 excluded from the analysis. Rats were anesthetized with sodium pentobarbital solution
 219 (35 mg/kg, intraperitoneally) during the light phase (16:00–18:00). They were placed in
 220 a stereotaxic device (KOPF, CA, USA), and 50 μ L of CSF was collected immediately

221 from the cisterna magna. Then, after decapitation, 5 mL of blood was collected in test
222 tubes containing EDTA-2Na. The plasma was collected after centrifugation of the blood
223 at $2,000 \times g$ for 15 min. Samples were stored at -80°C until use.

224 Hydrophilic metabolites were extracted using the MeOH- CHCl_3 method according to
225 the procedure detailed in previous reports (Tsugawa et al., 2011; Nishiumi et al., 2012).
226 Fifty μL of plasma or CSF was mixed with 250 μL of a solvent mixture
227 (MeOH: H_2O : CHCl_3 , 2.5:1:1, v/v/v) containing 20 μL of 0.25 mg/mL 2-isopropylmalic
228 acid (Sigma-Aldrich, Tokyo, Japan) as the internal standard. The mixture was then
229 shaken at 37°C for 30 min and centrifuged at $16,000 \times g$ for 5 min at 4°C . Then, 225 μL
230 of supernatant was mixed with 200 μL of distilled water, and the solution was
231 centrifuged at $16,000 \times g$ for 5 min at 4°C . The resultant supernatant (250 μL)
232 containing hydrophilic primary metabolites was collected and lyophilized using a freeze
233 dryer. For oximation, 40 μL of 20 mg/mL methoxyamine hydrochloride (Sigma-Aldrich,
234 Tokyo, Japan) dissolved in pyridine was mixed with a lyophilized sample, and the
235 mixture was then shaken at 30°C for 90 min. For derivation, 20 μL of
236 N-methyl-N-trimethylsilyl-trifluoroacetamide (MSTFA) (GL Science, Tokyo, Japan)
237 was added, and the mixture was shaken at 37°C for 30 min. The mixture was then
238 centrifuged at $16,000 \times g$ for 5 min at 4°C , and the resultant supernatant was subjected
239 to gas chromatography–mass spectrometry (GC–MS) analysis.

240 GC–MS analysis was performed by using a GCMS-QP2010 Ultra device (Shimadzu
241 Co., Kyoto, Japan) with a fused-silica capillary column (CP-SIL 8 CB low bleed/MS;
242 30 m \times 0.25 mm inner diameter, film thickness: 0.25 μm ; Agilent Co., Palo Alto, CA,
243 USA). The front inlet temperature was set at 230°C . The flow rate of helium gas
244 through the column was 39.0 cm/s. The column temperature was held at 80°C for 2 min

245 and then raised by 15°C/min to 330°C and held for 6 min. The transfer-line and
246 ion-source temperatures were 250°C and 200°C, respectively. Twenty scans per second
247 were recorded over the mass range of 85–500 m/z by using the Advanced Scanning
248 Speed Protocol (ASSP, Shimadzu Co., Kyoto, Japan).

249 Raw data were exported in netCDF format, and peak detection and alignment were
250 performed by using MetAlign software (Wageningen UR, The Netherlands). The
251 resulting data were exported in CSV format and then analyzed with in-house analytical
252 software (AIoutput), which enabled peak identification and semi-quantification by using
253 an in-house metabolite library (Tsugawa et al., 2011; Nishiumi et al., 2012). For
254 semi-quantification, the peak height of a particular ion for each metabolite was
255 normalized to the peak height of the specified ion of 2-isopropylmalic acid (the internal
256 standard).

257 *Brain sections*

258 According to previous reports, lesions in the medial PFC cause alterations in the
259 locomotor activity of rats (Jinks and McGregor, 1997; Fritts et al., 1998). Moreover, the
260 nucleus accumbens (NAcc) and striatum (CPu) receive input from the PFC and are
261 associated with locomotor activity and impulsivity (Moreno et al., 2013; Spencer et al.,
262 2015; Scofield et al., 2016; Zhu et al., 2016; Dahoun et al., 2017). Hence, sections of
263 PFC, NAcc, and CPu were subjected to gene expression and phospholipid analyses, and
264 immunohistochemistry. Male offspring of 9 weeks of age from the maternal
265 undernutrition experiment and male rats injected with PEL or CL at 14 weeks of age (at
266 10 hours after the elevated plus maze test) were anesthetized and euthanized with
267 sodium pentobarbital solution (100 mg/kg, ip). Brain samples from the AL (four dams
268 and eight litters), 40F (five dams and eight litters), PEL (n = 4), and CL (n = 4) groups

269 were immediately dissected out and frozen in dry ice. Cryosections of the brain were cut
 270 at a thickness of 10 μ m before use for gene expression and phospholipid analyses.
 271 Greater than eight sections of the prefrontal cortex (PFC) every 200 μ m, greater than six
 272 sections of NAcc (Nucleus accumbens) every 80 μ m, and greater than seven sections of
 273 CPu (Caudate putamen) every 80 μ m were placed on Platinum Pro (Matsunami, Osaka,
 274 Japan) or polyethylene naphthalate membrane slides (Leica Microsystems, Wetzlar,
 275 Germany) for immunohistochemistry or gene expression analysis, respectively. Sections
 276 at 0.8 mm and 2 mm from the frontal end of the cerebral cortex, a section at 0.32 mm
 277 from the anterior end of the NAcc, and a section at 0.48 mm from the anterior end of the
 278 CPu were placed on indium tin oxide (ITO)-coated glass slides (Bruker Daltronics,
 279 Bremen, Germany) for phospholipid analysis.

280 *Gene expression*

281 The PFC, NAcc, and CPu of male offspring in the AL and 40F groups were dissected
 282 and collected from brain sections by a laser microdissection system (LMD6000, Leica
 283 Microsystems, Wetzlar, Germany). Gene expression analysis was performed according
 284 to the protocol described in the previous report (Kimura et al., 2018). Isolated total RNA
 285 was converted to cDNA via reverse transcription (RT) and amplified using the Ovation
 286 PicoSL WTA system V2 (NuGEN Technologies, Inc., San Carlos, CA, USA). The
 287 mRNA expression levels were estimated using quantitative real-time PCR (RT-qPCR)
 288 analysis using a LightCycler 480 system (Roche Diagnostics GmbH, Mannheim,
 289 Germany) with SYBR Premix Ex Taq II polymerase (Takara Bio, Kusatsu, Japan). The
 290 RT-qPCR reaction was performed in duplicates, and comparative C_q values of the target
 291 genes (Table 1) normalized to B2m, a reference gene, were compared between the AL
 292 and 40F groups.

293 *Counting microglial cells and activated microglial cells in PFC*

294 In each rat, ten or more brain sections at approximately 4.8 to 2.6 rostral to the bregma
 295 were fixed with 4% paraformaldehyde at room temperature for 30 min, incubated with
 296 rabbit anti-Iba1 antibody (dilution, 1:500; Wako Cat# 019-19741, RRID:AB_839504)
 297 and mouse anti-CD11b (dilution, 1:300; Bio-Rad / AbD Serotec Cat# MCA275R,
 298 RRID:AB_321302), and then incubated with goat anti-rabbit IgG (H&L) conjugated
 299 with DyLight 488 (dilution, 1:500; Abcam Cat# ab96895, RRID:AB_10679405) and
 300 goat anti-mouse IgG (H&L) conjugated with DyLight 549 (dilution, 1:500; Rockland
 301 Cat# 610-142-121, RRID:AB_1057533). Sections were stained with
 302 4',6-diamidino-2-phenylindole (DAPI), and the numbers of all Iba1-expressing
 303 microglial cells and CD11b-labeled activated microglial cells were counted in the
 304 medial PFC by using a fluorescence microscope (IX83, Olympus, Tokyo, Japan). The
 305 density of those cells was then compared between offspring in the AL and 40F groups.

306 *Matrix-assisted laser desorption/ionization-imaging mass spectrometry*
 307 *(MALDI-IMS)*

308 MALDI-IMS was performed by using the PFC, NAcc, and CPu samples in both
 309 experiments as previously described (Hossen et al., 2015; Sugiyama et al., 2015).
 310 Tissues on ITO-coated glass slides were subjected to matrix application by the
 311 sublimation/deposition method, with 1 g of 9-AA sublimated at 210°C in order for the
 312 deposition thickness to reach 1.0 µm by using the iMLayer device (Shimadzu, Kyoto,
 313 Japan). Experiments were performed by using a mass microscope, a prototype of the
 314 iMScope equipped with a 355-nm Nd:YAG laser (Shimadzu, Kyoto, Japan). Negative
 315 ions from a sample area of 30 µm × 30 µm on the PFC, NAcc, and CPu samples were
 316 obtained in a mass range of m/z 400 to 1,000. Adjacent sections of a mouse brain as a

reference were laid together with rat brain sections of 40F and AL offspring, and the rats injected with PEL and CL and were used to correct for differences in peak intensity due to differences in sample preparation between slides (Fig. 2). The peaks of 52 PE, 16 phosphatidylserine (PS), 14 phosphatidylinositol (PI), 10 lysophosphatidylethanolamine (lysoPE), and 4 lysophosphatidylinositol (lysoPI) were detected (Taguchi and Ishikawa, 2010). The peak intensity of each individual phospholipid was corrected using the average intensity of the corresponding phospholipid of the mouse references. The average peak intensities of the PFC, NAcc, and CPu, respectively, were compared between the AL and 40F groups and between the PEL and CL groups.

To identify the peak assigned at m/z 774.5 as the peak of PE (18:0p-22:6), we performed MALDI tandem mass spectrometry (MS/MS) using the mass microscope described above (Sugiyama et al., 2015) and identified PE (18:0p-22:6) via collision-induced dissociation (CID) (Zemski Berry et al., 2014).

Blood cells

Blood was collected from 14-week-old male rats for the PEL and CL injection experiment. Whole blood cells were washed with saline, applied to a 12-well flexiPERM® plate (Sarstedt, Tokyo, Japan), and affixed to an ITO-coated glass slide (Matsunami Glass Industries, Osaka, Japan), of which the surface was coated with poly-L-lysine (Hossen et al., 2015) to ensure that the blood cells were confluent. Blood cells were centrifuged to attach the cells to the surface of the glass slide, fixed with 0.25% glutaraldehyde for 5 min, and rinsed three times with 150 mM ammonium acetate buffer (pH 7.5). Samples were dried and subjected to MALDI-IMS to examine the phospholipid content in the blood cells. Briefly, negative ions from a sample area of $10\ \mu\text{m} \times 10\ \mu\text{m}$ were obtained in a mass range of m/z 400 to 1,000. The ratio of the

341 peak intensity of PE (18:0p-22:6) to the total peak intensity of all lipids was compared
342 between the PEL and CL groups for the blood cells collected from the rats of these
343 groups.

344 *Statistical analysis*

345 All data are presented as means \pm standard deviations. Differences in body weight and
346 locomotor activities in the adult offspring between the 40F, 50F, LPD, or AL groups
347 were identified using one-way analysis of variance (one-way ANOVA) followed by
348 Dunnett's multiple comparison test to examine which treatment leads to the disturbance
349 of the body growth and the behavior in the offspring compared with *ad libitum* food
350 access. Analysis of covariance (ANCOVA) was applied to compare changes in
351 behaviors after liposome injection using the behavioral data at 14 weeks of age as the
352 dependent variable, data at 8 weeks of age as the covariate, and group allocation (CL
353 and PEL groups, or CL, POPEL, and saline groups) as the independent variable.
354 Metabolites in plasma and CSF, gene expression levels, the number of microglia, and
355 phospholipid levels in the brain were compared between the 40F and AL groups by
356 using the unpaired Student's *t*-test. Phospholipid levels in the blood cells were
357 compared among CL, PEL, and saline groups using one-way ANOVA followed by
358 Tukey's HSD test. Differences were considered significant when $p < 0.05$ and Cohen's
359 d was calculated to assess the effect size. ANCOVA was performed using JMP version
360 14.0 software (SAS Institute Inc., Cary, NC, USA). The other statistical analyses were
361 performed using IBM-SPSS Statistics 22.0 software (IBM-SPSS, Inc., Chicago, IL,
362 USA).

363

364 Results

365 *Body weight of male rat offspring*

366 No significant difference was observed between the body weight of male offspring of
 367 40F and that of AL, 50F, or LPD at 9 weeks of age. However, the body weight of the
 368 male offspring was significantly lower in the 50F and LPD groups ($p = 0.06$ and 0.10 ,
 369 respectively) compared with that of the male offspring in the AL group using Dunnett's
 370 test following one-way ANOVA ($p = 0.002$, $\eta_p^2 = 0.159$ for the main effect) (Table 3).
 371 The body weight of 50F and LPD male offspring became similar to that of AL offspring
 372 at 12 weeks of age (Table 3).

373 *Maternal undernutrition during early pregnancy leads to hyperactivity in rat offspring*

374 To study the effect of nutritional stress during early embryonic stages on postnatal
 375 behavior, the open-field test was performed for adult male offsprings delivered from
 376 dams that underwent food restriction from GD 5.5 to GD 10.5. In this study, the
 377 behavioral tests performed for offspring at 8 weeks of age revealed that the total
 378 distance traveled, the distance traveled in the center area, and the frequency of crossings
 379 were significantly increased for males from the 40F group ($p = 0.028$, $p = 0.036$, and p
 380 < 0.001 , respectively), the 50F group ($p < 0.001$, $p = 0.004$, and $p < 0.001$, respectively),
 381 and the LPD group ($p = 0.026$, $p = 0.030$, and $p = 0.016$, respectively) compared with
 382 the corresponding findings for males in the AL group, in which dams were fed *ad*
 383 *libitum*, using Dunnett's test following one-way ANOVA (total distance traveled: $p <$
 384 0.001 , $\eta_p^2 = 0.209$; distance traveled in the center: $p = 0.003$, $\eta_p^2 = 0.167$;
 385 frequency of crossing: $p < 0.001$, $\eta_p^2 = 0.439$ for the main effect; Fig. 1B–D). There
 386 was no significant difference in time spent in the center area among all groups in

Condition 1 of the open-field test (Fig. 1E), but the time spent in the center area was longer for both the 40F ($p = 0.015$) and 50F ($p = 0.037$) groups compared with the AL group in Condition 2, where the radius of the center area represented 70% of the open field (Dunnett's test following one-way ANOVA: $p = 0.011$, $\eta_p^2 = 0.135$ for the main effect; Fig. 1F). Maternal protein restriction during early pregnancy, in part, contributed to the hyperactivity of the offspring. Increased total distance traveled (LPD: $p < 0.001$, 50F: $p < 0.001$), the distance traveled in the center area (LPD: $p < 0.001$, 50F: $p = 0.006$), the frequency of crossings (LPD: $p = 0.001$, 50F: $p = 0.011$), and the time spent in the center area in Conditions 1 (LPD: $p = 0.020$, 50F: $p = 0.015$) and 2 (LPD: $p = 0.001$, 50F: $p = 0.019$) were observed even at 12 weeks of age for offspring from the 50F and LPD groups compared with the AL group (one-way ANOVA with Dunnett's test), although this parameter was not examined for the offspring of the 40F group (Fig. 1G–K). However, locomotor activity did not correspond to the body weight at 9 and 12 weeks of age (Fig. 1 and Table 3). The p value and effect size for the main effect in one-way ANOVA of locomotor activities at 12 weeks of age were as follows: total distance traveled: $p < 0.001$, $\eta_p^2 = 0.333$; distance traveled in the center: $p < 0.001$, $\eta_p^2 = 0.298$; frequency of crossing: $p = 0.001$, $\eta_p^2 = 0.237$; time spent in the center area: $p = 0.011$, $\eta_p^2 = 0.173$ in Condition 1 and $p = 0.001$, $\eta_p^2 = 0.246$ in Condition 2.

Glyco- and amino-metabolisms are altered in the offspring

In this study, metabolome profiling of the plasma and CSF was performed for offspring at 9 weeks of age (after behavioral tests) of the 40F group, which displayed the most severe behavioral changes among the experimental groups (Fig.3-1). The concentration

of glycerol, which is the source of diacylglycerol in phospholipids, was increased in blood plasma ($p = 0.014$, $d = 1.57$) and CSF ($p = 0.001$, $d = 1.50$) in the 40F group compared with the AL group (Fig. 3A). In plasma, 1,5-anhydro-D-glucitol ($p = 0.038$, $d = 1.61$) and 2-aminoethanol ($p = 0.014$, $d = 1.21$) were also increased in the offspring of the 40F group (Fig. 3A and Fig. 3-1). 2-aminoethanol is converted to O-phosphoethanol amine, and finally transferred to diacylglycerol or 1-O-alkyl-2-acyl-*sn*-glycerol to produce PE (Braverman and Moser, 2012; Vance, 2015).

Microglial cell activation in the PFC is not induced by prenatal undernutrition

The cell densities of Iba1-positive microglial cells and those of both Iba1- and CD11b-positive activated microglial cells in the PFC were not significantly different between the AL and 40F groups (Fig. 3B–D). Additionally, no change in the ratio of activated glial cells to total microglial cells was observed between the AL and 40F groups (Fig. 3E); therefore, microglial cell activation was not enhanced by prenatal undernutrition during the early embryonic period.

Expression of genes related to phospholipid biosynthesis is increased in the PFC of rats exposed to prenatal undernutrition

To examine the modulation of the phospholipid biosynthetic pathway in the PFC, NAcc, and CPu in offspring (Fig. 4) exposed to maternal undernutrition, gene expression of the enzymes involved in this pathway was examined. In the brain, dihydroxyacetone phosphate (DHAP) is the main precursor of phospholipids. DHAP is synthesized from glucose and serves as a precursor of diacylglycerol, which constitutes the hydrophobic tail of phospholipids such as PC, PE, and PS (Benjamins et al., 2011). Diacylglycerol may be, in part, formed from glycerol, mediated through glycerol 3-phosphate (Fig. 4) (Jenkins and Hajra, 1976). Further, DHAP serves as a precursor of the

434 ether-phospholipid, plasmalogen (Braverman and Moser, 2012). In the first step of the
 435 synthetic pathway of plasmalogen, as well as that for diacyl phospholipids, hexokinase
 436 1 (*Hkl*) is one of the key enzymes in the regulation of the carbohydrate metabolic rate,
 437 which converts glucose to glucose 6-phosphate (McKenna et al., 2011). The expression
 438 of *Hkl* was enhanced in 40F offspring ($p = 0.016$, $d = 1.68$), although
 439 phosphofructokinase (*Pfk1*), the rate-limiting enzyme in glycolysis (McKenna et al.,
 440 2011), was not different in the PFC between 40F and control offspring (Fig. 5A, 5B, and
 441 5-1). Regarding the synthesis of the hydrophobic tail of phospholipids, the gene
 442 expression of the following enzymes related to diacylglycerol and CDP-diacylglycerol
 443 synthesis were elevated in the PFC of the offspring of the 40F group:
 444 glyceronephosphate O-acyltransferase (*Gnpat*) ($p = 0.001$, $d = 2.32$), glycerol kinase (p
 445 $= 0.016$, $d = 1.60$), glycerol-3-phosphate transferase (*Gpat*) 1, 3, and 4 ($p = 0.027$, 0.032 ,
 446 and 0.030 , and $d = 1.46$, 1.50 , and 1.57 , respectively), and phosphatidate
 447 cytidyltransferase 2 ($p = 0.045$, $d = 1.25$). Regarding the synthesis of the hydrophilic
 448 head of phospholipids, the gene expression of ethanolamine kinase for PE ($p = 0.033$, d
 449 $= 1.37$), phosphate cytidyltransferase 1, choline, alpha (*Pcytl1a*) ($p = 0.013$, $d = 1.68$)
 450 and phosphate cytidyltransferase 1, choline, and beta (*Pcytl1b*) ($p = 0.010$, $d = 1.73$)
 451 for PC; and phosphatidylserine synthase 1 (*Ptdss1*) for PS ($p = 0.007$, $d = 1.86$) was
 452 elevated in the PFC of the offspring of the 40F group (Fig. 5B). In contrast,
 453 plasmalogen is synthesized from ethanolamine or choline and
 454 1-O-alkyl-2-acyl-*sn*-glycerol, which is produced from fatty alcohol and
 455 1-O-alkyl-DHAP (Braverman and Moser, 2012). In this pathway, 1-O-alkyl-DHAP is
 456 generated from DHAP by GNPAT and alkylglycerone phosphate synthase (AGPS)
 457 (Braverman and Moser, 2012). The genes, *AGPS* ($p = 0.013$, $d = 1.71$) and *GNPAT*, of

458 these enzymes were found to be activated whereas the fatty acyl-CoA reductase 1
 459 (*Far1*) gene expression, which is a potential rate-limiting enzyme (Honsho and Fujiki,
 460 2017), was not altered in the offspring of the 40F group compared with that of the
 461 offspring of the AL group (Fig. 5B and 5-1). In contrast to the PFC, *Gpat1* ($p = 0.036$, d
 462 $= 1.39$) and *Pcytlb* ($p = 0.008$, $d = 1.65$) expression was lower in the offspring of the
 463 40F group than in the offspring of the AL group in the NAcc and CPu, respectively,
 464 although the expression of the other genes, which showed altered expression in the PFC,
 465 was not changed in the NAcc and CPu (Fig. 5A, B and 5-1). Furthermore, between the
 466 offspring of the AL group and the 40F group, in the PFC, no significant difference was
 467 observed in the expression of the calcium-independent phospholipase A2 (*iPla2*) gene,
 468 which catalyzes phospholipids (Yeagle, 2016), or that of sphingomyelin synthase 1
 469 (*Sgms1*) and sphingomyelin phosphodiesterase 3 (*Smpd3*), which convert PC into
 470 sphingomyelin and ceramide (Vance, 2015), respectively (Fig. 5-1). As described above,
 471 our results indicated that the genes of enzymes related to plasmalogen, as well as diacyl
 472 phospholipids, were activated in the PFC of offspring exposed to prenatal
 473 undernourishment. Alternatively, gene expression was not affected with respect to
 474 aquaporin 9 (Aqp9; a channel permeable to glycerol and water) (Badaut and Regli,
 475 2004) or major facilitator superfamily domain containing 2A (Mfsd2a; the major
 476 transporter for docosahexaenoic acid [DHA]) (Nguyen et al., 2014). Additionally, the
 477 expression of apolipoprotein E, which is involved in lipid transport (Liao et al., 2017),
 478 was not altered by prenatal undernutrition (Fig. 5-1).

479 *Phospholipid composition is altered in the cerebrum by maternal undernutrition*

480 The peak intensity of m/z 774.5 PE in the PFC of the offspring of the 40F group was
 481 6.6-fold higher than that of the offspring of the AL group ($p = 0.012$, $d = 2.33$; Fig. 5C

and D). CID of m/z 774.5 yielded product ions at m/z 283.2, 327.2, and 464.3; therefore, m/z 774.5 was identified as PE (18:0p-22:6) (Fig. 5E). Additionally, lysoPE (20:1) in the PFC of 40F offspring was 1.7-fold that of the AL offspring ($p = 0.037$, $d = 1.36$); however, no significant difference was observed between groups for the other phospholipids examined (Fig. 5-2). In contrast, the peak intensities of PE (20:1-22:6), lysoPE (20:4), and lysoPE (22:6) in the NAcc ($p = 0.045$, 0.034 , and 0.031 , and $d = 1.30$, 1.39 , and 1.42 , respectively), and of lysoPE (20:4) and lysoPE (22:6) in the CPu ($p = 0.004$ and 0.019 , and $d = 2.07$ and 1.57 , respectively) were attenuated in the offspring of the 40F group compared with the offspring of the AL group. However, no phospholipid showed enhanced peak intensity in the NAcc and CPu of the offspring of the 40F group (Fig. 5D and 5-2). On the other hand, only PE (18:0p-22:6) in the PFC varied in amount among PlsEtn examined in this study by prenatal undernutrition (Fig. 5D, Table 4). From the perspective of the tail forms of the phospholipids, PE (18:0p-22:6) in the PFC exclusively increased in the offspring of the 40F group among DHA (22:6)-containing phospholipids, although some phospholipids containing DHA or arachidonic acid (AA, 20:4) decreased in the NAcc and CPu (Fig. 5D, Table 4).

PlsEtn affects the behavior of adult rats

To examine the effect of PE (18:0p-22:6) on rat behavior, rats were subjected to the open-field test and elevated plus maze test after intravenous injection of PEL or CL (Fig. 6A). The age-related decline in the frequency of crossing, as well as the difference in the time spent in the center area (Condition 2) was reduced in rats in the PEL group compared with rats in the CL group (Fig. 6B). However, no significant difference was observed in the results of the elevated plus maze test between the PEL and CL groups (Fig. 6D). At the same time, no significant difference was observed in the effect of

506 POPEL, CL, and saline on rat behavior (Fig. 6C and 6E). Four days after the second
 507 injection of PEL (Fig. 6A), the amount of PE (18:0p-22:6) in the PFC ($p = 0.019$, $d =$
 508 3.02), but not that in the NAcc or CPu, was still greater in the PEL group than in the CL
 509 group, while no significant difference was observed in the amount of the other PlsEtn
 510 between these groups, as determined by MALDI-IMS (Fig. 6F and Fig. 6-1). The
 511 increased amount of PE (18:0p-22:6) in the PFC of the PEL group was verified in
 512 another PE injection experiment (Fig. 6-2). Meanwhile, the amounts of PE (22:6-24:6)
 513 ($p = 0.019$, $d = 3.02$), PE (24:4-22:6) ($p = 0.013$, $d = 3.36$), PS (16:0-22:6) ($p < 0.001$, d
 514 $= 9.02$), PI (16:1-18:1) ($p = 0.004$, $d = 4.56$), and lysoPE (18:1) ($p < 0.001$, $d = 8.58$), all
 515 of which are acyl phospholipids, were lower in the PFC in the PEL group than in the CL
 516 group (Fig. 6F). The amounts of PE (16:0-18:1) (NAcc; $p = 0.041$, $d = 2.08$, CPu: $p =$
 517 0.016 , $d = 2.68$) and lysoPE (20:1) (NAcc; $p = 0.012$, $d = 2.91$, CPu: $p = 0.033$, $d =$
 518 2.64) were lower in the NAcc and CPu for the PEL group than in the CL-group. The
 519 amounts of PE (16:1-20:5), PI (18:0-22:4), and plasmalogen PE (18:1p-20:1) were also
 520 decreased in the NAcc of the PEL group ($p = 0.008$, 0.032 , and 0.023 , and $d = 3.18$,
 521 0.85 , and 2.51 , respectively), whereas those of lysoPE (18:1), lysoPE (22:6), and
 522 lysoPE (22:4) were decreased in the CPu of that group ($p = 0.039$, 0.033 , and 0.026 , and
 523 $d = 2.13$, 2.55 , and 2.42 , respectively; Fig. 6F). To verify that increased PE (18:0p-22:6)
 524 in the PFC was not ascribed to PE incorporated into the blood cells from liposomes
 525 inside blood vessels, PE (18:0p-22:6) in blood cells was measured by MALDI-IMS.
 526 The ratio of PE (18:0p-22:6) to total lipids for the PEL group did not differ from that for
 527 the CL group (Fig. 6G). Thus, PE (18:0p-22:6) was incorporated at least into the PFC,
 528 but not blood cells, after the injection of PE liposomes. Similar to prenatal
 529 undernutrition, PE (18:0p-22:6) injection did not increase the amount of other PE, PS,

530 and PI containing DHA or AA (Fig. 6F, Table 5). Furthermore, most phospholipids that
531 varied in amount after exposure to prenatal undernutrition were not altered by PEL
532 injection, although lyso PE (22:6) was reduced in both 40F offspring and PEL-injected
533 rats (Fig. 5D and 6F, Table 5).

534 Discussion

535 The findings of our study suggest that changes in phospholipid composition led by
536 prenatal undernutrition is associated with hyperactivity in rats, and plasmalogen PE
537 (18:0p-22:6) injection reproduces a part of hyperactive behaviors. Regarding the cell
538 membrane, plasmalogens constitute approximately 20% of total phospholipids, both in
539 the rat cerebral cortex and the human brain (Braverman and Moser, 2012). In humans,
540 ethanolamine plasmalogen constitute 57% and 84% of the glycerophosphoethanolamine
541 fraction of the gray and white matter of the frontal cortex, respectively (Braverman and
542 Moser, 2012). Neurons and myelin are rich in plasmalogens, which decrease membrane
543 fluidity, increase membrane rigidity, and allow tight packing of phospholipids in the
544 membrane (Dean and Lodhi, 2017). Plasmalogens play a role in membrane trafficking
545 and fusion processes, Schwann cell differentiation and function, molecule antioxidation,
546 and inhibition of neuronal apoptotic signaling. Hence, a deficiency of plasmalogens
547 induces impairments of neurotransmitter release from synaptosomes to the presynaptic
548 cleft, myelination and axonal sorting by Schwann cells, and neuronal apoptosis
549 signaling (Dean and Lodhi, 2017). These reports suggest the clinical importance of
550 plasmalogens to the nervous system. The amyloid β peptide, which is rich in the brains
551 of patients with Alzheimer's disease, reduces AGPS protein stability and decreases
552 plasmalogen PE levels in patients with Alzheimer's disease (Han et al., 2001; Grimm et
553 al., 2011). Recently, Hossain et al. reported that inflammatory stimuli, such as the
554 administration of lipopolysaccharide (LPS) or polyriboinosinic:polyribocytidylic acid,
555 reduce plasmalogens in murine glial cells through the activation of NF- κ B, which
556 downregulates *Gnpat* through increased c-Myc recruitment to the *Gnpat* promoter
557 (Hossain et al., 2017). Similar findings have been observed for the murine brain after

aging, exposure to chronic restraint stress, and injection of LPS; furthermore, the reduction of plasmalogen induced activation of microglial cells and elevated expression of proinflammatory cytokines (Hossain et al., 2017). In brains from transgenic mice model of Alzheimer's disease, and postmortem brain tissues from patients with Alzheimer's disease, *Gnpat* reduction via a similar mechanism has been observed (Hossain et al., 2017). Likewise, maternal infection, obesity, a high-fat diet, and restraint stress with bright-light exposure causes microglial cell activation and proinflammatory cytokine induction in the fetal and postnatal brain of rodents and monkeys, and maternal stress results in anxiety-like, depressive and aggressive behavior, and schizophrenia-like behavior in offspring (Bilbo and Tsang, 2010; Grayson et al., 2010; Matrisciano et al., 2012; Diz-Chaves et al., 2013; Sasaki et al., 2013; Marques et al., 2015). In addition, activation of microglia is augmented in the brain including the anterior and orbitofrontal cortices in young adults with ASD, although the distribution pattern of activated microglia is similar to that of healthy control subjects, as determined by positron emission tomography (Suzuki et al., 2013). Maternal obesity before pregnancy is considered a risk factor for ADHD and ASD in humans (Andersen et al., 2017). Obesity is involved in elevated inflammatory mediators, e.g. IL-6, which induces Th17 cell differentiation. IL-17A secreted from Th17 may act to promote ASD by affecting fetal neurodevelopment (Wong and Hoeffler, 2018). These results suggest that brain inflammation plays a key role in behavior and that plasmalogen alters brain function through its anti-inflammatory effects. However, in our study, microglial cell activation was not altered by maternal undernutrition, at least for the adult offspring. Further, injection of plasmalogen PE (18:0p-22:6) to adult rats in the 40F group altered the phospholipid composition and resulted in two characteristic behaviors: frequent

582 crossing and long time spent in the center area in the open-field test.

583 Therefore, the hyperactivity of the rat offspring that were exposed to prenatal

584 undernutrition may be attributable to the phospholipid composition of the brain rather

585 than a direct effect of undernutrition on inflammatory reactions. Patients with RCDP,

586 who display plasmalogen deficiency, have psychomotor retardation, and, in severely

587 affected cases, they display microcephaly and cerebellar atrophy (Berger et al., 2016).

588 Myelination and neuronal migration are thought to be causes of these features of

589 patients with RCDP (Berger et al., 2016). RCDP type 1, type 2, and type 3 are caused by

590 mutations of *PEX7*, *GNPAT*, and *AGPS* (Berger et al., 2016), respectively, all of which

591 contribute to plasmalogen synthesis. In our study, expression of the latter two genes was

592 elevated in the PFC. In RCDP fibroblasts with the *PEX7* mutation, peroxisome targeting

593 signal 2 protein, phytanoyl-CoA hydroxylase, and AGPS fail to be imported into the

594 peroxisome (Yu et al., 2013). The *PEX7* homozygous mutation has also been found in

595 three ASD children whose unaffected siblings were heterozygous or wild-type within

596 one family (Yu et al., 2013). Moreover, single-nucleotide-polymorphism fine mapping

597 has shown that *GNPAT* is a candidate gene for schizophrenia, as is *DISC1* (Liu et al.,

598 2006). These findings suggest that altered phospholipid metabolism, especially

599 plasmalogen metabolism, may be involved in a person's vulnerability to developmental

600 and psychiatric disorders. *Gnpat*-knockout mice show delayed migration of granule cell

601 precursors, enhanced apoptosis in the cerebellum, and hypo- and dysmyelination in the

602 neocortex, cerebellum, and corpus callosum (Berger et al., 2016). Aberrant myelination

603 may be one of the key factors in hyperactivity because MR findings suggest altered

604 myelination in the white matter of adults with ADHD (Wu et al., 2017). Skin fibroblasts

605 derived from patients with RCDP showed reduced PlsEtn, whereas the total amount of

606 PE was maintained by an increase in other PEs (Dorninger et al., 2015).
607 Polyunsaturated fatty acid (PUFA)-containing PlsEtn was reduced in these cells, and
608 AA-containing, but not DHA-containing, PE species mainly compensated for PlsEtn
609 deficiency (Dorninger et al., 2015). Similar findings have been observed for
610 *Gnpat*-knockout mice; therefore, the ratio among essential PUFAs, as well as the ratio
611 between PlsEtn and PE, may be critical for brain development, and a shift in these ratios
612 may be the cause of the psychomotor retardation of patients with RCDP. In contrast, in
613 our study, the levels of PE (18:0p-22:6) in the PFC and lyso PE (22:6) in CPu were both
614 altered by exposure to prenatal undernutrition and PE (18:0p-22:6) injection.
615 Additionally, injection of POPE, which is not a plasmalogen, did not alter rat behavior.
616 A specific plasmalogen, such as PE (18:0p-22:6), and a specific DHA-containing PE,
617 such as lyso PE (22:6), coupled with the level of plasmalogen in the brain may have a
618 function for behavior. Exogenous administration of plasmalogen can be considered as a
619 potential therapeutic strategy as it results in changes in the phospholipid composition of
620 the brain. Regarding psychiatric disorders, the levels of PUFAs (e.g., PE22:5n6,
621 PC20:3n6, and PC22:5n6) were lower in the white matter adjacent to the dorsolateral
622 PFC of a patient with schizophrenia, while the level of PE20:2n6 was higher, and those
623 of PE22:5n6, PC20:4n6, and PC22:5n6 were lower, in the white matter of the patient
624 with bipolar disorder, despite no alterations in the plasmalogen level for both disorders
625 (Ghosh et al., 2017). A subset combination of the head group (e.g., ethanolamine,
626 choline, serine, or inositol) and a fatty acid tail, such as DHA or AA, which are
627 incorporated into the phospholipid, may be critical to evoke the behavioral alterations
628 that are characteristic of developmental or psychiatric disorders, and thus, could be a
629 therapeutic target to improve conditions for patients with these diseases. As in one of

the trials, the administration of a PS supplement in a chewable tablet presentation improved ADHD for children aged 4–14 years (Hirayama et al., 2014). Exogenous PS can cross the blood–brain barrier (BBB) and function in the brain (Glade and Smith, 2015). Regarding the delivery of plasmalogen into the brain across the BBB, liposomes with PC may be one of the preferred carriers, since the intravenous injection of liposomes has been the preferred delivery route for many previous studies (Vieira and Gamarra, 2016). However, PE has a cationic head, and the liposome may not be able to reach the brain because of nonspecific binding to the peripheral tissues and serum proteins (Vieira and Gamarra, 2016) if the ethanolamine head is exposed on the surface of the liposome membrane. In our study, the amount of PE (18:0p-22:6) was not altered in blood cells, suggesting that little of the injected PE was taken into the peripheral tissues during circulation in the brain before liposomes were captured by the liver. Plasmalogen is asymmetrically localized to the inner leaflet of the myelin membrane bilayer (Kirschner and Ganser, 1982), and plasmalogen-rich membranes tend to form non-lamellar inverse hexagonal structures compared with the membrane, which exclusively consists of diacyl phospholipids (Dean and Lodhi, 2017). The surface of the liposomes that were used in our study may be electrically neutral, and their nonspecific binding to the peripheral tissues and serum proteins may be prevented.

In summary, maternal undernutrition during early pregnancy led to the hyperactivity of male rat offspring, and the behavioral changes observed may be, in part, caused by an alteration of the plasmalogen composition in the PFC, which was induced by the activation of the phospholipid synthetic pathway. Ethanolamine plasmalogen (18:0p-22:6) appears to play a critical role in behavior. Thus, plasmalogens could be candidate therapeutic molecules for improving behavioral disorders. Further study of

654 their complex functions is warranted.

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870

871 Figure legends

872

- 873 Fig. 1 Study design and behavior of rat offspring subjected to prenatal undernutrition.
- 874 (A) The experimental schedule is shown. The daily food intake was restricted from
- 875 GD5.5 to GD10.5 or from the day of blastocyst implantation to the day just before the
- 876 closure of the neural tube. (B–F) Behavioral tests at 8 weeks of age: male rat offspring
- 877 exhibited hyperactivity in the LPD (n = 18), 50F (n = 13), and 40F (n = 17) groups
- 878 compared with the offspring in the AL group (n = 32). (G–K) Behavioral tests at 12
- 879 weeks of age: behavioral disturbances continued at 12 weeks of age in the LPD (n = 19)
- 880 and 50F (n = 14) offspring compared with the AL offspring (n = 18). The statistical
- 881 analysis was conducted using one-way ANOVA with Dunnett's test. * $p < 0.05$, ** $p <$
- 882 0.01 , *** $p < 0.001$; A1: the area of the center region, A2: the area of the peripheral
- 883 region.

884

885 Fig. 2 Preparation of the brain sections for MALDI-IMS (A) Rat brain sections were
 886 aligned between adjacent mouse brain sections (references). (B) Peak intensity of each
 887 phospholipid was corrected using the average peak intensity of the same phospholipid
 888 of the mouse reference sections.

889

890 Fig. 3 Metabolome profiling of the plasma and CSF, and the observation of microglia in
 891 the PFC. (A) 2-aminoethanol and glycerol increased in the rat offspring exposed to
 892 prenatal undernutrition (plasma: $n = 7$ in AL and $n = 6$ in 40F; CSF: $n = 7$ in AL and $n =$
 893 5 in 40F) (See also Figure 3-1). (B) The dotted area, which was the middle third of the
 894 box area, was examined. (C) Microglial cell activation in the PFC Iba1-positive
 895 microglia (green), CD11b-positive cells (red), and activated microglia (yellow) are
 896 shown in the PFC of the offspring of the AL and 40F groups. Scale bar, 20 μ m. (D) The
 897 densities of microglia and activated microglia were not increased in the PFC of 40F ($n =$
 898 8) offspring compared with AL ($n = 8$) offspring. (E) The ratio of the number of
 899 activated microglia to the total number of microglia was not altered by prenatal
 900 undernutrition. $*p < 0.05$, $**p < 0.01$, Student's t -test.

901

902 Fig. 4 Biosynthetic pathway of plasmalogens and diacyl phospholipids. The enzymes
 903 related to phospholipid synthesis, with intermediates, are shown. The genes indicated
 904 by underlined bold italic characters were activated. Abbreviations of the
 905 genes are noted as follows: *Hk*:Hexokinase, *GPI*:Glucose-6-phosphate isomerase,
 906 *Pfk*:Phosphofructokinase, *ald*:aldolase, *ald*:aldolase, *Gpd*:Glycerol-3-phosphate dehydrogenase, *Far*:fatty acyl-CoA reductase, *Gk*:Glycerol
 907 kinase, *Gnpat*:Glyceronephosphate O-acyltransferase, *Agps*:Alkylglycerone phosphate

909 *synthase, ADHAPAR:Alkyl/acyl-glycerophosphate acyltransferase, Plpp:Phosphatidic*
 910 *acid phosphatase, Cept:Choline/ethanolamine phosphotransferase,*
 911 *Plasmenylethanolamine desaturase, Ept:Ethanolamine phosphotransferase,*
 912 *Chpt:Choline phosphotransferase, Gpat:Glycerol-3-phosphate acyltransferase, Lpcat:*
 913 *LysoPA-acyltransferase, Chk:choline kinase, Etnk:Ethanolamine kinase,*
 914 *Pemt:Phosphoethanolamine N-methyltransferase, Pcyt:Phosphate cytidyltransferase,*
 915 *Pisd:Phosphatidylserine decarboxylase, Ptdss:Phosphatidylserine synthase,*
 916 *CDS:CDP-diacylglycerol synthase, Sgms:Sphingomyelin synthase,*
 917 *Smpd:Sphingomyelin phosphodiesterase, Pis:Phosphatidylinositol synthase,*
 918 *Pgps:Phosphatidylglycerophosphate synthase, Cls:Cardiolipin synthase,*
 919 *PE:Phosphatidylethanolamine, PC: Phosphatidylcholine, PS: Phosphatidylserine, PI:*
 920 *Phosphatidylinositol, CL:Cardiolipin, PG, Phosphatidylglycerol, SM:Sphingomyelin.*

921

922 Fig. 5 Gene expression of the enzymes related to phospholipid synthesis and
 923 phospholipid composition of the rat brain. (A) The area examined by gene expression
 924 analysis (B) The ratios of the expression levels of the enzymes in the synthetic pathway
 925 of plasmalogens and diacyl phospholipids in the PFC, NAcc, and CPu for the
 926 40F-group offspring (n = 8) were compared with those for the AL-group offspring (n =
 927 8) (See Figure 5-1). (C) The area examined by MALDI-IMS. The dotted areas were
 928 examined in the PFC, NAcc, and CPu. In a section of the PFC, the dotted area is the
 929 middle third of the box area indicated by the solid line. Signal intensity was indicated
 930 by color. (D) The ratios of peak intensities of phospholipids of 40F offspring (n = 7) to
 931 those of AL offspring (n = 7) are shown (See Figure 5-2). (E) The peaks of product ions
 932 by collision-induced dissociation of m/z 774.5 in MALDI tandem mass spectrometry.

933 * $p < 0.05$, ** $p < 0.01$, Student's t -test.

934

935 Fig. 6 Behavioral tests and phospholipid composition in the brain of the rats injected
 936 with PE. (A) The experimental schedule of the liposome injection at 14 weeks of age.
 937 Changes in the behavior of rats by using the open-field test after (B) PE (18:0p-22:6)
 938 (PEL: $n = 6$, CL: $n = 7$), or (C) POPE injection (POPEL: $n = 5$, CL: $n = 5$, saline: $n = 4$)
 939 were examined by ANCOVA. Similarly, behavioral changes evaluated according to the
 940 elevated plus maze test after (D) PE (18:0p-22:6) (PEL: $n = 6$, CL: $n = 6$), or (E) POPE
 941 injection (POPEL: $n = 5$, CL: $n = 5$, saline: $n = 4$) were examined. (F) The ratios of peak
 942 intensities of phospholipids in the brains of PEL-injected rats ($n = 4$) significantly
 943 increased compared with those of CL-injected rats ($n = 4$) using Student's t -test (See
 944 also Figure 6-1). The increased ratio of peak intensity of PE (18:0p-22:6) in the PFC of
 945 PEL-injected rats was verified in an additional experiment (See Figure 6-2). (G) The
 946 ratio of PE (18:0p-22:6) to total lipids in the blood cells was compared among CL PEL
 947 and saline groups using one-way ANOVA with Tukey's HSD test. * $p < 0.05$, ns: not
 948 significant.

949

950 Fig. 3-1 Metabolomics of the plasma and CSF in male offspring aged 9 weeks. All data
 951 identified in this study are shown. The peak height of a particular ion for each
 952 metabolite was normalized to the peak height of the specified ion of 2-isopropylmalic
 953 acid in metabolic profiling. * $p < 0.05$, Student's t -test, ND: not detected.

954

955 Fig. 5-1 Gene expression profiles in male offspring aged 9 weeks. Comparative Cq
 956 values of the target genes normalized to B2m are shown. * $p < 0.05$, Student's t -test,

957 ND: not detected, NE: not examined.

958

959 Fig. 5-2 Change in phospholipid composition in PFC, NAcc, and CPu by undernutrition.

960 All data examined in this study are shown. $*p < 0.05$, Student's *t*-test.

961

962 Fig. 6-1 Phospholipid composition in PFC, NAcc, and CPu of the rats injected with

963 liposomes. All data examined in this study are shown. $*p < 0.05$, Student's *t*-test.

964

965 Fig. 6-2 Phospholipid composition in PFC of the rats injected with liposomes in the

966 verification experiment. To verify whether PE (18:0p-22:6) was elevated in the brain

967 following PE injection, phospholipid composition was analyzed in PFC sections of male

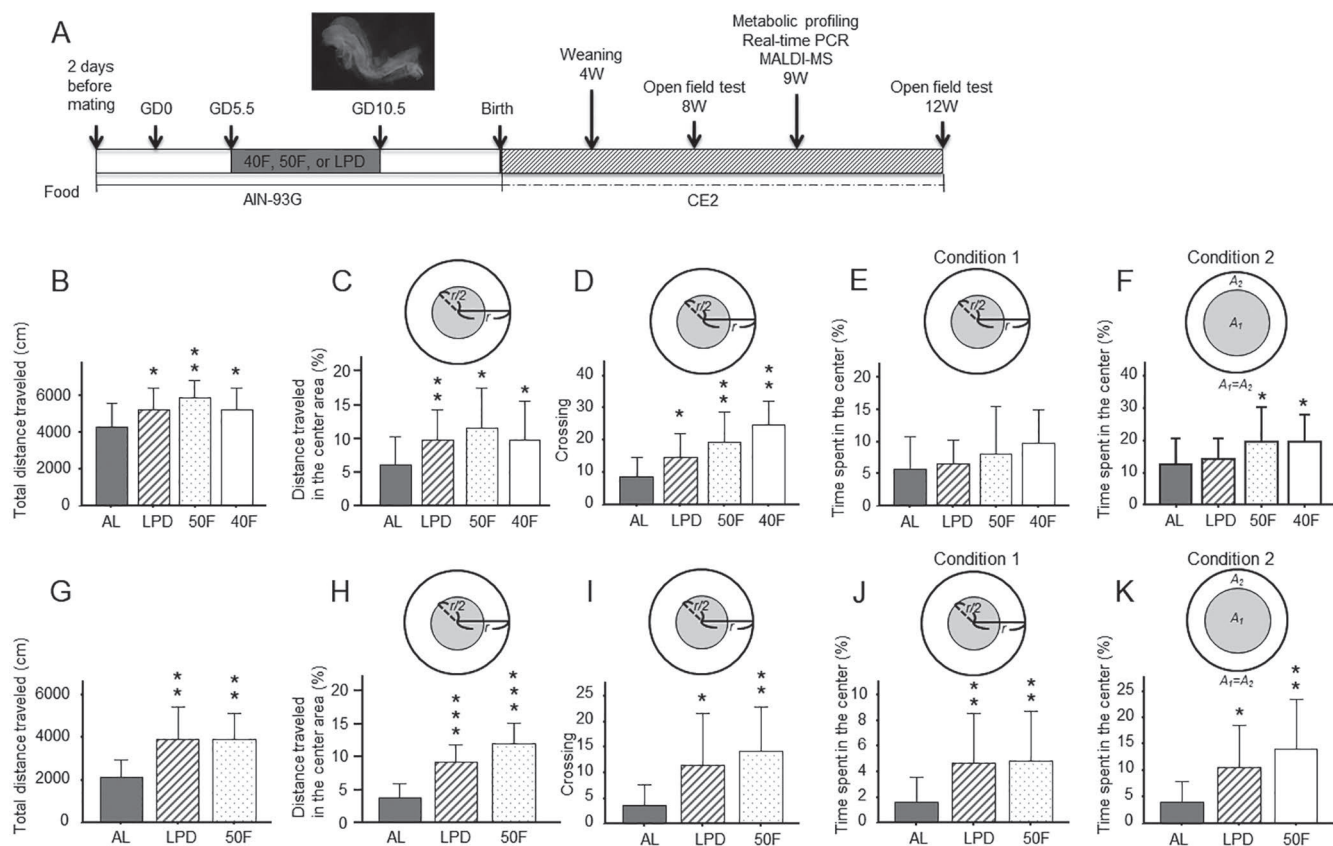
968 rats in the PEL ($n = 5$) and CL ($n = 5$) after two shots of PE (18:0p-22:6) using the

969 method described in the Materials and Methods section. This experiment was conducted

970 separately from the PEL injection study in Fig. 6F. PE (18:0p-22:6) significantly

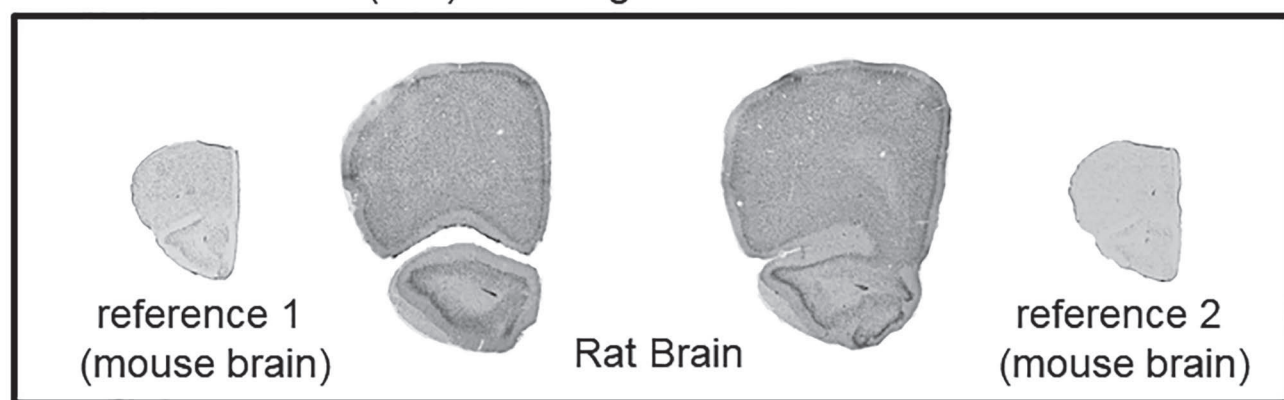
971 increased in the PFC of the rats injected with PEL liposomes ($p = 0.032$, $d = 1.83$). $*p$

972 < 0.05 , Student's *t*-test.



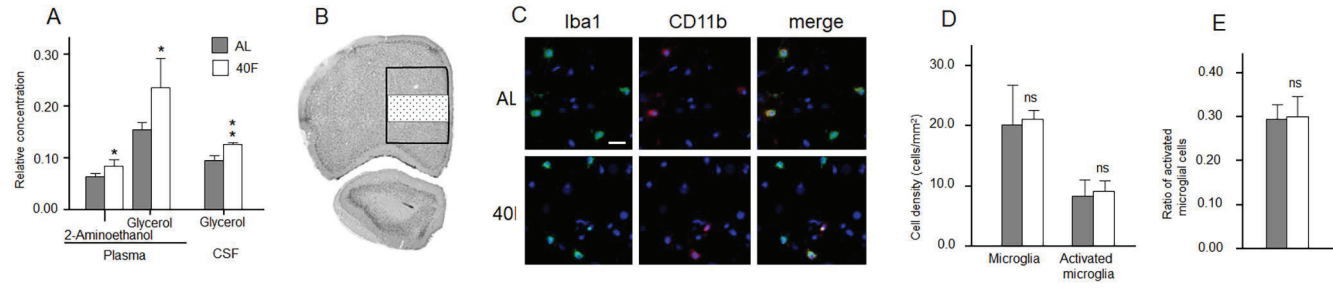
A

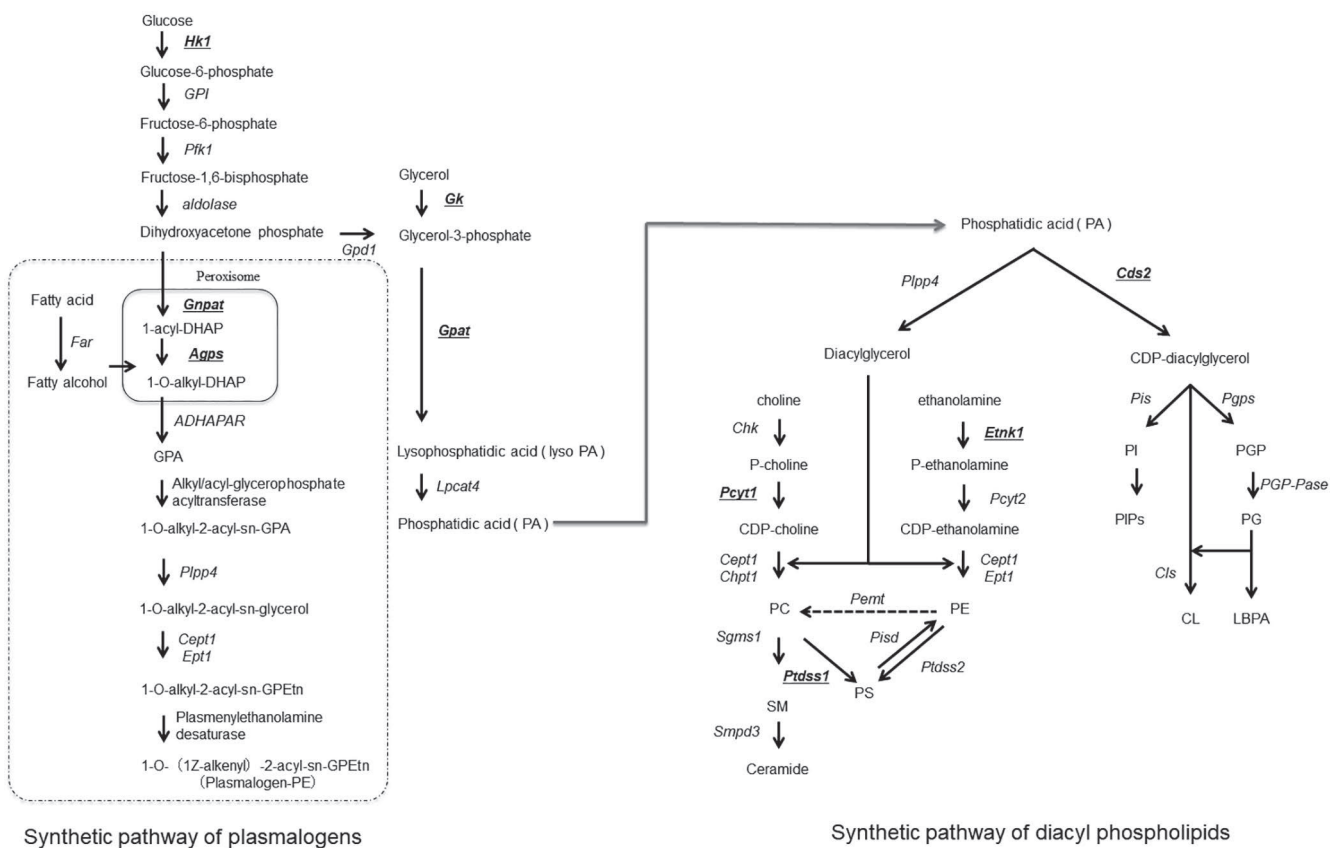
Indium tin oxide (ITO)-coated glass slide

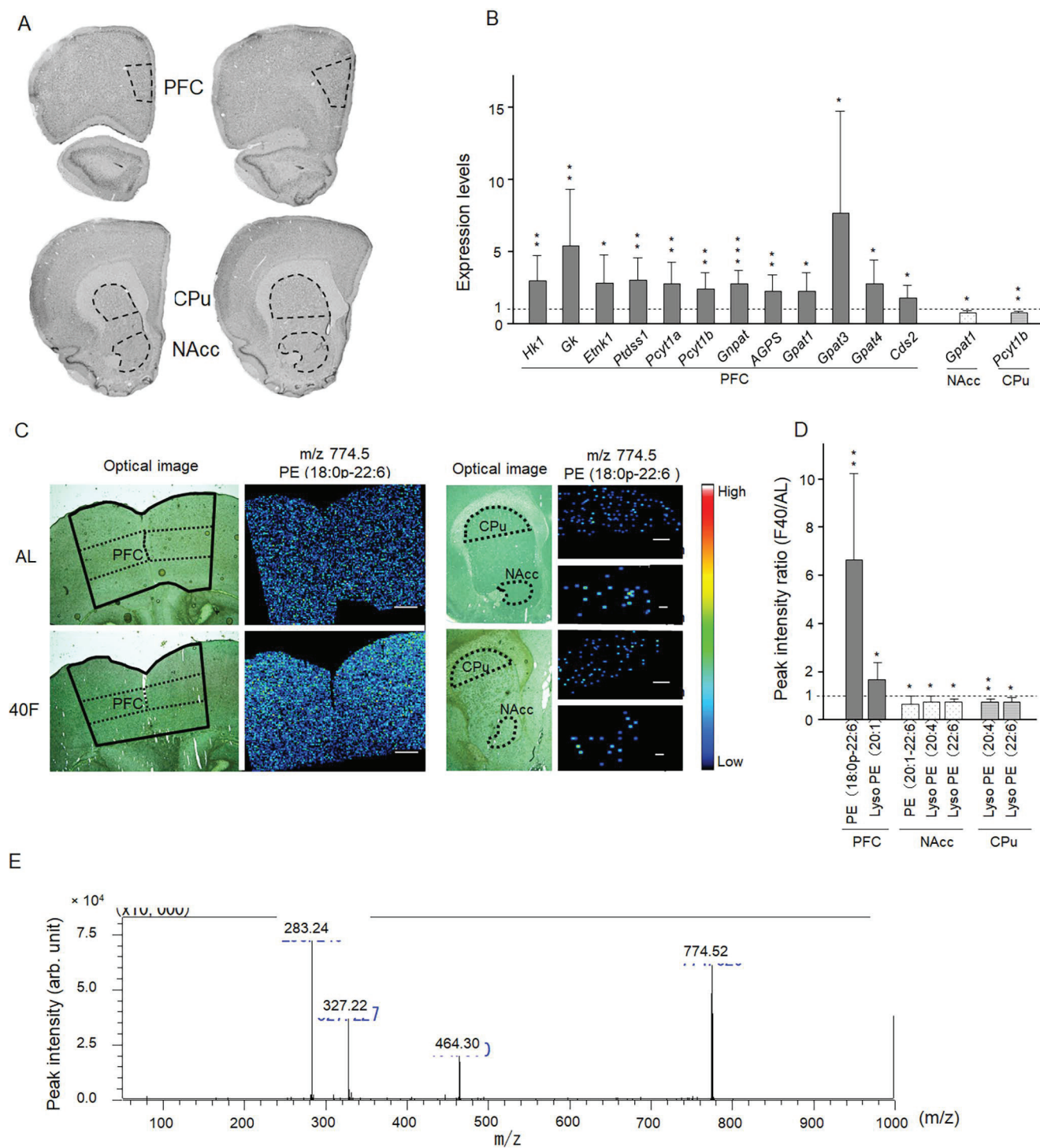


B

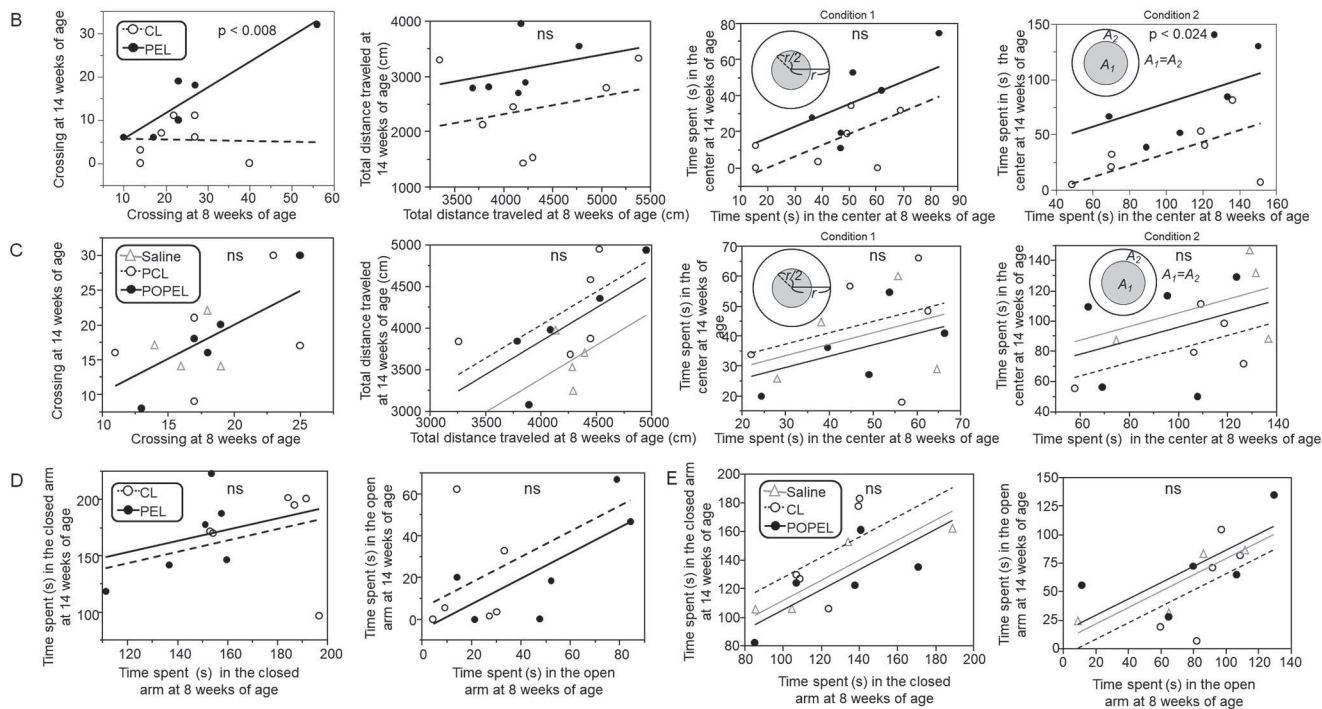
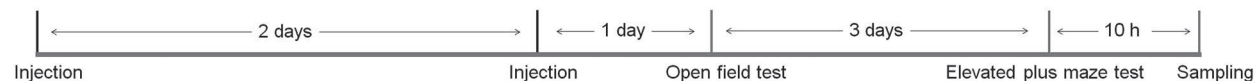
$$\text{Peak intensity} = \frac{\text{Peak intensity of phospholipid X in the rat brain}}{\text{Average raw peak intensity of phospholipid X in references 1 and 2}}$$



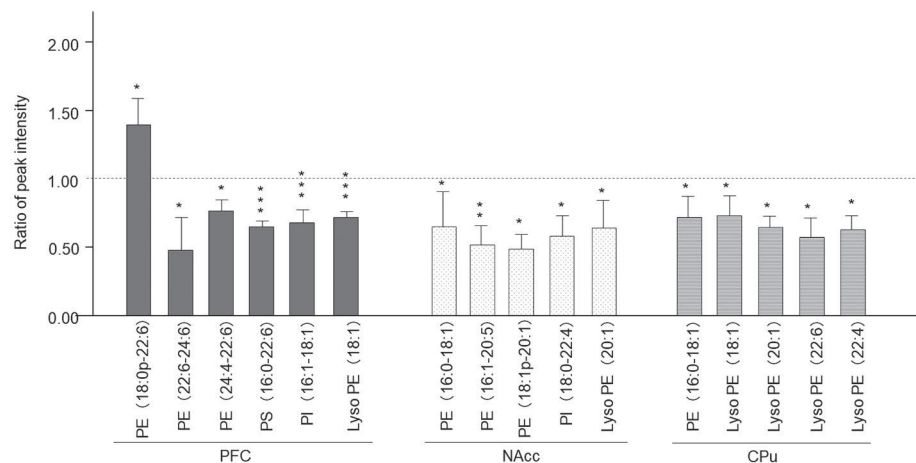




A



F



G

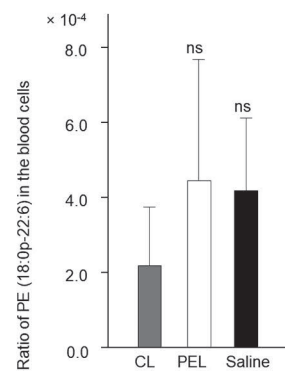


Table 1 Primer sequences for real-time PCR

Gene symbol	Forward primer sequence (5'>3')	Reverse primer sequence (5'>3')	Acc. number	Amplicon size (bp)
<i>Agps</i>	TCCTACTCACAAGACGCAGA	AGGAATCCGCTCAAACATCC	NM_053350.2	99
<i>ApoE</i>	CCGCAACGAGGTAAACACCA	ATCATCCGCATCCCGCATC	NM_001270684.1	109
<i>Aqp9</i>	AGCCGGATAGCGAAGGAGA	AGTGATGATCCCGCCAAAAC	NM_022960.2	120
<i>B2M</i>	CGAGACCGATGTATATGCTTGC	GTCCAGATGATTCAGAGCTCCA	NM_012512.2	114
<i>Cds2</i>	CCACCGGTTTCATCTCCTTTAC	GCGTTACGACAATAAGCAAGG	NM_053643.1	134
<i>Cept1</i>	AGTCTTCTACTGCCCTACAGC	TTCTTCTGGCCTGTTTCCCG	NM_001007.1	119
<i>Chka</i>	CAACAACTGCACAAGTTCCTC	CTCTTGGCCTTCCAACAATAAG	NM_017127.1	144
<i>Chkb</i>	GTCCACTAGCCTTCCCCAGA	CCTGGATGTCATTGTGGCAG	NM_017177.1	125
<i>Etnk1</i>	AGCTCGTCAGCTTGCTAAAATC	CTTCATCAGCAAATCCTGTGGG	NM_001107894.1	122
<i>Far1</i>	CCCTTGCGATCTCGTTCTCT	AGGATTAGTGCTGCCTGTTGT	XM_006230020.3	125
<i>Gk</i>	GAAACTTCGTTGGCTCCTCG	GAACACCGCCATTGATTCCC	NM_024381.2	128
<i>Gnpat</i>	GTGTGTGTGAATGAAGAAGGCA	GGACAAAGGACAGCATGAGGA	NM_053410.1	115
<i>Gpat1</i>	GGCAACAACCTCAACATCCC	TTGCGTCCATCTGGAGTTTC	NM_017274.1	98
<i>Gpat3</i>	AACTGTTGGCCAGCTTC	AGACAGGAGCGAACACAGA	XM_008770021.2	96
<i>Gpat4</i>	GCTCAAACCAGACATGGGGG	TTGCTAACCATACGTCGCCC	NM_001047849.1	141
<i>Gpd1</i>	AGCATCCTCCAACACAAGGG	GCAGCAGATGAACACACCA	NM_022215.2	102
<i>Hk1</i>	ACCCGGAATTCGTTCTCTCC	TCTTTTGTCCGGAGCATCCC	NM_012734.1	80
<i>iPLA2</i>	CCCATCCACACAGCCATGAA	AGAAGCATTCGGGCCATCTC	XM_006242004.2	149
<i>Lpcat4</i>	GGAGCAGCTTCAGGAACCAA	CGAAGGAAGCCAAGCAGGAA	NM_001106494.1	102
<i>Mfsd2a</i>	GCTTCTGCATCAGCAAGTCC	GGGAAGTCAGGCACAAACCA	NM_001106683.1	118
<i>Mfsd3</i>	GGTGCTTCTGCCTCAGATTT	GCACTGACCTCAACAGCTTC	NM_001024908.1	142
<i>Plpp4</i>	TGCTTTCCAGATGGGGTGATG	CAACTTGCCAGCCAGGTAGAA	NM_001191631.1	150
<i>Pcyt1a</i>	ATCCCTACTCTTCGGCAGG	GGACAATGCGGGTGATGAGG	NM_078622.2	121
<i>Pcyt1b</i>	TCATCTCGGGGTCTGATGAC	ACGGACAATTCTGGTGATGATG	NM_173151.1	114
<i>Pcyt2</i>	ATGTGGCTGGTGCCTTTGA	AAAGTGATAGCCCGCGATGA	NM_053568.1	104
<i>Pemt</i>	GTAAGGGGAAGTACAGCCAAC	CTTCAAACAGGAGAGCAACCAC	NM_013003.1	116
<i>Pfkfb</i>	AAGTACCTGGAGCACCTCTCT	TGTATATTTCCATGCGCACCA	NM_206847.1	115
<i>Pisd</i>	ATGTGGGCTCTATCCGCATC	TGGGAATGCCCTCCTTGTTG	XM_002725000.4	120
<i>Ptdss1</i>	TATGGGCTCTGCTGGACAATC	CCACCACCATTACACAACAGG	NM_001012133.1	131
<i>Ptdss2</i>	AACCCCTCAGGATACAGCTAC	CTAACACACAGCCAAAACCGC	NM_001106316.1	144
<i>Smpd3</i>	TGTCTCAACAGCGGTCTCTTC	GTACAGGCGATGTACCAACA	NM_053605.1	176
<i>Sgms1</i>	ATGCTAACGCTCACCTACCTG	GTGTAGTGGTCATGCGCTAA	NM_181386.2	128

Table 2 Allocation of experimental groups

Treatment	Experiment	Age (weeks)	
Under nutrition	Open-field test	8	AL, 40F, 50F, LPD
	Open-field test	12	AL, 50F, LPD
	Metabolic profiling (plasma, CSF)	9	AL, 40F
	Counting microglial cells (PFC)	9	AL, 40F
	Gene expression (PFC, NAcc, CPu)	9	AL, 40F
	MALDI-IMS (PFC, NAcc, CPu)	9	AL, 40F
Liposome injection	Open-field test	8 and 14	CL, PEL, POPEL, saline
	Elevated plus maze test	8 and 14	CL, PEL, POPEL, saline
	MALDI-IMS (PFC, NAcc, CPu, blood cells)	14	CL, PEL

AL: *ad libitum* group, 40F or 50F: the group fed 40% or 50% of the daily food intake of the AL, respectively, LPD: low protein diet group

CL, PEL, POPEL, or saline: the group injected with control, PE (18:0p-22:6), or POPE liposome, or saline

Table 3 Body weight of the offspring

Group	Body weight (g)	
	9 weeks of age	12 weeks of age
AL	271.4 ± 26.5 (n = 32)	315.5 ± 26.0 (n = 18)
50F	248.7 ± 20.6* (n = 18)	317.7 ± 26.2 (n = 18)
40F	267.8 ± 28.8 (n = 18)	Not measured
LPD	250.4 ± 17.3* (n = 19)	325.7 ± 18.5 (n = 19)

Values are means ± SD, * $p < 0.05$ vs AL (One-way ANOVA with Dunnett's test)

Table 4 Change in amounts of DHA- or AA-containing phospholipids in PFC, NAcc, and CPu by undernutrition

Phospholipid	sn1_sn2	PFC		NAcc		CPu	
		AL	40F	AL	40F	AL	40F
		Mean \pm SD	Mean \pm SD	Mean \pm SD	Mean \pm SD	Mean \pm SD	Mean \pm SD
Lyso PE_AA	20:4 (Acyl)	3.86 \pm 1.20	6.30 \pm 2.70	3.60 \pm 0.67	2.60 \pm 0.77*	3.76 \pm 0.57	2.75 \pm 0.39*
Lyso PE_DHA	22:6 (Acyl)	2.54 \pm 0.84	3.67 \pm 1.17	2.23 \pm 0.46	1.70 \pm 0.26*	2.48 \pm 0.33	1.92 \pm 0.38*
PE_AA_Pls	16:0p-20:4 (Alkenyl_acyl)	2.83 \pm 0.95	3.97 \pm 1.70	4.52 \pm 1.46	4.52 \pm 0.86	3.27 \pm 0.74	3.46 \pm 0.76
	18:1p-20:4 (Alkenyl_acyl)	2.07 \pm 0.72	2.48 \pm 1.03	2.15 \pm 0.41	2.02 \pm 0.43	1.94 \pm 0.27	1.90 \pm 0.42
PE_AA_Acyl	18:1-20:4 (Alkyl_acyl)	2.71 \pm 0.86	3.39 \pm 1.31	3.65 \pm 0.82	3.38 \pm 0.63	2.89 \pm 0.50	3.00 \pm 0.62
	18:0-20:4 (Alkyl_acyl)	2.95 \pm 0.73	4.29 \pm 1.87	3.83 \pm 0.95	3.21 \pm 0.72	3.14 \pm 0.76	2.97 \pm 0.59
	18:2-20:4 (Diacyl)	1.37 \pm 0.47	1.67 \pm 0.67	1.21 \pm 0.22	1.19 \pm 0.26	1.18 \pm 0.17	1.19 \pm 0.27
	18:1-20:4 (Diacyl)	1.67 \pm 0.54	2.10 \pm 0.88	1.64 \pm 0.31	1.45 \pm 0.26	1.67 \pm 0.20	1.49 \pm 0.27
	18:0-20:4 (Diacyl)	1.84 \pm 0.56	2.35 \pm 0.96	2.38 \pm 0.32	2.11 \pm 0.42	1.74 \pm 0.19	1.72 \pm 0.40
PE_DHA_Pls	18:0p-22:6 (Alkenyl_acyl)	1.41 \pm 0.43	9.39 \pm 4.62*	1.74 \pm 0.32	1.67 \pm 0.30	1.79 \pm 0.28	1.75 \pm 0.32
	18:1p-22:6 (Alkenyl_acyl)	1.30 \pm 0.36	1.63 \pm 0.65	1.28 \pm 0.24	1.16 \pm 0.20	1.34 \pm 0.26	1.26 \pm 0.19
	20:1p-22:6 (Alkenyl_acyl)	3.11 \pm 1.38	3.46 \pm 0.85	2.09 \pm 0.69	1.77 \pm 0.82	2.48 \pm 1.14	2.13 \pm 0.49
PE_DHA_Acyl	16:1-22:6 (Diacyl)	1.36 \pm 0.38	1.64 \pm 0.84	1.80 \pm 0.41	1.55 \pm 0.26	1.86 \pm 0.24	1.66 \pm 0.20
	18:0-22:6 (Alkyl_acyl)	1.67 \pm 0.46	2.19 \pm 0.92	1.94 \pm 0.45	1.70 \pm 0.34	1.93 \pm 0.35	1.93 \pm 0.41
	18:2-22:6 (Diacyl)	1.75 \pm 0.66	2.27 \pm 1.09	1.56 \pm 0.29	1.37 \pm 0.31	1.93 \pm 0.32	1.73 \pm 0.42
	18:0-22:6 (Diacyl)	1.31 \pm 0.42	1.67 \pm 0.63	1.46 \pm 0.28	1.46 \pm 0.31	1.44 \pm 0.22	1.48 \pm 0.31
	20:1-22:6 (Diacyl)	3.00 \pm 1.17	3.65 \pm 1.96	2.72 \pm 0.71	1.76 \pm 0.77*	3.08 \pm 0.91	2.69 \pm 1.20
	20:0-22:6 (Diacyl)	2.16 \pm 0.85	3.42 \pm 1.71	3.28 \pm 1.08	2.58 \pm 1.73	4.47 \pm 1.45	3.74 \pm 2.08
	22:6-22:6 (Diacyl)	1.86 \pm 0.72	2.35 \pm 0.98	1.87 \pm 0.35	1.76 \pm 0.40	1.74 \pm 0.29	1.73 \pm 0.39
	22:6-24:6 (Diacyl)	2.74 \pm 2.55	2.90 \pm 0.55	1.41 \pm 0.40	1.05 \pm 0.28	3.52 \pm 1.30	3.23 \pm 0.81

	24:4-22:6 (Diacyl)	2.97 ± 1.20	3.55 ± 1.41	1.74 ± 0.58	2.01 ± 1.46	2.24 ± 0.80	2.49 ± 1.23
PS_AA_Acyl	18:1-20:4 (Diacyl)	2.00 ± 0.63	2.94 ± 1.50	3.50 ± 1.28	3.16 ± 0.76	6.01 ± 1.91	6.38 ± 1.87
PS_DHA_Acyl	16:0-22:6 (Diacyl)	1.46 ± 0.47	2.39 ± 1.13	3.91 ± 0.94	3.35 ± 0.68	7.26 ± 1.62	7.21 ± 1.56
	18:1-22:6 (Diacyl)	2.64 ± 0.97	3.12 ± 1.57	1.73 ± 0.63	1.63 ± 0.43	2.14 ± 0.46	1.88 ± 0.38
	20:0-22:6 (Diacyl)	2.11 ± 0.67	4.29 ± 2.46	2.12 ± 0.92	1.89 ± 0.54	2.72 ± 1.02	2.47 ± 0.50
	22:4-22:6 (Diacyl)	2.10 ± 1.19	3.56 ± 2.64	1.56 ± 0.22	1.56 ± 0.44	1.91 ± 0.29	1.77 ± 0.45
PS_AA_DHA	20:4-22:6 (Diacyl)	2.09 ± 0.77	2.52 ± 1.08	2.02 ± 0.57	1.82 ± 0.44	2.01 ± 0.69	2.16 ± 0.77
Lyso PI_AA	20:4 (Acyl)	3.40 ± 1.67	4.53 ± 2.83	4.39 ± 1.03	4.06 ± 2.43	4.22 ± 0.64	3.99 ± 2.32
PI_AA_Acyl	16:1-20:4 (Diacyl)	5.79 ± 8.33	4.14 ± 2.30	1.92 ± 0.39	1.87 ± 0.50	2.44 ± 0.66	2.21 ± 0.22
	16:0-20:4 (Diacyl)	2.07 ± 0.84	2.38 ± 1.07	4.77 ± 7.19	1.44 ± 0.49	4.67 ± 6.92	1.52 ± 0.50
	18:0-20:4 (Alkyl_acyl)	2.83 ± 1.19	2.67 ± 1.19	1.98 ± 0.56	1.82 ± 0.28	2.27 ± 0.53	2.11 ± 0.30
	18:1-20:4 (Diacyl)	1.89 ± 0.77	2.23 ± 1.01	1.70 ± 0.36	1.42 ± 0.49	1.59 ± 0.32	1.43 ± 0.42
	18:0-20:4 (Diacyl)	2.28 ± 0.83	2.79 ± 1.19	2.71 ± 0.77	2.66 ± 0.78	2.80 ± 0.39	2.69 ± 0.75
PI_DHA_Acyl	18:0-22:6 (Diacyl)	2.22 ± 0.87	2.82 ± 1.80	1.17 ± 0.33	1.17 ± 0.35	1.64 ± 0.33	1.34 ± 0.56

* $p < 0.05$

Table 5 Change in amounts of DHA- or AA-containing phospholipids in PFC, NAcc, and CPu of the rats injected with liposomes

Phospholipid	sn1_sn2	PFC		NAcc		CPu	
		PEL	CL	PEL	CL	PEL	CL
		Mean \pm SD	Mean \pm SD	Mean \pm SD	Mean \pm SD	Mean \pm SD	Mean \pm SD
Lyso PE_AA	20:4 (Acyl)	1.94 \pm 0.21	2.17 \pm 0.24	1.44 \pm 0.79	2.15 \pm 0.88	1.31 \pm 0.15	1.87 \pm 0.58
Lyso PE_DHA	22:6 (Acyl)	1.29 \pm 0.23	1.55 \pm 0.09	1.00 \pm 0.45	1.24 \pm 0.28	0.88 \pm 0.19*	1.55 \pm 0.32
PE_AA_Pls	16:0p-20:4 (Alkenyl_acyl)	4.29 \pm 0.71	3.66 \pm 0.59	3.02 \pm 1.03	3.36 \pm 0.50	2.19 \pm 0.43	2.58 \pm 0.62
	18:1p-20:4 (Alkenyl_acyl)	2.25 \pm 0.41	2.07 \pm 0.32	1.13 \pm 0.36	1.36 \pm 0.14	1.06 \pm 0.24	1.24 \pm 0.20
PE_AA_Acyl	18:1-20:4 (Alkyl_acyl)	3.22 \pm 0.54	2.70 \pm 0.58	2.26 \pm 0.73	2.48 \pm 0.19	1.88 \pm 0.42	2.14 \pm 0.36
	18:0-20:4 (Alkyl_acyl)	3.11 \pm 0.58	2.28 \pm 0.27	2.11 \pm 0.65	2.56 \pm 0.32	1.82 \pm 0.40	2.30 \pm 0.29
	18:2-20:4 (Diacyl)	1.87 \pm 0.28	1.50 \pm 0.24	0.65 \pm 0.18	0.73 \pm 0.05	0.68 \pm 0.13	0.78 \pm 0.10
	18:1-20:4 (Diacyl)	1.69 \pm 0.39	1.43 \pm 0.28	1.02 \pm 0.24	1.12 \pm 0.17	1.06 \pm 0.20	1.19 \pm 0.24
	18:0-20:4 (Diacyl)	2.60 \pm 0.56	1.97 \pm 0.60	1.30 \pm 0.35	1.50 \pm 0.12	1.07 \pm 0.20	1.22 \pm 0.16
PE_DHA_Pls	18:0p-22:6 (Alkenyl_acyl)	2.62 \pm 0.31*	1.88 \pm 0.08	1.06 \pm 0.30	1.18 \pm 0.05	1.08 \pm 0.18	1.18 \pm 0.13
	18:1p-22:6 (Alkenyl_acyl)	2.13 \pm 0.31	1.99 \pm 0.16	0.73 \pm 0.23	0.88 \pm 0.11	0.84 \pm 0.15	1.00 \pm 0.15
	20:1p-22:6 (Alkenyl_acyl)	2.99 \pm 1.29	2.39 \pm 0.22	1.00 \pm 0.46	1.04 \pm 0.26	1.33 \pm 0.30	1.92 \pm 0.64
PE_DHA_Acyl	16:1-22:6 (Diacyl)	1.10 \pm 0.31	1.03 \pm 0.19	0.97 \pm 0.27	1.20 \pm 0.28	1.26 \pm 0.26	1.32 \pm 0.27
	18:0-22:6 (Alkyl_acyl)	2.52 \pm 0.38	1.87 \pm 0.35	0.99 \pm 0.31	1.23 \pm 0.08	1.17 \pm 0.22	1.33 \pm 0.15
	18:2-22:6 (Diacyl)	1.29 \pm 0.24	1.27 \pm 0.20	0.86 \pm 0.17	1.08 \pm 0.22	1.04 \pm 0.34	1.31 \pm 0.31
	18:0-22:6 (Diacyl)	1.63 \pm 0.32	1.28 \pm 0.33	0.82 \pm 0.20	0.92 \pm 0.10	0.84 \pm 0.14	0.93 \pm 0.11
	20:1-22:6 (Diacyl)	1.94 \pm 0.36	2.05 \pm 0.46	0.88 \pm 0.31	1.08 \pm 0.24	0.92 \pm 0.44	1.19 \pm 0.39
	20:0-22:6 (Diacyl)	1.85 \pm 0.46	2.12 \pm 0.21	1.90 \pm 1.23	3.23 \pm 0.69	2.68 \pm 0.40	3.44 \pm 0.58
	22:6-22:6 (Diacyl)	1.91 \pm 0.41	1.71 \pm 0.40	1.05 \pm 0.32	1.31 \pm 0.13	1.02 \pm 0.21	1.20 \pm 0.21
	22:6-24:6 (Diacyl)	3.74 \pm 1.60*	7.77 \pm 0.78	0.61 \pm 0.28	0.93 \pm 0.59	1.68 \pm 0.44	2.23 \pm 0.50

	24:4-22:6 (Diacyl)	2.28 ± 0.22*	2.98 ± 0.19	1.08 ± 0.75	1.54 ± 0.20	0.98 ± 0.22	1.43 ± 0.55
PS_AA_Acyl	18:1-20:4 (Diacyl)	1.04 ± 0.14	1.34 ± 0.14	1.73 ± 0.67	2.03 ± 0.25	2.77 ± 0.69	3.32 ± 0.49
PS_DHA_Acyl	16:0-22:6 (Diacyl)	0.60 ± 0.03*	0.91 ± 0.04	1.60 ± 0.62	2.41 ± 1.04	3.56 ± 0.92	4.71 ± 0.98
	18:1-22:6 (Diacyl)	1.80 ± 0.39	1.58 ± 0.18	1.44 ± 0.79	1.23 ± 0.15	1.26 ± 0.41	1.55 ± 0.29
	20:0-22:6 (Diacyl)	1.74 ± 0.29	2.36 ± 0.49	1.00 ± 0.45	1.65 ± 0.45	1.63 ± 0.36	1.69 ± 0.45
	22:4-22:6 (Diacyl)	7.68 ± 2.36	5.82 ± 1.23	3.02 ± 1.03	1.21 ± 0.25	1.16 ± 0.30	1.37 ± 0.22
PS_AA_DHA	20:4-22:6 (Diacyl)	2.33 ± 0.40	2.51 ± 0.32	1.13 ± 0.36	1.54 ± 0.49	1.14 ± 0.34	1.22 ± 0.25
Lyso PI_AA	20:4 (Acyl)	2.80 ± 0.19	3.27 ± 0.32	2.26 ± 0.73	1.33 ± 0.22	1.13 ± 0.31	1.34 ± 0.30
PI_AA_Acyl	16:1-20:4 (Diacyl)	1.88 ± 0.47	1.31 ± 0.08	2.11 ± 0.65	1.71 ± 0.11	1.49 ± 0.42	1.67 ± 0.43
	16:0-20:4 (Diacyl)	2.08 ± 0.43	1.88 ± 0.37	0.65 ± 0.18*	1.32 ± 0.22	0.94 ± 0.21	1.32 ± 0.19
	18:0-20:4 (Alkyl_acyl)	2.13 ± 0.37	2.63 ± 0.45	1.02 ± 0.24	1.25 ± 0.15	1.17 ± 0.31	1.23 ± 0.25
	18:1-20:4 (Diacyl)	2.43 ± 0.24	2.27 ± 0.33	1.30 ± 0.35	1.11 ± 0.23	0.95 ± 0.30	1.11 ± 0.33
	18:0-20:4 (Diacyl)	2.41 ± 0.38	2.25 ± 0.35	1.11 ± 0.20	1.84 ± 0.22	1.49 ± 0.32	1.70 ± 0.30
PI_DHA_Acyl	18:0-22:6 (Diacyl)	1.64 ± 0.12	1.75 ± 0.29	0.73 ± 0.23	1.96 ± 0.16	1.43 ± 0.58	1.54 ± 0.17

* $p < 0.05$