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Changes in the lipid composition of *Bradyrhizobium* cell envelope reveal a rapid response to water deficit involving lysophosphatidylethanolamine synthesis from phosphatidylethanolamine in outer membrane

Adriana B. Cesari, Natalia S. Paulucci, María A. Biasutti, Gustavo M. Morales, Marta S. Dardanelli

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1 **Changes in the lipid composition of *Bradyrhizobium* cell envelope reveal a rapid**
2 **response to water deficit involving lysophosphatidylethanolamine synthesis from**
3 **phosphatidylethanolamine in outer membrane**

4
5 Adriana B. Cesari^a, Natalia S. Paulucci^{a*}, María A. Biasutti^b, Gustavo M. Morales^b, Marta
6 S. Dardanelli^a

7 a- Departamento de Biología Molecular, Facultad de Ciencias Exactas, Físico-
8 Químicas y Naturales, Universidad Nacional de Río Cuarto, Río Cuarto, Córdoba,
9 Argentina

10 b- Departamento de Química, Facultad de Ciencias Exactas, Físico-Químicas y
11 Naturales, Universidad Nacional de Río Cuarto, Río Cuarto, Córdoba, Argentina

12
13
14
15 acesari@exa.unrc.edu.ar

16 npaulucci@exa.unrc.edu.ar “*Correspondence and reprints”

17 abiasutti@exa.unrc.edu.ar

18 gmorales@exa.unrc.edu.ar

19 mdardanelli@exa.unrc.edu.ar

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26 **Abstract**

27 We evaluate the behavior of the membrane of *Bradyrhizobium* sp. SEMIA6144 during
28 adaptation to polyethylene glycol (PEG). A dehydrating effect on the morphology of the
29 cell surface, as well as a fluidizing effect on the membrane was observed 10 min after PEG
30 shock; however, the bacteria were able to restore optimal membrane fluidity.

31 Shock for 1 h caused an increase of lysophosphatidylethanolamine in the outer membrane
32 at the expense of phosphatidylcholine and phosphatidylethanolamine, through an increase
33 in phospholipase activity. The amount of lysophosphatidylethanolamine did not remain
34 constant during PEG shock, but after 24 h the outer membrane was composed of large
35 amounts of phosphatidylcholine and less amount of lysophosphatidylethanolamine similar
36 to the control. The inner membrane composition was also modified after 1 h of shock,
37 observing an increase of phosphatidylcholine at the expense of phosphatidylethanolamine,
38 the proportions of these phospholipids were then modified to reach 24 h of shock values
39 similar to the control.

40 Vesicles prepared with the lipids of cells exposed to 1 h shock presented higher rigidity
41 compared to the control, indicating that changes in the composition of phospholipids after 1
42 h of shock restoring fluidity after the PEG effect and would allow cells to maintain surface
43 morphology.

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45

46

47 **Keywords:** *Bradyrhizobium* peanut-nodulating, water deficit, envelope cell topography,
48 membrane fluidity, phospholipids remodeling.

49

50 **1. Introduction**

51

52 *Bradyrhizobium* sp. SEMIA6144 is a Gram-negative bacterium that is able to fix
53 atmospheric nitrogen (N) in symbiotic association with peanut plants to enhance plant N-
54 content [1,2]. However, physiological stresses such as water deficit, salinity and extreme
55 temperature negatively affect this symbiotic interaction. Water deficit is the most important
56 limiting factor for crop production in many parts of the world, and fluctuating water
57 availability is one of the primary environmental factors that affects the activity and viability
58 of soil microorganisms [3,4]. High-molecular-weight (i.e., a molecular weight of 6000 Da)
59 polyethylene glycol (PEG) has been utilized extensively in microbial and plant studies
60 examining responses to reduced water content and is believed to simulate water deficit [5].

61 The lipid matrix of bacterial membranes comprises structured phospholipid (PL) assemblies
62 in a bimolecular layer and integrated proteins [6]. There is general agreement that
63 phosphatidylethanolamine (PE), phosphatidylcholine (PC), phosphatidylglycerol (PG) and
64 cardiolipin (CL) are the major PL in rhizobia [6]. However, PC is restricted to a limited
65 number of genera, being a clear example of this is the genus *Bradyrhizobium* [2]. Microbial
66 fatty acids (FA) are typically 14-20 carbons in length [7, 8, 9]. Acyl chains of this length
67 balance the stability and fluidity needs of the membrane [6]. Paulucci [10] reported that the
68 major fatty acid (FA) in SEMIA6144 are the unsaturated FA (UFA) 18:1 Δ 11 and the
69 saturated FA (SFA) 16:0 and 18:0.

70 Environmental factors can alter the packing order of the bacterial cell membrane and
71 change the bilayer structure [6]. The cell envelope is in constant contact with the
72 environment, and thus flexibility of the membranes and its components to facilitate

73 adaptation is a primary determinant of cell survival. There are several techniques for
74 evaluating membrane integrity within which atomic force microscopy (AFM) is used to
75 monitor changes in the cell surface produced by different environmental factors [11,12].
76 Lipid membranes can undergo changes upon exposure to adverse environmental factors
77 through alterations in the polar head groups or acyl chains of PL [2, 13, 14]. These changes
78 may be accomplished by varying the amount of total PL and/or their class distribution,
79 changing the type of FA incorporated into PL during synthesis or turnover via
80 phospholipase activity, or regulating the degree of unsaturation of FA moieties [15, 13, 14].
81 As intermediates in PL metabolism and turnover, lysophospholipids, as
82 lysophosphatidylethanolamine (LPE), are produced via phospholipase A (PLA) mediated
83 hydrolysis of the acyl group of a PL to lysophospholipids and FA [16].
84 Water deficits affect membranes integrity. Scherber [17] reported an increase in the
85 membrane phase temperature transition of *Escherichia coli* after extensive cell water deficit
86 (1 week at 20 % - 40 % relative humidity). These authors also found that *E. coli*
87 membranes exposed to water deficit exhibit high levels of SFA compared to undried cells.
88 In addition, Cesari [14] showed that *Azospirillum brasilense* can perceive osmotic changes
89 in medium by sensing modifications in membrane fluidity. Under these conditions, the
90 bacterium responds by increasing PC and SFA levels to maintain membrane integrity.
91 Several studies in bacteria have elucidated the roles of membrane components in water
92 deficit tolerance [17, 18, 13, 14]; however, there have been few studies exploring how
93 within the cell envelope, the membranes and its components change during a stress
94 situation to achieve optimal fluidity and ensure survival. In addition, many questions
95 remain regarding the physiological and biochemical responses of rhizobia to water deficit;

96 prior work in this area has focused on improving the long-term tolerance of inoculant-based
97 rhizobia rather than investigating bacterial physiology under water deficit conditions.

98 *Escherichia coli* has been widely used as a model for predicting behaviors of both outer and
99 inner membrane of other bacteria against different environmental conditions. However, in
100 the case of rhizobia it is very difficult to make such comparisons due to the fact that they
101 contain PC in their membranes, which is absent in *E. coli*.

102 The objective of this work was to elucidate the dynamic changes that occur in the lipid
103 components of the cellular envelope of a strain of the genus *Bradyrhizobium* peanut
104 nodulant, exposed to water deficit.

105 We have placed special emphasis on elucidating the commitment of each of the membranes
106 (outer and inner) in the response to water deficit. Moreover, in this paper, we observed a
107 dehydrating effect of PEG on the cell surface and explained how *B. sp.* SEMIA6144
108 responded rapidly to the presence of PEG and counteracted that effect on the membrane by
109 modifying the lipid composition. We also demonstrate that this first response is modified
110 during adaptation to achieve optimal membrane fluidity.

111 The identification of physical and physiological factors that determine the survival of *B. sp.*
112 SEMIA6144 under water deficit conditions may aid in the strategic development of
113 rhizobial strains with improved ability to survive water deficit.

114

115 **2. Materials and Methods**

116

117 **2.1. Bacteria and growth conditions**

118 *Bradyrhizobium sp.* SEMIA6144, provided by MIRCEN/FEPAGRO, Brazil (SEMIA6144
119 in the rest of text), was maintained in yeast extract mannitol (YEM) medium [18] and

120 grown on B-modified medium [2]. Bacterial media were supplemented with the
121 nonpermeating solute polyethylene glycol (PEG, average MW 5489 Da, Sigma Chemical
122 Co., St. Louis, MO, USA) to simulate moderate water deficit [14].

123 All cultures were incubated at 28 °C with shaking at 150 rpm (Allied Fisher Scientific). The
124 water potential of 15 mmol l⁻¹ (90 g l⁻¹) PEG medium at 25 °C was measured using an
125 osmometer (Semi Micro K-7400, Knauer) and was found to be -0.8 MPa.

126 Viable SEMIA6144 cells were counted as colony forming units per ml of culture (CFU ml⁻¹)
127 using the droplet technique [20].

128

129 **2.2. Effects of PEG Shock on SEMIA6144 Viability**

130 SEMIA6144 cells were grown until the exponential growth phase was reached (48 h); then
131 to induce osmotic shock, 15 mmol l⁻¹ PEG was added to the cultures. The cultures were
132 then incubated for 1 h, 5 h and 24 h, and viability was determined using the droplet
133 technique [20].

134

135 **2.3. Atomic Force Microscopy Imaging of SEMIA6144 Cells**

136 We used AFM to investigate if PEG caused changes in cell morphology. Coverslips coated
137 with polyethylenimine (0.1 %) were used to firmly immobilize bacterial cells so they could
138 withstand the lateral forces inflicted by the cantilever during AFM imaging. Briefly,
139 coverslips were cleaned in 70 % ethanol for 10 min, rinsed with MilliQ water, and air-dried
140 in upright position. Then polyethylenimine (0.1 %) was added on each coverslip, air-dried
141 20 min at room temperature. Finally, excessive polyethylenimine were removed by
142 submerging coverslips in MilliQ water. The coverslips were air-dried, and stored for a
143 maximum of 2 weeks at 4 °C.

144 We immobilized cells by applying 20 μ L cell suspension to the coverslips covered with
145 polyethylenimine, and let the cells settle for 15 min at room temperature before removing
146 non-adhered cells by rinsing with MilliQ water.

147 The images were obtained through the use of an atomic force microscope (AFM, Agilent,
148 Technologies, SPM model 5500) working in acoustic mode. AFM probes (Micromasch,
149 HQ:XSC11/A1 BS) with a cantilever resonance frequency and force constant of 155 kHz
150 and 7 N/m were used in all the measurements.

151 Images were processed by using the Gwyddion v2.39 free software for scanning probe
152 microscopy data visualization and analysis (<http://gwyddion.net/>).

153

154 **2.4. Incorporation of Labeled Acetate**

155 A total of 0.5 μ Ci of [14 C] sterilized acetate sodium salt (43 mCi mmol $^{-1}$, New England
156 Nuclear) was added to 25 ml bacterial cultures at the time of inoculation. Cells grown under
157 the above conditions were harvested by centrifugation at 8,000 rpm for 10 min in a
158 Beckman Allegra 64R refrigerated centrifuge. Pellets were washed twice with 0.9 % NaCl
159 and used for subsequent studies.

160

161 **2.5. Preparation of Outer and Cytoplasmic Membranes**

162 Fractions of internal and external membrane were obtained according to the technique of
163 Mizuno et al., (1978) [21]. Cells control and PEG shocked were harvested by centrifugation
164 at 8,000 rpm for 10 min at room temperature and washed with saline buffer pH 7. Cells (ca.
165 1.5 g wet weight) were suspended in 18ml of ice-cold 20 % (w/v) sucrose and ice-cold
166 reagents were slowly added to the suspension in an ice-bath in the following order: 9 ml of
167 2 M sucrose, 10 ml of 0.1 M Tris-HCl (pH 7.8), 0.8 ml of 1 % Na-EDTA (pH 7.0), and 1.8

168 ml of 0.5 % lysozyme. Then the mixture was warmed to 30 °C within a few minutes and
169 kept at that temperature for 60 min. During the incubation, the mixture became viscous due
170 to breakage of a part of the cells. Then the suspension was centrifuged to remove the
171 spheroplasts at 13,000 rpm for 15 min at 30 °C. Crude outer membranes were recovered
172 from the supernatant by centrifugation at 30,000 rpm for 60 min. The spheroplasts were
173 burst in 40 ml of 5mM MgCl₂ and the spheroplast membranes were recovered by
174 centrifugation at 15,000 rpm for 20 min.

175 Two membrane markers, the KDO (2-ketodeoxyoctonate) content for the outer membrane
176 and the NADH oxidase activity for the inner membrane [22] were used. The KDO content
177 was 4.16 µg/100 µg protein and 0.63 µg/100 µg protein for the outer and inner membrane
178 respectively. NADH oxidase activity was 760 U/mg protein/min and 17.6 U/mg
179 protein/min for the inner and outer membrane respectively. The content of these markers
180 demonstrates efficient separation and purity of both membrane fractions.

181

182 **2.6. Lipid extraction**

183 Lipids were extracted from bacterial cells and outer and inner membrane and washed with
184 chloroform/methanol/water [23]. The lower phase containing lipids was dried under N₂ and
185 dissolved in an appropriate volume of 2:1 chloroform/methanol (v/v).

186

187 **2.7. Separation and Quantification of [1-14C]-labelled Phospholipids**

188 Thin layer chromatography (TLC) plates (silica gel HLF, 250 ml) were purchased from
189 Analtech. Aliquots of total lipid extracts were analyzed by TLC using a solution of
190 40:15:14:12:7 chloroform/acetone/methanol/acetic acid/water (v/v/v/v/v) as the solvent. All
191 solvents were of analytical or high-performance liquid chromatography grade. Lipids were

192 detected with iodine vapors, and the separated lipids were identified by comparing with
193 authentic purified standards purchased from Sigma. The TLC plates were scraped, and
194 fractions were quantified by performing radioactivity measurements using a liquid
195 scintillation counter (Beckman LS 60001 C).

196

197 **2.8. Analysis of Fatty Acids by GC-MS**

198 FA methyl esters (FAME) from SEMIA6144 cells were analyzed using a gas
199 chromatograph (GC) Agilent 7890B coupled to a Mass Spectrometer Agilent 5977A (MS)
200 equipped with a ZB-WAX (30 mm x 0.25 mm ID) Zebron column. The following GC-MS
201 conditions were used: injector temperature, 240 °C; column temperature, 180 °C,
202 maintained for 30 min; increase of 5 °C/min to 240 °C, maintained for 10 min. Run time:
203 46 min. MS: full SCAN, 40-500. Injection volume: 1 µl. Split: 1:10.

204 The peaks for each FA were identified using a mixture of commercial standards obtained
205 from Sigma-Aldrich Chemical Co. and Supelco [24]. FAME were identified by comparing
206 retention times to commercial standards (Sigma Chemical Co., St. Louis, MO, USA).

207 To determine the FA composition of every specific PL class, PL were separated as
208 described above, and each spot was scraped from plates and methyl-esterified. The
209 resulting FAME were analyzed by GC-MS as described above.

210

211 **2.9. Preparation of Phospholipid Vesicles**

212 Multilamellar vesicles (MLVs) were prepared according to Yamazaki [25] by adding 1 ml
213 of distilled water to dry SEMIA6144 lipids (total, outer or inner membrane) and then
214 heating the mixture to approximately 10 °C above the phase transition temperature for 20
215 min. The suspension was vortexed several times for approximately 5 min each.

216 2.10. Determination of Membrane Fluidity

217 Fluidity of cells and MLVs was determined by measuring the fluorescence polarization of
218 the 1,6-diphenyl-1,3,5-hexatriene (DPH) probe (Invitrogen) inserted into the membrane.
219 Fluorescence polarization quantifies the degree of depolarization of light emitted by the
220 embedded fluorescence probe and is a measure of membrane state [26]. The probe
221 polarization ratio and membrane fluidity are inversely correlated. As the fluidity of the
222 bacterial membrane decreases, the polarization ratio increases and vice versa [27].
223 Following procedures described by Trevors [28], SEMIA6144 cells were grown in B-
224 modified medium to exponential phase, harvested, washed in sterile 15 mmol l⁻¹ Tris-HCl
225 buffer (pH 7.0) and resuspended in the same buffer up to 0.2 OD at 600 nm. Then, 1 µl of
226 the fluorescent probe (stock solution diluted to 12 mmol l⁻¹ in tetrahydrofuran) was added
227 to 3 ml aliquots of resuspended cells to obtain a final probe concentration of 4 µmol l⁻¹.
228 Aliquots were incubated with a magnetic stirrer at 200 rpm for 10 min in the dark at room
229 temperature. The culture aliquots were then divided; one part was maintained as the control
230 and the other part was subjected to 15 mmol l⁻¹ PEG treatment for different periods of time
231 (10, 20, 30 and 60 min). Aliquots were collected, and the degree of polarization was
232 determined.

233 To measure the fluidity of the vesicles directly 1 µl of DPH was added to 3 ml of the
234 suspension of MLVs and the polarization was measured.

235 Fluorescence polarization measurements were performed using a FluoroMax®-4 Horiba
236 spectrofluorometer with a polarizer. The excitation wavelength for the DPH probe was 358
237 nm, and the emission wavelength was 428 nm. The slit widths for the excitation and
238 emission beams were 12 and 10 nm, respectively. The degree of polarization was calculated
239 from the polarization ratio (P) using the equation $P = (IVV - IVHG)/(IVV + IVHG)$, where

240 IVV and IVH are the intensities of vertically and horizontally polarized light components
241 emitted after excitation by vertically polarized light, respectively, and G is the sensitivity
242 factor of the detection system [29].

243

244 **2.11. Phospholipase Assay**

245 Total protein extracts were obtained from SEMIA6144 cultures (200 ml) grown to
246 exponential phase under control conditions or PEG shock for 1 h, 5 h and 24 h. Outer
247 membrane was extracted as explained above and used as enzyme source. Protein
248 concentrations were determined using the Bradford method [30]. To determine the PLA
249 activity in the extracts, PLA assays were performed at 30 °C. The standard reaction mixture
250 contained 50 mM Tris-HCl (pH 9), 10 mM CaCl₂, 450 μM Triton X-100, and [1-¹⁴C]-
251 labeled PE of *E. coli* DH5α as the substrate [16, 31]. Assays were initiated by adding 100
252 μg of protein (enzyme extract) to the reaction mixture and incubating for 60 min. The
253 reaction was stopped with the addition of 150 μl of methanol and 125 μl of chloroform.
254 Following centrifugation, the bottom layer was transferred to a glass tube and dried under
255 an N₂ stream. The residue was dissolved in 20 μl of chloroform/methanol 2:1 (v/v) and
256 subjected to TLC detection. PE and LPE fractions were scraped and quantified by
257 performing radioactivity measurements using a liquid scintillation counter. Finally, the
258 radioactivity present in the LPE fraction was used to calculate the enzymatic activity of
259 phospholipase A, which was expressed as percent of substrate hydrolyzed.

260

261 **2.12. Statistical Analyses**

262 All experiments were carried out in triplicate. One-way ANOVA was used to analyze the
263 results. When ANOVA indicated a significant treatment effect, the least significant
264 differences test (Tukey, $p < 0.05$) was applied to compare the mean values.

265 **3. Results**

266

267 **3.1. PEG Shock Negatively Affects the Viability of SEMIA6144 Cells**

268 The viability (measured as CFU ml⁻¹) of SEMIA6144 cells under water deficit conditions
269 was determined (Fig. 1). PEG shock from 48 h growth caused a decrease in bacterial
270 growth compared to control conditions, with bacterial counts reaching $7.3 \cdot 10^8$ CFU ml⁻¹
271 and $2.2 \cdot 10^{12}$ CFU ml⁻¹, respectively, after 110 h of growth.

272

273 **3.2. PEG Shock Modifies the Surface Morphology of the Cells of SEMIA6144**

274 AFM were used to determine the effect of PEG on cell surface morphology (Fig. 2). The
275 control cells were rod-shaped with a relatively rough surface (Fig. 2A). Perturbations on the
276 surface and membrane dehydration were observed 10 min after the addition of PEG (Fig.
277 2B). However, after 1 h of shock (Fig. 2C, D), we found that many of the cells had a similar
278 surface morphology to control cells, without the marked effects observed at 10 min. For
279 this reason, we decided to study the physical state of the membrane after PEG shock.

280

281 **3.3. PEG shock fluidizes the SEMIA6144 membrane, but the cells are able to restore 282 fluidity to optimal values**

283 The effects of PEG addition on the physical state of the membrane in SEMIA6144 cells
284 were evaluated by measuring the fluorescence polarization (P) of the probe DPH.

285 We found that the P value decreased from 0.20 ± 0.01 to 0.15 ± 0.02 immediately (10 min)
286 after the addition of PEG (Fig. 3). However, after 30 min of PEG exposure, the P value
287 returned to a value similar to that obtained in the absence of PEG. These results suggest
288 that PEG exerts a fluidizing effect on the SEMIA6144 membrane, and this effect reverts
289 after 30 min of PEG exposure, after reaching 1 h of shock values similar to the control.

290

291 **3.4. SEMIA6144 Cells Change the Composition of Totals Phospholipid and Fatty** 292 **acids**

293 Based on the results described above, the following question can be posed: how does the
294 SEMIA6144 membrane return to optimal fluidity values after undergoing the fluidizing
295 effects of PEG? To investigate this question, we studied the total PL and FA composition in
296 PEG shocked cells for different periods of time (1 h, 5 h and 24 h).

297 During exponential growth, the predominant labeled PL (Fig. 4) were PC (34.25 %), PE
298 (30.6 %) and PG (21.3 %), followed in descending order by CL (7.15 %) and LPE (6.7 %).
299 This implies that the PL forming the bilayer (PC and PG) comprised 55 % of the total PL,
300 while non-bilayer-forming PL (PE and LPE) comprised 37.3 % of the total. In addition, the
301 FA detected were 18:1 Δ 11 (78.1 %), 18:0 (5.2 %), 16:1 (0.5 %) and 16:0 (15.9 %) (Table
302 1).

303 The response to PEG shock for 1 h revealed an increase in PC (22 %) and a marked
304 increase in LPE (205 %), while PE decreased (76.6 %) with respect to control conditions
305 (Fig. 4). Unlike SEMIA6144 control cells, shocked cells exhibited membranes containing
306 63.8 % bilayer-forming PL and 29.3 % non-bilayer-forming PL. When SEMIA6144 was
307 exposed to PEG for 1 h, the percentage of FA 18:1 Δ 11 declined from 78.1 to 72.9 %, while
308 the percentage of FA 16:0 increased from 15.9 to 18.5 %. The UFA/SFA ratio decreased

309 from 3.7 to 2.9 with respect to control conditions due to a decrease in UFA (Table 1). These
310 changes in PL and FA composition may be responsible for the restoration of membrane
311 fluidity of SEMIA6144 after undergoing the fluidizing effects of PEG (Fig. 3).

312 To determine whether the response discussed above is constant or dynamic in time, we
313 evaluated the effects of prolonged exposure to PEG (5 h and 24 h) on total PL and FA
314 composition.

315 After 5 h of exposure to PEG, rearrangements in the levels of different PL were observed
316 with respect to the composition after 1 h of exposure. The amount of LPE decreased from
317 20.47 to 14.18 %, but the amount of PE was constant, unlike PC, whose percentage
318 increased from 41.8 to 48.8 % (Fig. 4). Interestingly, after 24 h of PEG shock, a novel
319 rearrangement in PL was observed compared to 5 h of shock. Under these conditions, the
320 amount of PE significantly increased from 8.5 to 23.7 %, the amount of PC decreased from
321 48.8 to 41 %, and the amount of LPE remained constant (Fig. 4). The total membrane
322 composition comprised 55 % bilayer-forming PL and 38 % non-bilayer-forming PL.

323 When the FA composition of SEMIA6144 cells exposed to PEG for 5 h and 24 h was
324 evaluated and compared to exposure for 1 h, FA 18:1 Δ 11 increased from 72.9 to 82 % and
325 from 72.9 to 84.6 %, respectively (Table 1). These results suggest that SEMIA6144 rapidly
326 responds to the presence of PEG, and this response is modified as the exposure time
327 increases.

328 Because SEMIA6144 is a Gram-negative bacterium and has a complex cell envelope with
329 two membranes (outer and inner), and due to the effect, that PEG exerts on the cell surface,
330 we decided to study the role of each of them in the response to PEG.

331

332

333

334

335 **3.5. Both Outer and Inner Membranes Modify Their Phospholipid Composition in**
336 **Response to PEG Shock**

337 The PL composition of both membranes (outer and inner) of SEMIA6144 cells was
338 evaluated (Fig. 5). Both membranes presented PC, PE, PG, CL and LPE as main PL,
339 however the outer membrane was more enriched in LPE than the inner membrane.

340 PEG shock for 1 h, caused an increase in LPE in both membranes, being more evident in
341 the outer membrane (165 %) compared to the control (Fig. 5A). In addition, the percentage
342 of PE decreased in both membranes (49 % in the outer membrane and 26.5 % in the inner
343 membrane). Regarding the behavior of PC was different in both membranes, increased in
344 inner membrane (9.3 %) and decreased in outer membrane (24 %) with respect to the
345 control. Compared to 1 h of shock, a decrease in LPE in both membranes (51 % in the outer
346 membrane and 34 % in the inner membrane) was observed at 5 h of shock, which resulted
347 in an increase of PE in both membranes (33 % in outer membrane and 13 % in inner
348 membrane), in this condition the percentage of PC remained constant in both membranes.
349 24 h of shock caused again changes in both membranes with respect to the previous
350 condition, obtaining a composition of PL similar to the control without PEG. For this PC,
351 increased external membrane (34 %) and decreased inner membrane (9 %) in which PE
352 increased (8 %). An increase in the anionic PL, PG and CL, was observed in outer
353 membrane, after 1 h (21 % and 8 % respectively) and 5 h (37 % and 31 % respectively)
354 shock. At 24 h the percentages of PG and CL decreased in outer membrane, reaching
355 values similar to control. In inner membrane, the percentages of PG and CL were not
356 modified during the shock.

357 Due to the complexity of the cellular envelopes of this type of bacteria, the use of model
358 systems is very useful to be able to study the behavior of the lipid components without the
359 interference of other components. For this reason, we studied the fluorescence polarization
360 of DPH in MLVs synthesized with lipids of different cell fractions.

361

362 **3.6. The MLVs synthesized with lipids from shocked cells 1 h have greater rigidity**

363

364 **MLVs From Total Lipids**

365 When the fluidity of the vesicles prepared with the total lipids of the SEMIA6144 cells was
366 analyzed (Fig. 6), greater P (less fluidity) was observed in the MLVs derived from lipids of
367 cells exposed to 1 h with PEG (0.20) as compared to control MLVs (0.14). MLVs prepared
368 with total lipids of cells exposed to 5 h and 24 h at PEG, showed lower P (0.17 and 0.12
369 respectively) than the MLVs prepared with total lipids from cells exposed 1 h to PEG. In
370 addition, MLVs prepared with total lipids of cells exposed to 24 h to PEG presented
371 practically the same polarization as the control MLVs.

372

373 **MLVs From Outer and Inner Membrane Lipids**

374 When the fluidity of the MLVs prepared with the outer membrane lipids of the
375 SEMIA6144 cells was analyzed (Fig. 6), higher P value (greater rigidity) was observed in
376 the MLVs derived from the lipids of cells exposed by 1 h to PEG (0.14) compared to
377 MLVs control (0.12). These results suggest that the PL composition of these vesicles would
378 contribute a greater rigidity to the outer membrane after 1 h of shock compared to the
379 control. However, the MLVs prepared with the outer membrane lipids of cells shocked for
380 5 h and 24 h showed fluidity similar to the control. The fluorescence polarization of DPH in

381 MLVs prepared with the internal membrane lipids of the SEMIA6144 showed a similar
382 behavior to that observed in MLVs from outer membrane lipids (Fig. 6), observing an
383 increase in rigidity in those vesicles prepared with the lipids of bacteria shocked for 1 h.

384

385 **3.7. LPE increase after 1 h of PEG shock involved an increase in PLA activity in the**
386 **outer membrane**

387 Our results suggested that the turnover between LPE and PE during PEG shock may
388 involve PLA activity and may be affected by PEG shock.

389 When outer membranes of SEMIA6144 grown under control conditions were used as
390 enzyme extract, PLA activity was observed, indicating that under optimum conditions of
391 growth SEMIA6144 produces lysophospholipids, as was demonstrated when the PL
392 composition of outer membrane was evaluated. Interestingly an increase in PLA activity
393 (70 %) was observed in outer membranes of SEMIA6144 shock for 1 h with PEG
394 compared to the control. Outer membranes from 24 h shock-cell presented a decrease in
395 enzyme activity (55 %) respect to 1 h of shock (Table 2).

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406 **4. Discussion**

407

408 Lipid bilayers are essential for the survival of microorganisms that are not isolated from
409 extrinsic physical and chemical factors [6, 32, 33, 34]. PEG dehydrates the lipid bilayer,
410 leading to formation of nonlamellar structures [35], raising the gel-to-fluid phase transition
411 temperature [36] and enhancing membrane permeability [37]. In addition, PEG creates
412 osmotic imbalance that leads to mechanical stress in membranes and may induce local
413 membrane curvature [38]. The visualization of SEMIA6144 cells through AFM showed
414 alteration of the surface morphology with areas of evident membrane dehydration by effect
415 to PEG shock. A fluidizing was observed 10 min after the addition of PEG, this effect was
416 maximal after 20 min of exposure and was then reversed until optimum fluidity was
417 achieved. The membrane fluidity of SEMIA6144 cells exposed to PEG for 1 h was
418 equivalent, indicating that the cells were able to alter this parameter to adjusting after the
419 immediate effects induced by PEG. In this condition, many of the cells presented a
420 superficial morphology similar to the control cells, indicating that they have strategies to
421 mitigate the effects of PEG.

422 The response of the SEMIA6144 membrane to quickly revert to optimal fluidity after
423 encountering the fluidizing effects of PEG involved changes in both PL and FA
424 composition.

425 When evaluating the total PL composition of the SEMIA6144 cells after 1 h of PEG shock,
426 we observed an increase in the amount of bilayer-forming PL (PC) and a reduction in non-
427 bilayer-forming PL (PE and LPE) compared to control. Among the non-bilayer-forming
428 PL, a surprising increase in LPE was observed as well as a decrease in its precursor (PE).

429 The LPE content in bacterial membranes usually does not exceed 2-5 %; however, various
430 stressors generate a dramatic increase in LPE content [39]. In *Yersinia pseudotuberculosis*
431 exposed to phenol, increased LPE caused an increase in the phase transition temperature of
432 the membrane, resulting in greater membrane rigidity [39]. In SEMIA6144 shocked cells
433 for 1 h, LPE increased at the expense of PE, which appeared to be a rapid response since
434 we find that the total lipid MVLS of exhibited higher rigidity compared to control,
435 indicating that the turnover between LPE and PE contributes to rigidify the membranes.
436 Besides an increase in PC after shock PEG for 1 h may also help counteract the fluidizing
437 effects, stabilizing the membrane. Furthermore, this rapid response of SEMIA6144
438 involved a change in the composition of FA with a decrease in the FA unsaturation index,
439 which may promote increased membrane rigidity in response to the fluidizing effects of
440 PEG.

441 When SEMIA6144 was shocked with PEG for longer periods of time (5 h and 24 h), LPE
442 was not maintained at a high percentage in the membrane. LPE is an intermediate in the
443 metabolism of PL [40] derived from the PLA activity on PE [41] and is quickly recycled
444 and likely metabolized due to its ability to create membrane instability [42, 43, 39].

445 Although the amount of LPE decreases, the rearrangements that occur in the membrane
446 after SEMIA6144 is exposed to PEG for longer periods of shock allow optimal fluidity to
447 be maintained ($P: 0.23$), as well as the balance between bilayer-forming and non-bilayer-
448 forming PL.

449 The envelope of Gram-negative bacteria is composed of two distinct lipid membranes: an
450 inner membrane and outer membrane. Both, the outer and the inner membrane of
451 SEMIA6144 presented PC and PE as main PL. The most important difference was the high

452 content of LPE in the outer membrane, in addition to a greater proportion of anionic PL, PG
453 and CL.

454 The detailed analysis of the PL composition of each SEMIA6144 membrane under PEG
455 shock, indicated that 1 h of shock caused a large increase of LPE in the outer membrane
456 and concomitant decrease of PC and PE. This increase in LPE was related to an increase in
457 the outer membrane PLA activity on PE. Nevertheless, after 24 h of PEG shock, the outer
458 membrane was composed of large amounts of PC and PE and less amount of LPE, due to a
459 decrease in PLA activity. The inner membrane composition was also modified after 1 h of
460 shock, observing an increase of PC at the expense of PE. The proportions of these PL were
461 then modified to reach 24 h of shock values similar to the control.

462 The role of lysophospholipids in bacteria remains poorly studied, it has been proposed that
463 the geometric form of inverted cone that these shows could alleviate the curvature stress
464 caused on the membrane in certain situations [44]. LPE was also found to become
465 dominant (12 %) in the outer membrane of *Y. pseudotuberculosis* after shifting of the
466 growth temperature from 8 °C to 37 °C or after exposure of the cells to heat shock [45], it is
467 unclear why heat-stressed cells are able to retain LPE at such a high level. In this paper, we
468 demonstrate that in SEMIA6144 cells after PEG shock, the amount of LPE does not remain
469 high for a long time, but other phospholipids are synthesized (PC and PE).

470 All the MLVs synthesized with lipids of cells exposed 1 h to shock presented a greater
471 polarization, which could indicate that the increase of LPE and PC at the expense of PE
472 could be the response that allows the cells to recover from the fluidizing effect of PEG.
473 Although the use of model systems such as MLVs are very important to predict the
474 behavior of the lipid component of cell membranes, we must not forget that they are much

475 more complex and other components such as proteins and lipopolysaccharides could also
476 be involved in the bacterial response to water deficit.

477 Finally, we conclude that SEMIA6144 is able to tolerate moderate water deficiency by a
478 rapid response in the outer membrane, product of an increase in PLA activity, which
479 generates LPE from PE. This response is not maintained over time, but the cells modify
480 again the composition of both membranes to maintain homeostasis and ensure the integrity
481 of the cellular envelope under this condition. The findings reported in this study are
482 significant because they describe for the first time the importance of phospholipases as key
483 enzymes in PL turnover for rapid adaptation to stress in this bacterial genus.

484

485 **Conflict of interest**

486 The authors do not have any possible conflicts of interest.

487 **Acknowledgment**

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612 membranes of *Yersinia pseudotuberculosis*. *Biochimie* 2016;123:103-109.

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619 **Figure legends**

620

621 **Fig. 1.** Effects of PEG shock on the viability of *B. sp.* SEMIA6144.

622 Viability is expressed as LOG CFU.ml⁻¹. Values represent the mean \pm SEM of three
623 independent experiments. (■) Control without PEG; (●) control cells grown for 48 h and
624 then shocked with PEG.

625

626 **Fig.2.** Topography AFM images of *B. sp.* SEMIA6144. Three-dimensional images of
627 control (A) and exposed to PEG by 10 min (B) and 1 h (C). Two-dimensional image of the
628 population of *B. sp.* SEMIA6144 cells exposed for 1 h to PEG (D).

629

630 **Fig. 3.** Fluorescence polarization of DPH in *B. sp.* SEMIA6144 cells after PEG shock for
631 different amounts of time.

632 Cells cultivated in B-modified medium were harvested at the exponential phase (48 h),
633 washed and resuspended in Tris-HCl buffer to an OD 660 nm of 0.2. The fluorescent
634 membrane probe was added to the resuspended culture and incubated to facilitate
635 incorporation of the probe. Fluorescence polarization was measured in one aliquot as the
636 control sample (absence of PEG). The remainder of the culture was treated with PEG.
637 Aliquots were collected at different times (10, 20, 30 and 60 min) to determine the degree
638 of polarization. Values represent the mean \pm SEM of three independent experiments.

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643 **Fig. 4.** Effects of PEG shock on the composition of total phospholipids of *B. sp.*
644 SEMIA6144.

645 Cells grown at exponential phase with [$1\text{-}^{14}\text{C}$] acetate sodium salt (37KBq ml^{-1}) followed
646 by exposure to PEG for 1 h, 5 h and 24 h were harvested and the total phospholipids
647 analyzed by TLC.

648 Lysophosphatidylethanolamine (LPE), phosphatidylcholine (PC), phosphatidylglycerol
649 (PG), phosphatidylethanolamine (PE), cardiolipin (CL). Values represent the mean \pm SEM
650 of three independent experiments. * indicates a statistically significant difference ($p < 0.05$)
651 for each condition studied with respect to control conditions.

652

653 **Fig. 5.** Effects of PEG shock on the composition of phospholipids of outer (a) and inner (b)
654 *B. sp.* SEMIA6144 membrane.

655 Cells grown at exponential phase with [$1\text{-}^{14}\text{C}$] acetate sodium salt (37KBq ml^{-1}) followed
656 by exposure to PEG for 1 h, 5 h and 24 h were harvested and the outer and inner membrane
657 fraction were obtained. The phospholipids were analyzed by TLC.

658 Lysophosphatidylethanolamine (LPE), phosphatidylcholine (PC), phosphatidylglycerol
659 (PG), phosphatidylethanolamine (PE), cardiolipin (CL). Values represent the mean \pm SEM
660 of three independent experiments. * indicates a statistically significant difference ($p < 0.05$)
661 for each condition studied with respect to control conditions.

662

663 **Fig. 6.** Fluorescence polarization of DPH in MLVs prepared with lipids of *B. sp.*
664 SEMIA6144.

665 Cells grown at exponential followed by exposure to PEG for 1 h, 5 h and 24 h were
666 harvested and the outer and inner membrane fraction were obtained. Phospholipids of each

667 fraction were used to prepare MLVs as indicated in the materials and methods section. The
668 fluorescent membrane probe (DPH) was added to the suspension of vesicles and incubated
669 to facilitate incorporation of the probe. Fluorescence polarization was measured. Values
670 represent the mean \pm SEM of three independent experiments.

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692 **Table 1.** Effects of PEG shock on the fatty acid composition of *B. sp.* SEMIA6144.

Fatty acid type (%)	Exponential Phase (48 h)			
	Control	PEG shock		
		1 h	5 h	24 h
Saturated				
16:0	15.9 ± 1.3	18.2 ± 1.1*	14.7 ± 1.1	13.7 ± 1*
18:0	5.22 ± 1.7	7.2 ± 1.4*	2.4 ± 0.9*	1.2 ± 0.8*
Unsaturated				
16:1Δ9	0.53 ± 0.2	0.5 ± 0.1	0.6 ± 0.1	0.8 ± 0.1*
18:1Δ11	78.1 ± 3	72.9 ± 3*	82 ± 3.2	84.6 ± 3.1*
UFA/SFA†	3.7	2.9	4.8	5.7

693 To study the effects of growth under PEG on FA composition, *B. sp.* SEMIA6144 cells
 694 were grown until the exponential phase was reached and then were exposed to PEG for 1 h,
 695 5 h and 24 h. Cells were harvested, and FAME were obtained from total lipids and analyzed
 696 by GC-MS. The percentage of each FA is relative to the total FA defined as 100 %.

697 † Ratio between the sums of unsaturated and saturated FA (UFA/SFA). Values represent
 698 the mean ± SEM of three independent experiments. * indicates a statistically significant
 699 difference ($P < 0.05$).

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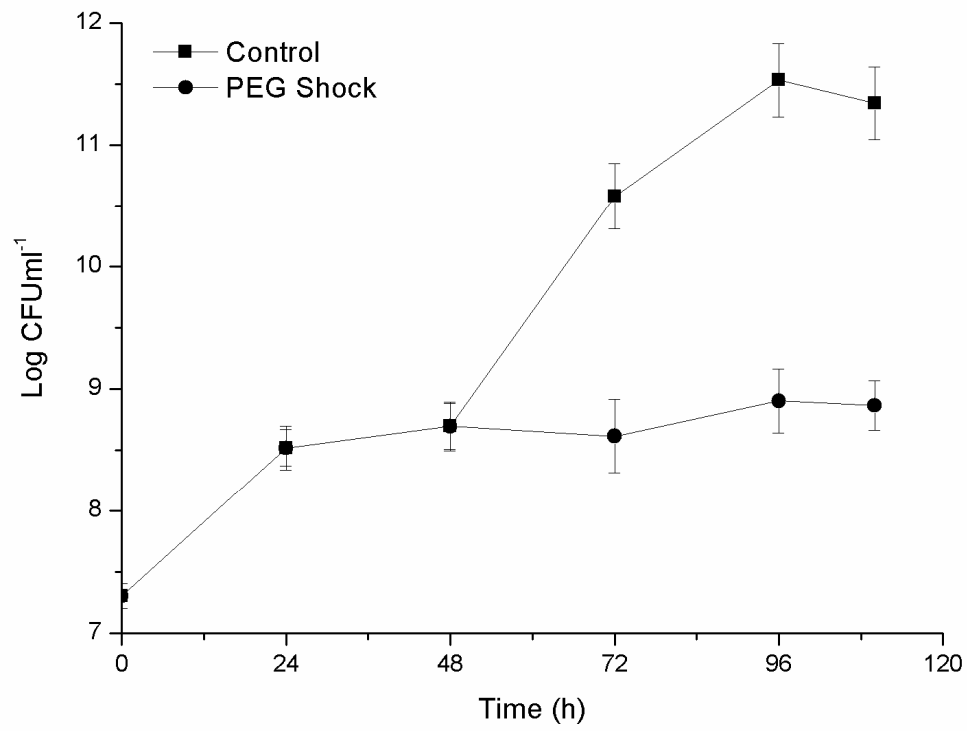
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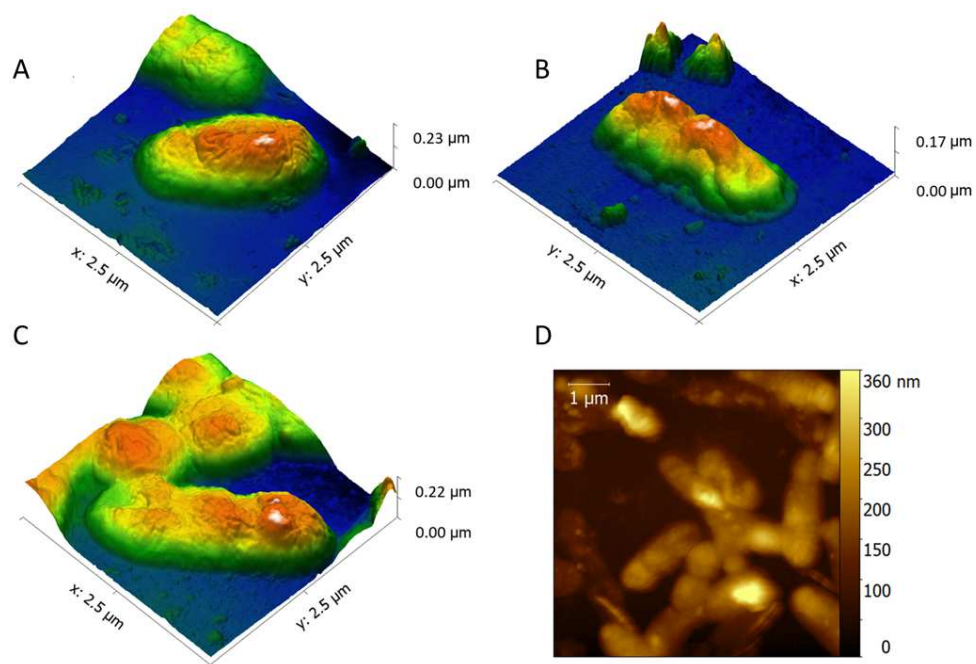
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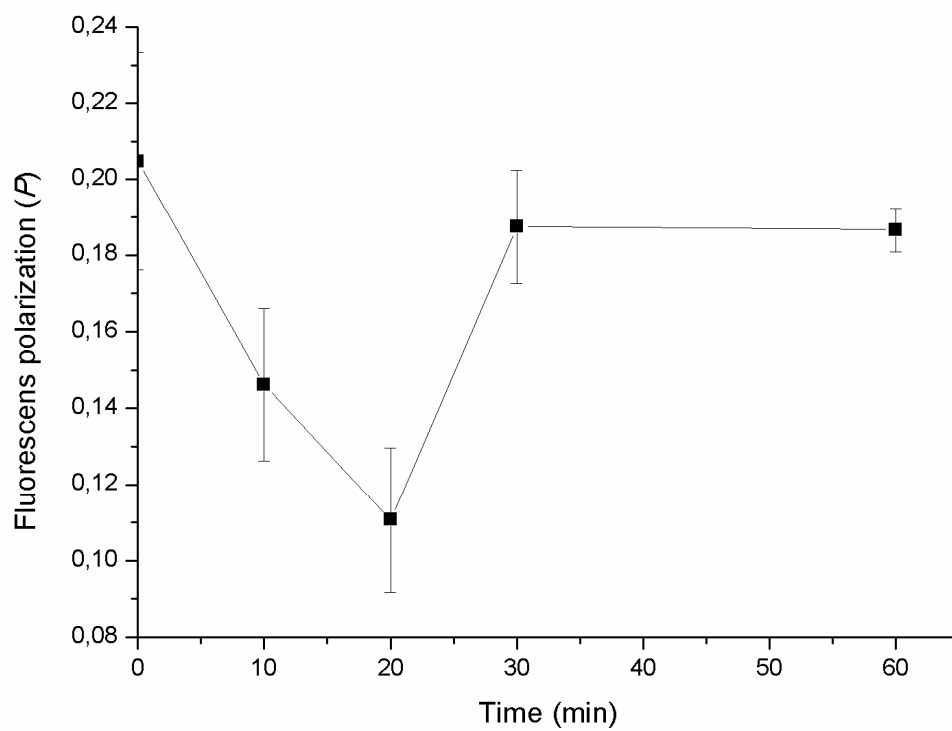
706 **Table 2.** Phospholipase A activity of *B. sp.* SEMIA6144 outer membrane of cells exposed
707 to PEG.

Growth conditions	PLA activity (% of substrate hydrolyzed)
Control	7.2 ± 0.7
1 h PEG	12.5 ± 0.5
5 h PEG	11.9 ± 0.9
24 h PEG	5.7 ± 0.5

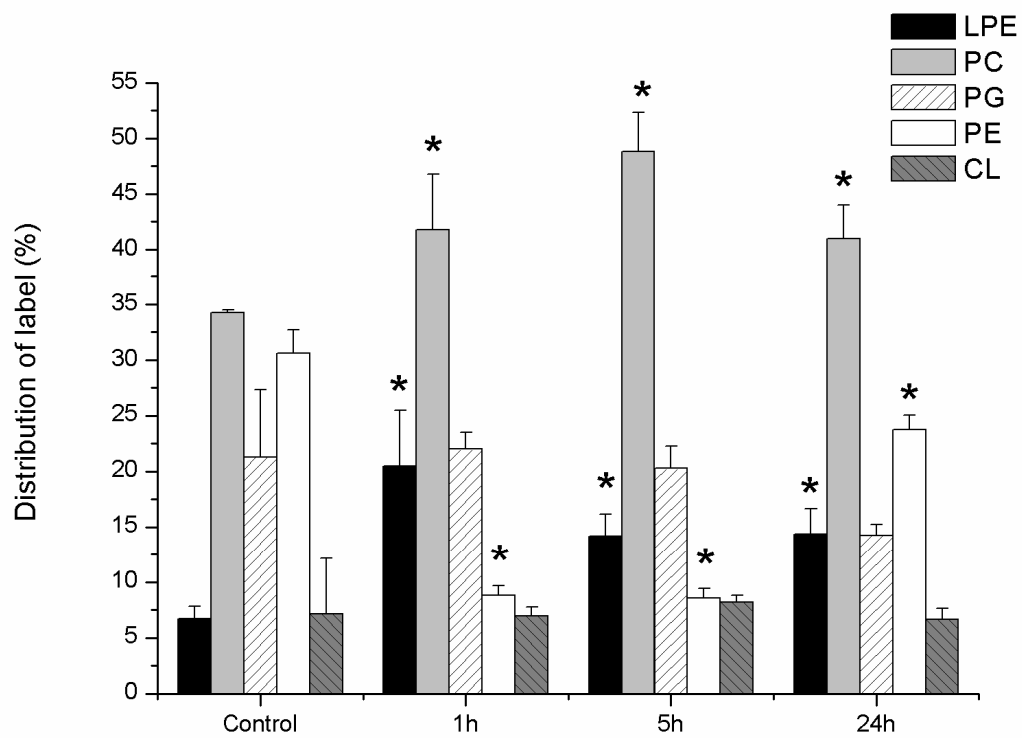
708 Cells were grown until the exponential phase was reached and then were exposed to PEG
709 for 1 h, 5 h and 24 h; cells were then harvested, and protein fraction of the outer membrane
710 was used as enzymatic extracts. [1-¹⁴C]-labeled PE of *E. coli* DH5α was used as substrate
711 and after expiry of the reaction time, the fraction of PE was scraped and the fraction
712 quantified by radioactivity measurements, activity is expressed as percent of substrate
713 hydrolyzed. Values represent mean ± SEM of two independent experiments.

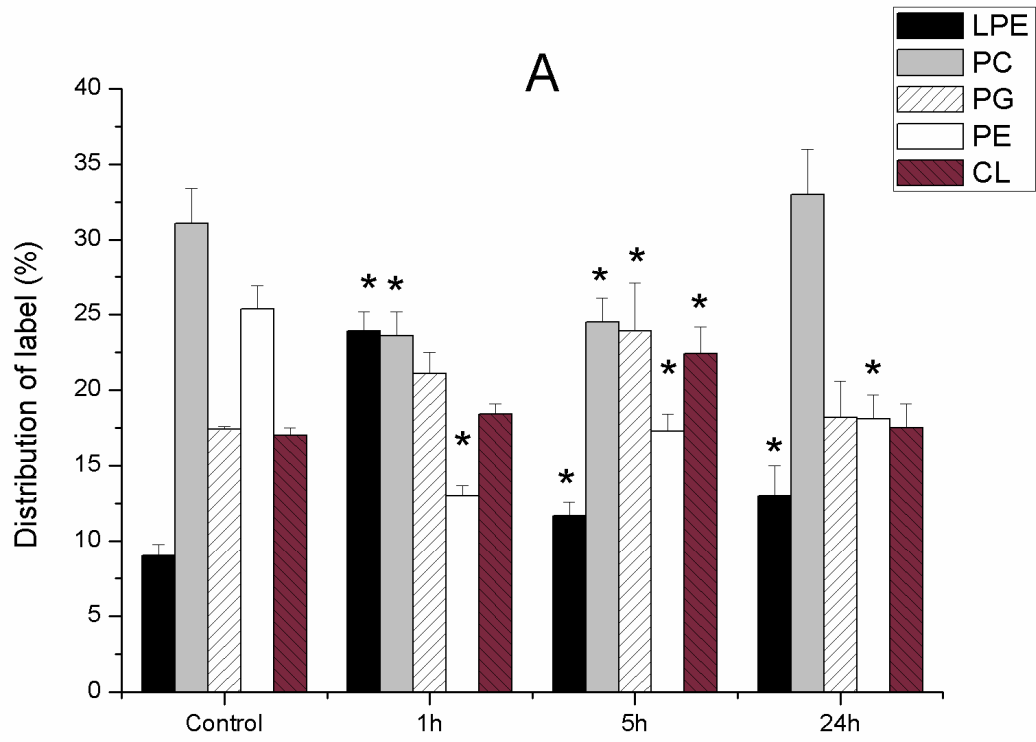




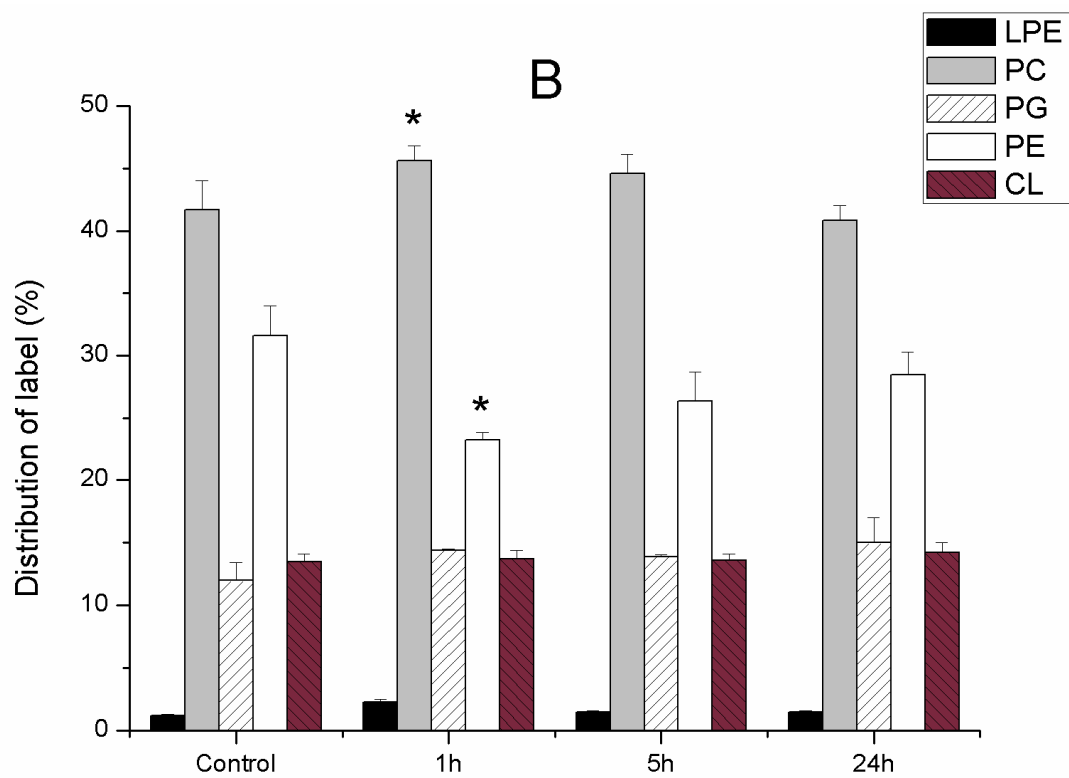


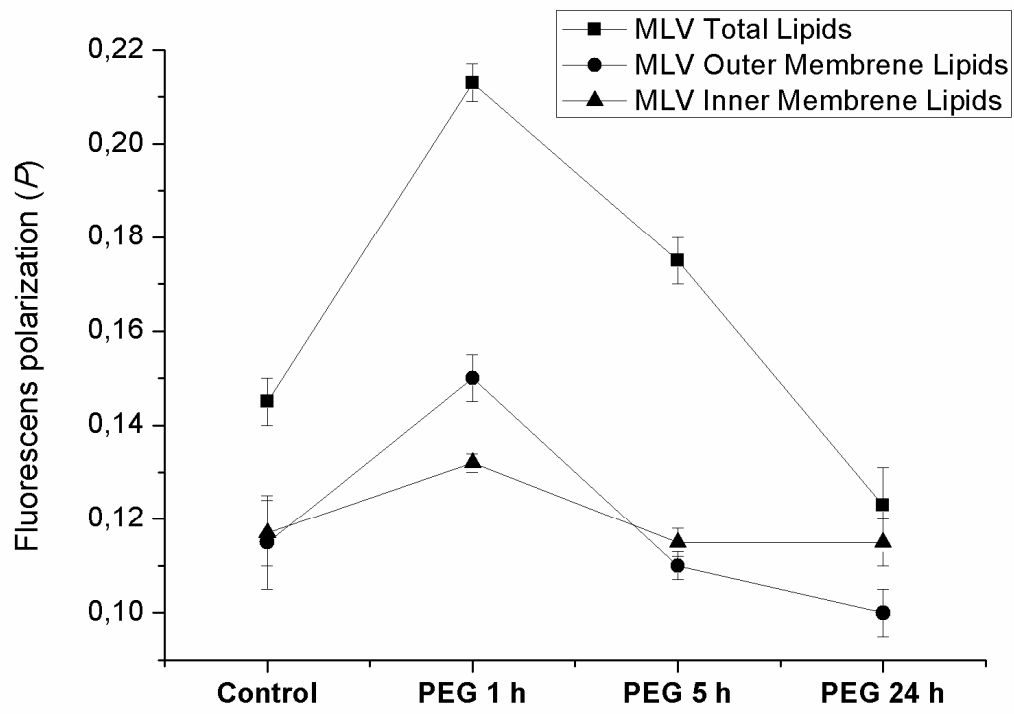
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