

Lipidomics Analysis of Multilamellar Bodies Produced by *Amoeba Acanthamoeba castellanii* in Co-Culture with *Klebsiella aerogenes*

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Supplementary Materials

1. The MLBs production inside *Acanthamoeba castellanii*

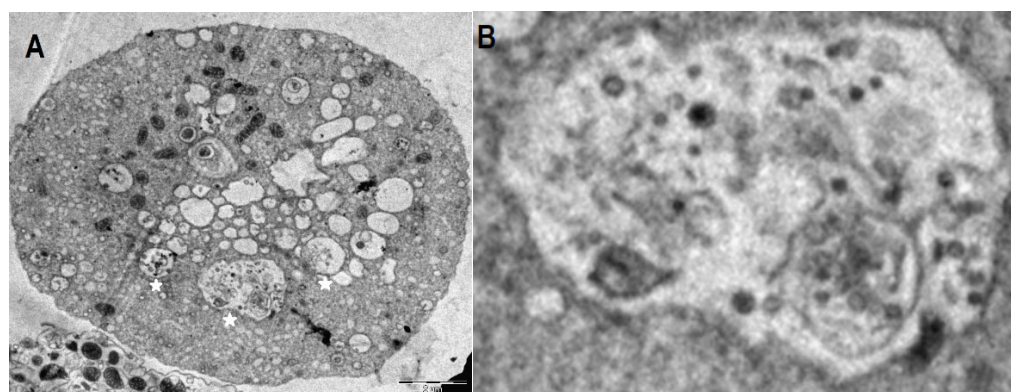


Figure S1. Representative TEM image of *A. castellanii* trophozoite cultured axenically for 48 hours on NNA agar plates. A. and B. magnified image of an autophagic-like vacuole; The white stars denote autophagic-like vesicles containing completely or partially digested mitochondria or cytosolic compartments.

2. Lipidomic analysis of MLBs derived of *Acanthamoeba castellanii*

Table S1. Classes and content (given in mol %) of ester bound fatty acids residues identified in lipids extracted from MLBs using the Bligh and Dyer protocol.

Fatty acid	mol %
14:0	1.055 ± 0.18
15:0	tr
16:1 ^{Δ7}	1.87 ± 0.72
16:0	21.43 ± 3.7
17:1*	3.25 ± 0.7
17:0	tr
18:3*	tr
18:2 ^{Δ9,12}	1.36 ± 0.47
18:1^{Δ9}	38.46 ± 0.87
18:1*	4.4 ± 1.8
18:0	20.75 ± 2.08
20:4 ^{Δ5,8,11,14}	2.41 ± 0.56
20:3 ^{Δ8,11,14}	1.15 ± 0.11

20:2 ^{Δ11,14}	1.4 ± 0.07
20:0	tr

Data were determined by GC-MS analysis of the FAs methyl esters subsequent to saponification (0.8 M NaOH/50 % MeOH; 1 h/ 80°C). The results are presented as ± S.D. of three independent measurements. *

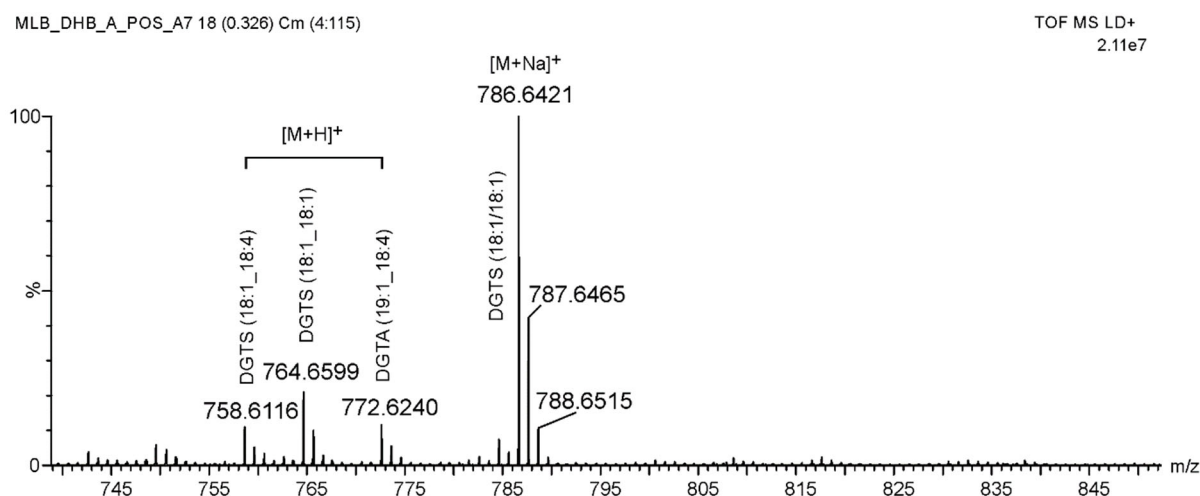


Figure S2. Spectrometric analysis of betaine lipids from MLBs of *A. castellanii*. Ions registered as $[M + H]^+$ and $[M + Na]^+$ adducts by MALDI-TOF in m/z range 700 - 850.

3. Lipids derived of whole cells *Klebsiella aerogenes*

3.1. HPTLC analysis

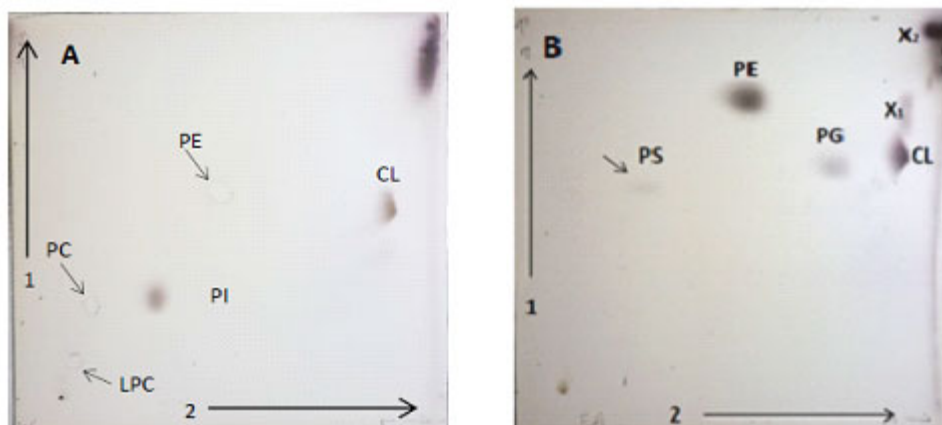


Figure S3. 2-D HPTLC chromatograms of **A)** commercial standards **B)** phospholipids extracted from whole cells of *Klebsiella aerogenes*. Abbreviations: PC - phosphatidylcholine, PE - phosphatidylethanolamine, CL - cardiolipin, PS - phosphatidylserine, LPC - lysophosphatidylcholine, PI - phosphatidylinositol; Used solvents: 1 – first direction - chloroform/methanol/water (14:6:1, v/v/v), 2- second direction chloroform/methanol/acetic acid (13:5:2) [15].

3.2. Fatty acids identified in total lipids of *K. aerogenes*

Among fatty acids liberated by saponification (conditions as for data in Table 1S) from total lipids of *K. aerogenes* and derivatized to methyl esters identified by GC-MS: C16:0 (41.97 ± 0.6 mol%), *cyc*17:0 (29.53 ± 0.19 mol%), and *cyc*19:0 (14.92 ± 1.31 mol%), C18:1⁴⁹ (5.14 ± 0.09 mol%), C14:0 (3.9 ± 0.55 mol%). In trace amount (< 2.0 %) also registered: C15:0, C16:1, C17:0, C18:0. The results are presented as ± S.D. of three independent measurements.

3.3. DI-ESI-MS and MALDI-TOF MS² analysis of lipids *K. aerogenes*

Table S2. Lipids identified in whole cells *K. pneumoniae* by DI-ESI-MS. For some molecular species fatty acids were established by MALDI-TOF MS² fragmentation. Abbreviations for PLs are given in Figure 3S. *c* - cyclopropane ring structure.

Lipid (acyl carbons: double bonds)	Adduct	Formula	Observed <i>m/z</i>	Calculated <i>m/z</i>
PE (30:0)	[M + H] ⁺	C ₃₅ H ₇₁ NO ₈ P	664.4926	664.4917
	[M + Na] ⁺	C ₃₅ H ₇₀ NO ₈ PNa	686.4781	686.4737
	[M - H] ⁻	C ₃₅ H ₆₉ NO ₈ P	662.4811	662.4761
PE (31:1)	[M + H] ⁺	C ₃₆ H ₇₁ NO ₈ P	676.4960	676.4917
	[M + Na] ⁺	C ₃₅ H ₇₀ NO ₈ PNa	698.4815	698.4737
	[M - H] ⁻	C ₃₆ H ₆₉ NO ₈ P	674.4771	674.4761
PE (32:1)	[M + H] ⁺	C ₃₇ H ₇₃ NO ₈ P	690.5097	690.5074
	[M + Na] ⁺	C ₃₇ H ₇₂ NO ₈ PNa	712.4986	712.4893
	[M - H] ⁻	C ₃₇ H ₇₁ NO ₈ P	688.4937	688.4917
PE (32:0)	[M + H] ⁺	C ₃₇ H ₇₅ NO ₈ P	692.5300	692.5230
	[M + Na] ⁺	C ₃₇ H ₇₄ NO ₈ PNa	714.5040	714.5050
	[M - H] ⁻	C ₃₇ H ₇₃ NO ₈ P	690.5127	690.5074
PE (16:0/<i>c</i>17:0)	[M + H] ⁺	C ₃₈ H ₇₅ NO ₈ P	704.5281	704.5230
	[M + Na] ⁺	C ₃₈ H ₇₄ NO ₈ PNa	726.5137	726.5050
	[M - H] ⁻	C ₃₈ H ₇₃ NO ₈ P	702.5155	702.5074
PE (34:1)	[M + H] ⁺	C ₃₉ H ₇₇ NO ₈ P	718.5469	718.5387
	[M + Na] ⁺	C ₃₉ H ₇₆ NO ₈ PNa	740.5228	740.5206
	[M - H] ⁻	C ₃₉ H ₇₅ NO ₈ P	716.5226	716.5230
PE (35:1)	[M + H] ⁺	C ₄₀ H ₇₉ NO ₈ P	732.5620	732.5543
	[M + Na] ⁺	C ₄₀ H ₇₈ NO ₈ PNa	754.5458	754.5363
	[M - H] ⁻	C ₄₀ H ₇₇ NO ₈ P	730.5462	730.5387
PE (<i>c</i>17:0_<i>c</i>19:0)	[M + H] ⁺	C ₄₁ H ₇₉ NO ₈ P	744.5551	744.5538
	[M + Na] ⁺	C ₄₁ H ₇₈ NO ₈ PNa	766.5355	766.5397
PG (<i>c</i>17:0_16:0) PG (<i>c</i>19:0_14:0)	[M - H] ⁻	C ₃₉ H ₇₇ NO ₁₀ P	733.5012	733.5020
PG (35:1)	[M - H] ⁻	C ₄₁ H ₇₈ NO ₁₀ P	788.5470	788.5442
PS (36:1)	[M - H] ⁻	C ₄₂ H ₇₉ NO ₁₀ P	788.5470	788.5442

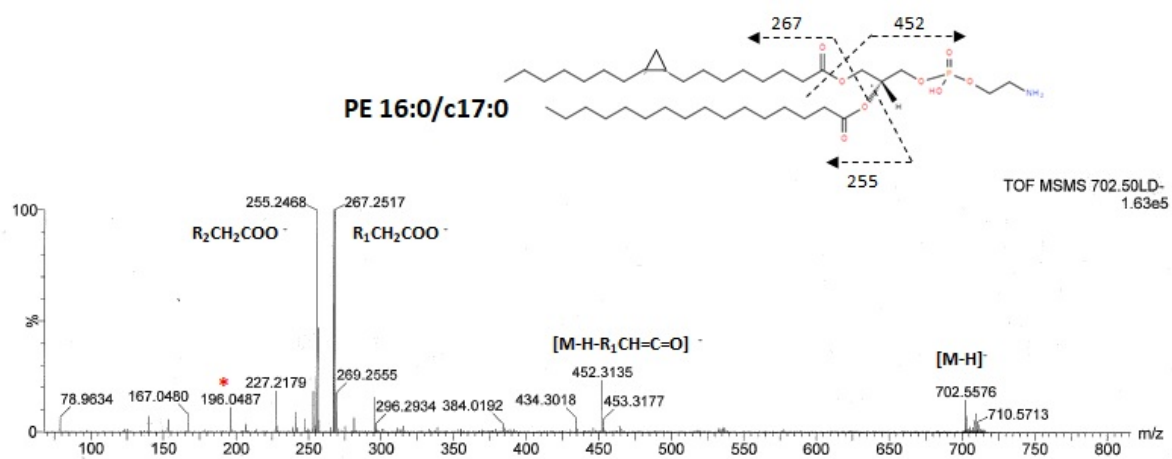


Figure S4. The MS² mass spectrum and fragmentation pattern for a selected ion corresponding to phospholipid PE (16:0/c17:0) $[M - H]^-$ obtained from total lipid extract from whole cells of *K. aerogenes*.