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Shiga Toxin Binding Alters Lipid Packing and Domain Structure of Gb₃-Containing Membranes: A Solid-State NMR Study

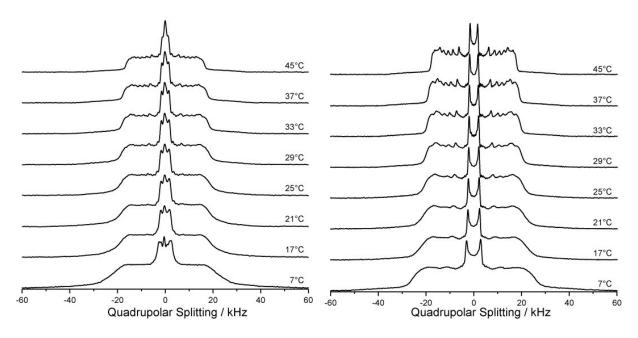
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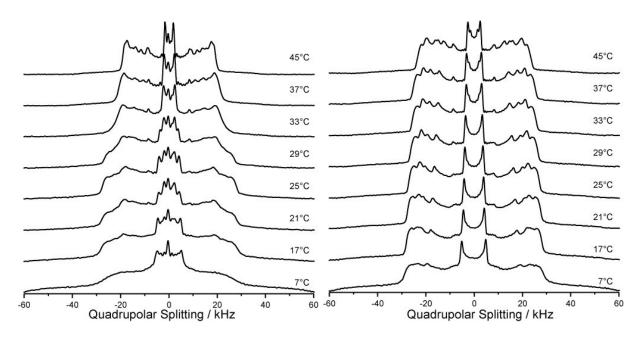
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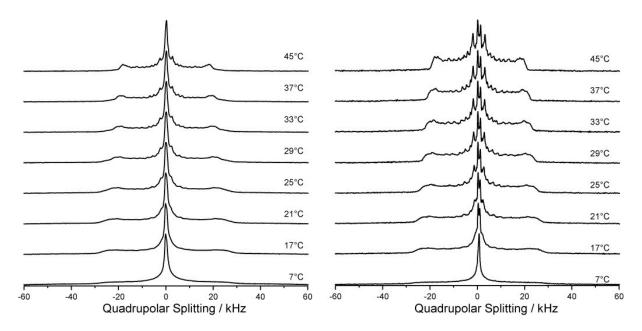
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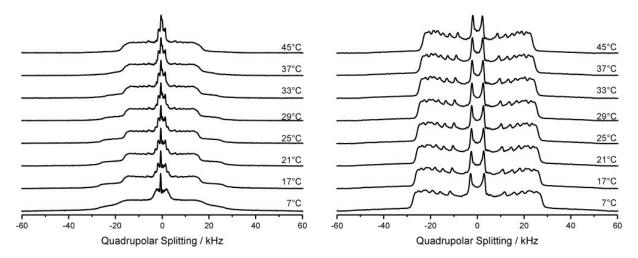
Supplementary Figure S1. ²H NMR spectra for the deuterated POPC- d_{31} of the quaternary lipid mixture lignoceryl-Gb₃/POPC- d_{31} /PSM/cholesterol (molar ratio: 5:40:35:20) in the presence of STxB (molar protein to lipid ratio of 1:120, left) and in the absence of STxB (right) at the given temperatures.



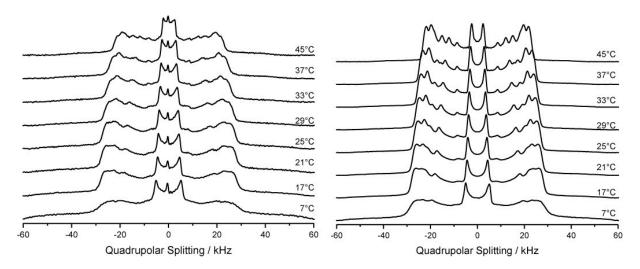
Supplementary Figure S2. ²H NMR spectra for the deuterated PSM- d_{31} of the quaternary lipid mixture lignoceryl-Gb₃/POPC/PSM- d_{31} /cholesterol (molar ratio: 5:40:35:20) in the presence of STxB (molar protein to lipid ratio of 1:120, left) and in the absence of STxB (right) at the given temperatures.



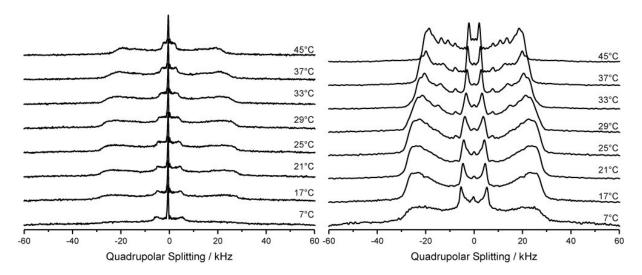
Supplementary Figure S3. ²H NMR spectra for the deuterated lignoceryl-Gb₃- d_{47} of the quaternary lipid mixture lignoceryl-Gb₃- d_{47} /POPC/PSM /cholesterol (molar ratio: 5:40:35:20) in the presence of STxB (molar protein to lipid ratio of 1:120, left) and in the absence of STxB (right) at the given temperatures.



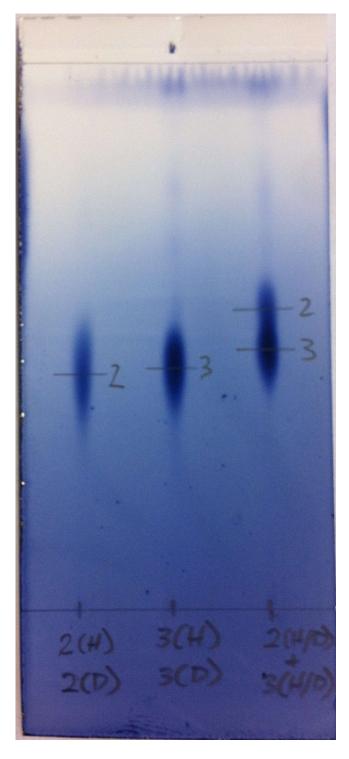
Supplementary Figure S4. ²H NMR spectra for the deuterated POPC- d_{31} of the quaternary lipid mixture palmitoyl-Gb₃/POPC- d_{31} /PSM/cholesterol (molar ratio: 5:40:35:20) in the presence of STxB (molar protein to lipid ratio of 1:120, left) and in the absence of STxB (right) at the given temperatures.



Supplementary Figure S5. ²H NMR spectra for the deuterated PSM- d_{31} of the quaternary lipid mixture palmitoyl-Gb₃/POPC/PSM- d_{31} /cholesterol (molar ratio: 5:40:35:20) in the presence of STxB (molar protein to lipid ratio of 1:120, left) and in the absence of STxB (right) at the given temperatures.



Supplementary Figure S6. ²H NMR spectra for the deuterated palmitoyl-Gb₃- d_{31} of the quaternary lipid mixture palmitoyl-Gb₃- d_{31} /POPC/PSM/cholesterol (molar ratio: 5:40:35:20) in the presence of STxB (molar protein to lipid ratio of 1:120, left) and in the absence of STxB (right) at the given temperatures.



Supplementary Figure S7. Thin layer chromatogram of all Gb_3 lipids used in this study. The plate was developed using H_2O , CH_3OH and CH_2Cl_2 (1, 30, 100, v/v/v). After the end of the chromatographic run and careful drying the different compounds were rendered visible by spraying with cerium (IV)/ammonium molybdate solution.

Supplementary Table S1: Calculated chain extent L_c^* in Å for the respective deuterated lipid in lignoceryl-Gb₃ /POPC/PSM /cholesterol (5:40:35:20 mol/mol/mol/mol) membranes. The experimental error of the L_c^* calculation is ~0.2 Å for Gb₃ and ~0.1 for POPC or PSM.

Temperature	Gb ₃ -d ₄₇		POPC-d ₃₁		PSM- <i>d</i> ₃₁	
/°C		+Shiga		+Shiga		+Shiga
25	21.0	21.0	14.1	13.5	15.8	15.4
37	19.5	18.9	13.6	13.1	15.4	14.6
45	18.9	18.5	13.3	13.0	14.9	14.1

Supplementary Table S2: Mean order parameter for the respective deuterated lipid in lignoceryl- Gb_3 - d_{47} /POPC/PSM /cholesterol (5:40:35:20 mol/mol/mol/mol) membranes. The typical experimental error on the average order parameter is <0.01.

Temperature	Gb ₃ -d ₄₇		POPC-d ₃₁		PSM- <i>d</i> ₃₁	
/°C		+Shiga		+Shiga		+Shiga
25	0.26	0.27	0.26	0.23	0.35	0.33
37	0.21	0.21	0.23	0.21	0.33	0.28
45	0.20	0.19	0.22	0.21	0.30	0.26

Supplementary Table S3: Calculated chain extent L_c^* in Å for the respective deuterated lipid in palmitoyl-Gb₃/POPC/PSM/cholesterol (5:40:35:20 mol/mol/mol/mol) membranes. The experimental error of the L_c^* calculation is ~0.2 Å for Gb₃ and ~0.1 for POPC or PSM.

Temperature	Gb ₃ -d ₃₁		POPC-d ₃₁		PSM- <i>d</i> ₃₁	
/°C		+Shiga		+Shiga		+Shiga
25	16.4	15.6	15.4	12.8	16.2	16.3
37	15.3	15.1	15.0	12.9	15.6	15.6
45	14.6	15.0	14.9	12.9	15.2	14.9

Supplementary Table S4: Mean order parameter for the respective deuterated lipid in palmitoyl-Gb₃/POPC/PSM/cholesterol (5:40:35:20 mol/mol/mol/mol) membranes. The typical experimental error on the average order parameter is <0.01.

Temperature	Gb ₃ -d ₃₁		POPC-d ₃₁		PSM- <i>d</i> ₃₁	
/°C		+Shiga		+Shiga		+Shiga
25	0.39	0.35	0.33	0.20	0.38	0.39
37	0.32	0.31	0.31	0.20	0.34	0.34
45	0.28	0.29	0.30	0.20	0.31	0.30