

Supplementary Information

Table S1. High resolution mass spectrometry (HRMS) analysis of the Antarctic microcystins and mass deviations from the proposed structures.

Microcystin	Measured <i>m/z</i> ^a	Molecular Formula	Calculated <i>m/z</i>	Error (ppm)
[Gly ¹ , Asp ³ , Dhb ⁷] MC-LR (3)	967.5285	C ₄₇ H ₇₁ N ₁₀ O ₁₂	967.5247	+3.86
[Gly ¹ , Asp ³ , Dhb ⁷] MC-LHar (4)	981.5413	C ₄₈ H ₇₃ N ₁₀ O ₁₂	967.5404	+0.87
[Gly ¹ , Asp ³ , ADMAdda ⁵ , Dhb ⁷] MC-LR (5)	995.5206	C ₄₈ H ₇₁ N ₁₀ O ₁₃	995.5197	-0.94
[Gly ¹ , Asp ³ , ADMAdda ⁵ , Dhb ⁷] MC-LHar (6)	1009.5348	C ₄₉ H ₇₃ N ₁₀ O ₁₃	1,009.5353	+0.47
[Gly ¹ , Asp ³ , Dhb ⁷] MC-RR (7)	1010.5442	C ₄₇ H ₇₂ N ₁₃ O ₁₂	1,010.5418	-2.44
[Gly ¹ , Asp ³ , Dhb ⁷] MC-RHar (8)	1024.5569	C ₄₈ H ₇₄ N ₁₃ O ₁₂	1,024.5574	+0.53
[Gly ¹ , Asp ³ , ADMAdda ⁵ , Dhb ⁷] MC-RR (9)	1038.5405	C ₄₈ H ₇₂ N ₁₃ O ₁₃	1,038.5367	-3.82
[Gly ¹ , Asp ³ , ADMAdda ⁵ , Dhb ⁷] MC-RHar (10)	1052.5541	C ₄₉ H ₇₄ N ₁₃ O ₁₃	1,052.5524	-1.60

^a All HRMS measurements were made using the singularly-protonated molecular ion.

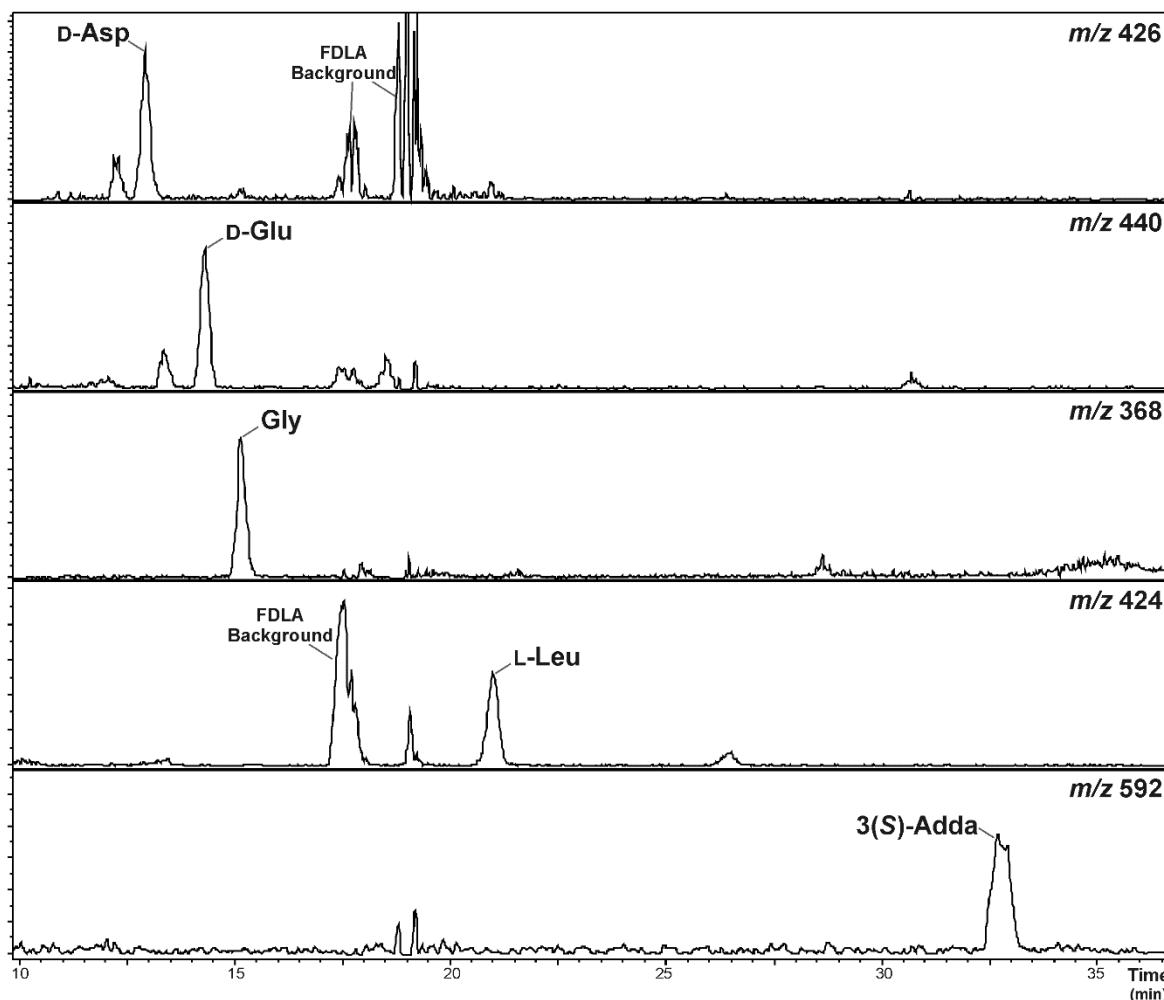


Figure S1. Advanced Marfey's amino acid analysis of the Antarctic-LR microcystin congeners; extracted ion chromatograms of hydrolysates derivatized with L-FDLA (1-fluoro-2,4-dinitrophenyl-5-leucine).

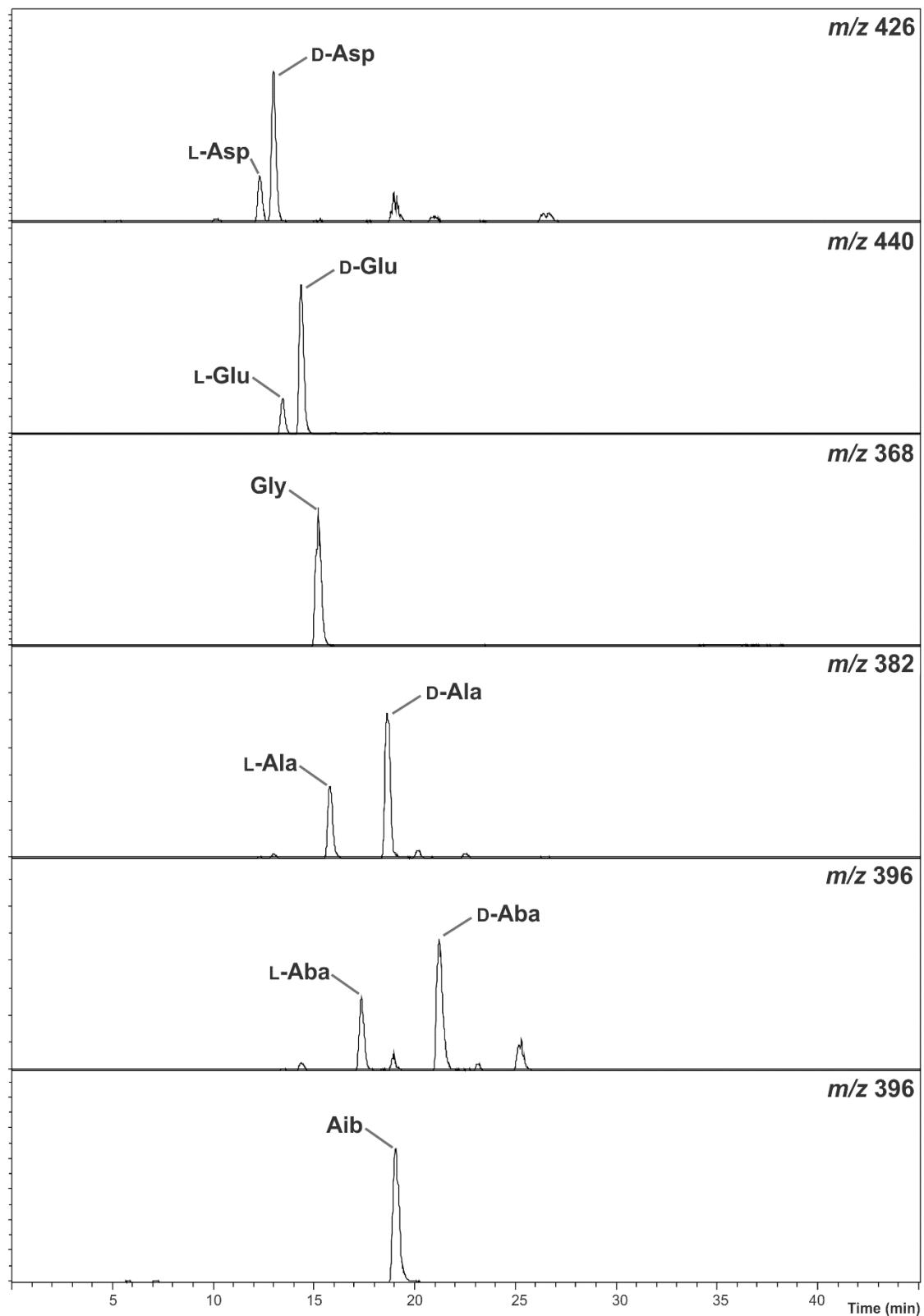


Figure S2. Cont.

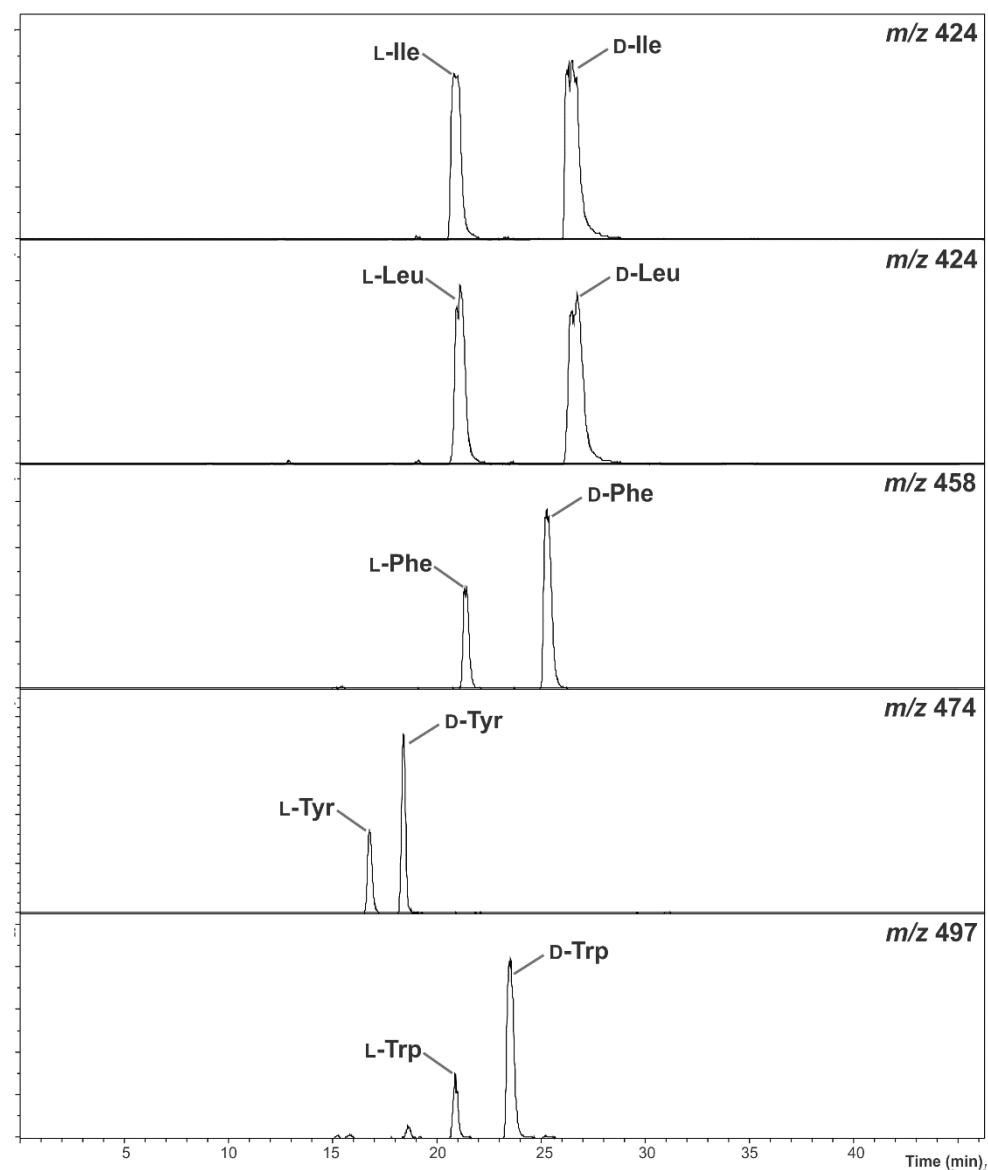


Figure S2. Advanced Marfey's amino acid analysis of standard amino acids; extracted ion chromatograms of amino acids derivatized with DL-FDLA.

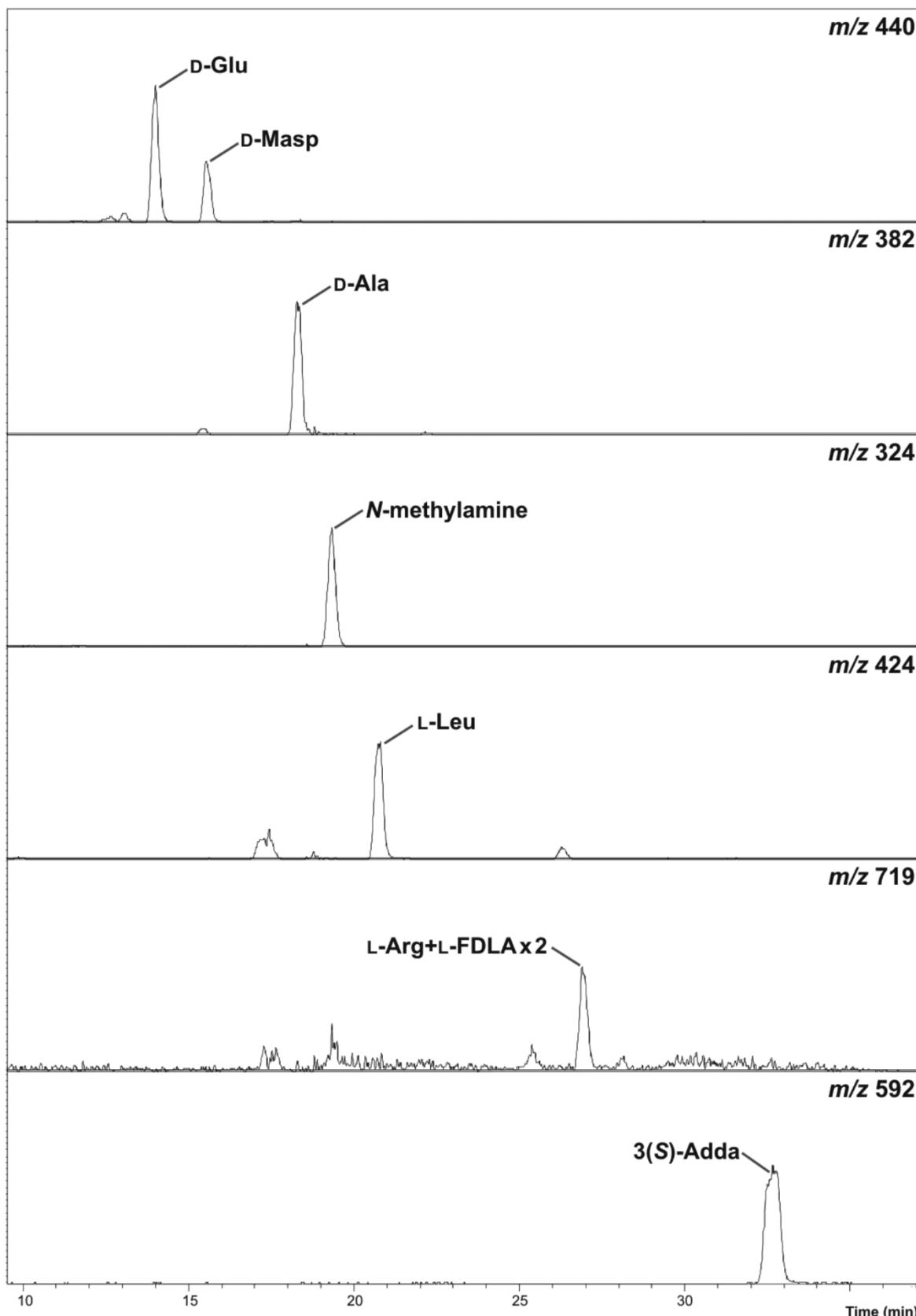


Figure S3. Advanced Marfey's amino acid analysis of MC-LR; extracted ion chromatograms of hydrolyzed MC-LR derivatized with L-FDLA.

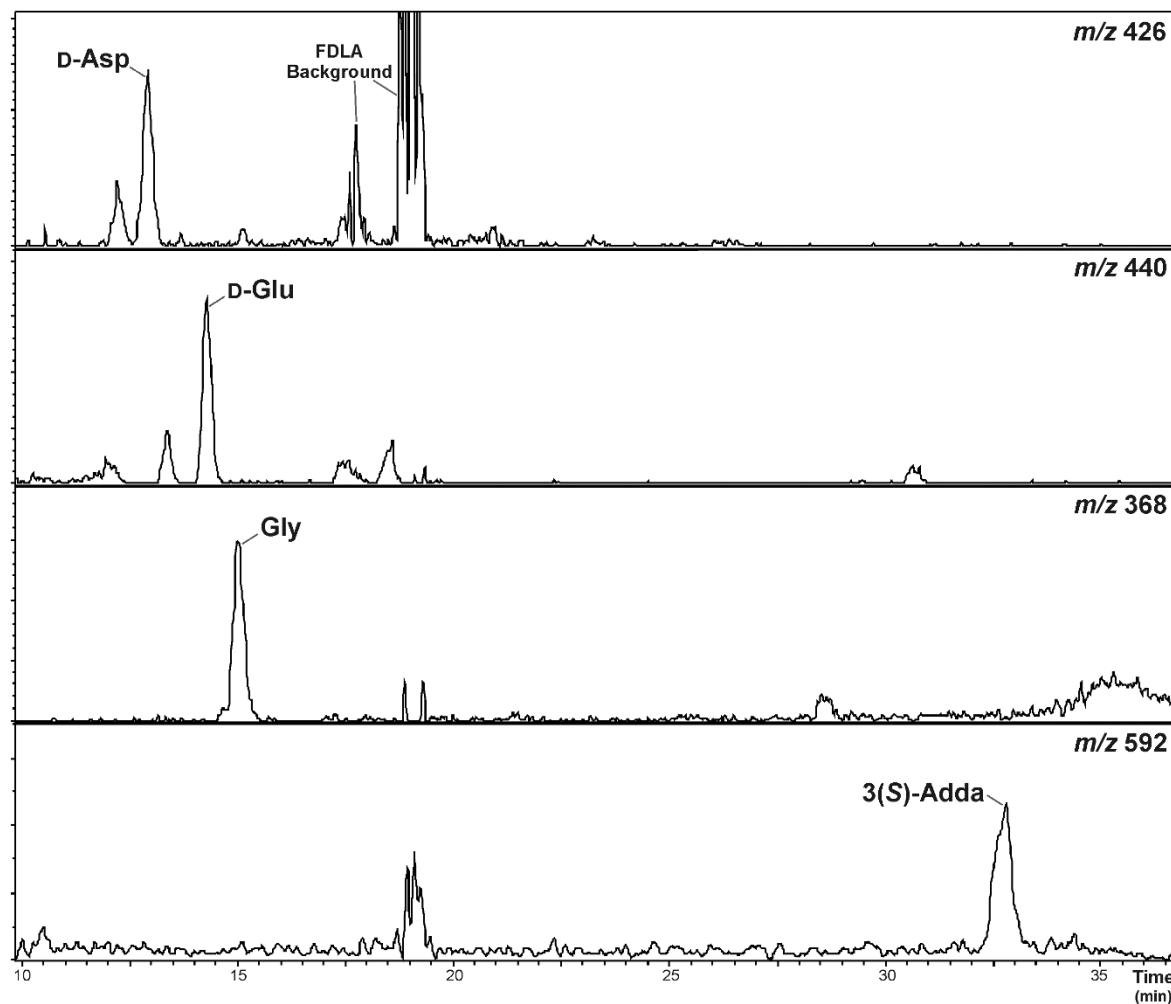


Figure S4. Advanced Marfey's amino acid analysis of the Antarctic-RR microcystin congeners; extracted ion chromatograms of hydrolysates derivatized with L-FDLA.