

Supporting information to the original publication titled: Mass Spectrometric Detection and Characterization of Metabolites of Gemini Surfactants Used as Gene Delivery Vectors. J. Am. Soc. Mass Spectrom. 2020, 31, 2, 366-378. doi.org/10.1021/jasms.9b00004

Supporting Information

Mass spectrometric detection and characterization of metabolites of gemini surfactants used as gene delivery Vectors

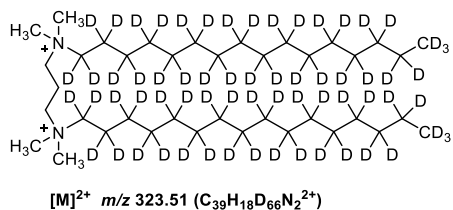
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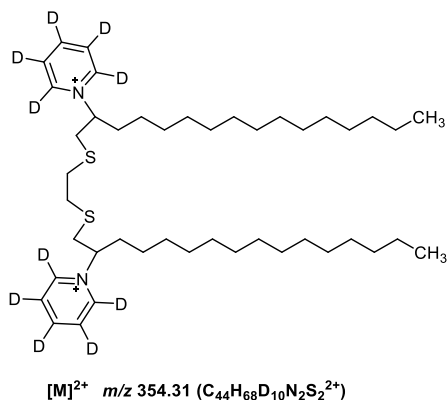
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a 16-3-16-D₆₆



b 16(Py)-S-2-S-16(Py)-D₁₀



c 16-7N(GK)-16-D₄

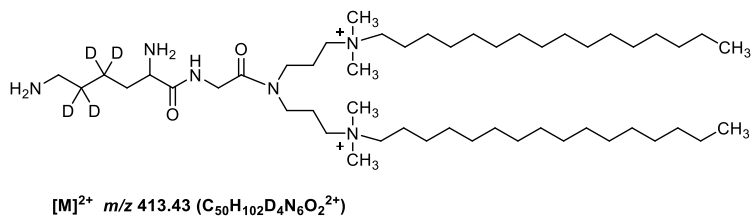


Figure S1, The structures of deuterated gemini surfactants, a) 16-3-16-D₆₆, b) 16(Py)-S-2-S-16(Py)-D₁₀, and c) 16-7N(GK)-16-D₄

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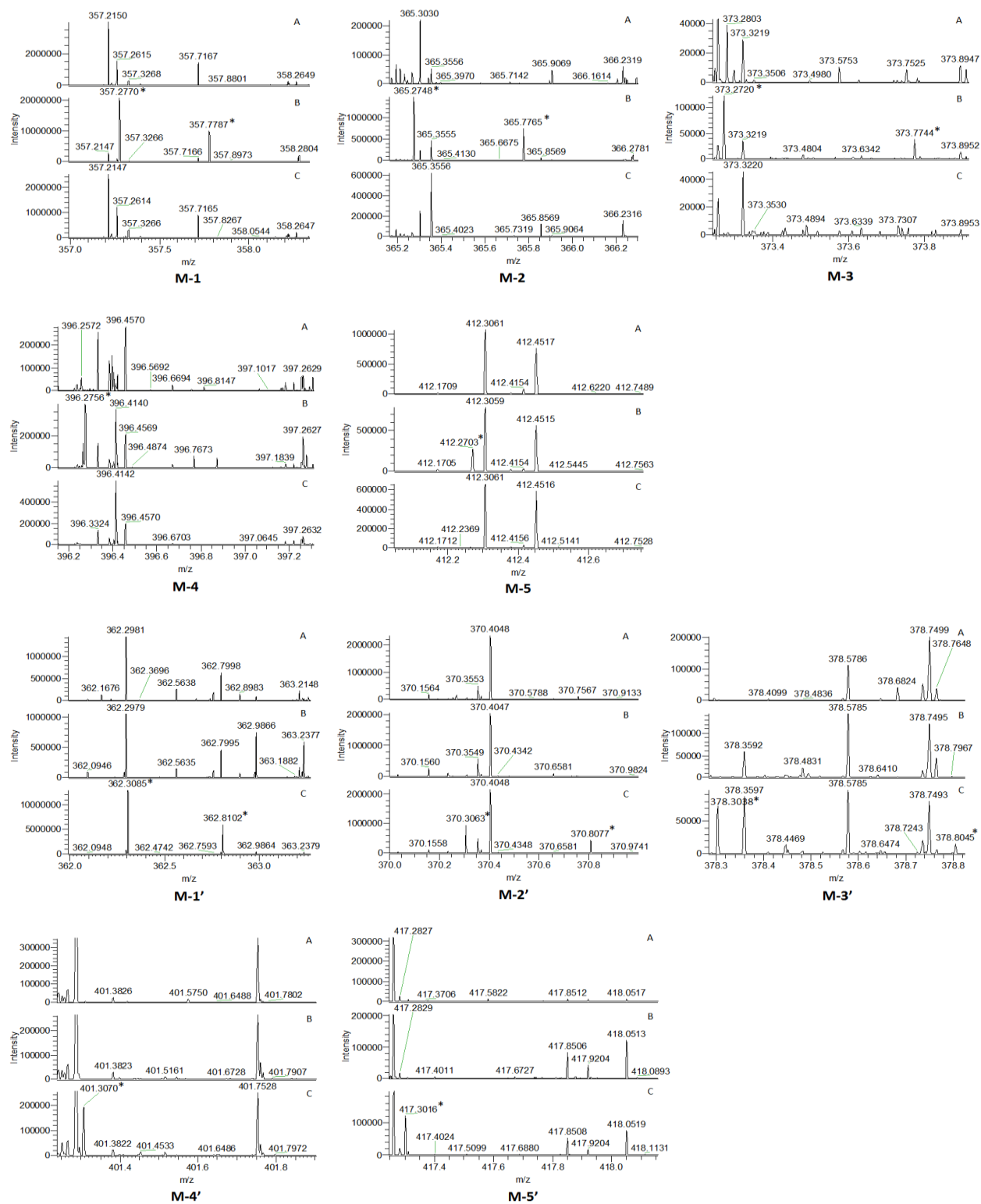


Figure S2. MS spectra of the metabolites of 16(Py)-S-2-S-16(Py) and 16(Py)-S-2-S-16(Py)-D₁₀ based on accurate mass measurement at 5 h of treatment. (* denotes the peak of the metabolite. panel A- control, panel B- cells

treated with 16(Py)-S-2-S-16(Py) nanoparticles, and panel C- cells treated with 16(Py)-S-2-S-16(Py)-D₁₀ nanoparticles. Zoomed in all three panels to show the peaks of the metabolites).

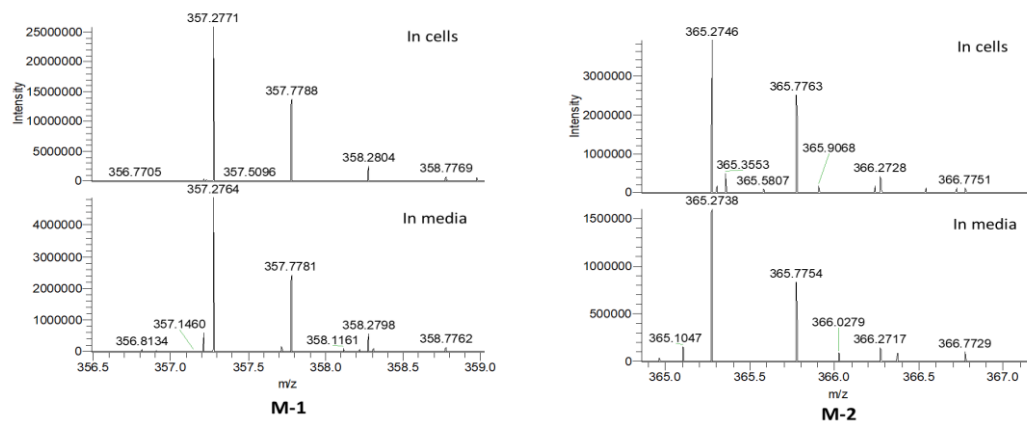


Figure S3. MS spectra of metabolite M-1 (m/z 357) and M-2 (m/z 365) of 16(Py)-S-2-S-16(Py) produced in PAM 212 cells and media at 10 h of treatment. The ion intensity of M-1 and M-2 generated by natural oxidation of 16(Py)-S-2-S-16(Py) in media is lower compared with that within cells. Ion count was 18% in the media for M-1 in comparison to treated cells while it was 41% for M-2. This indicates that oxidation metabolism occurred in cells (i.e, biological system).

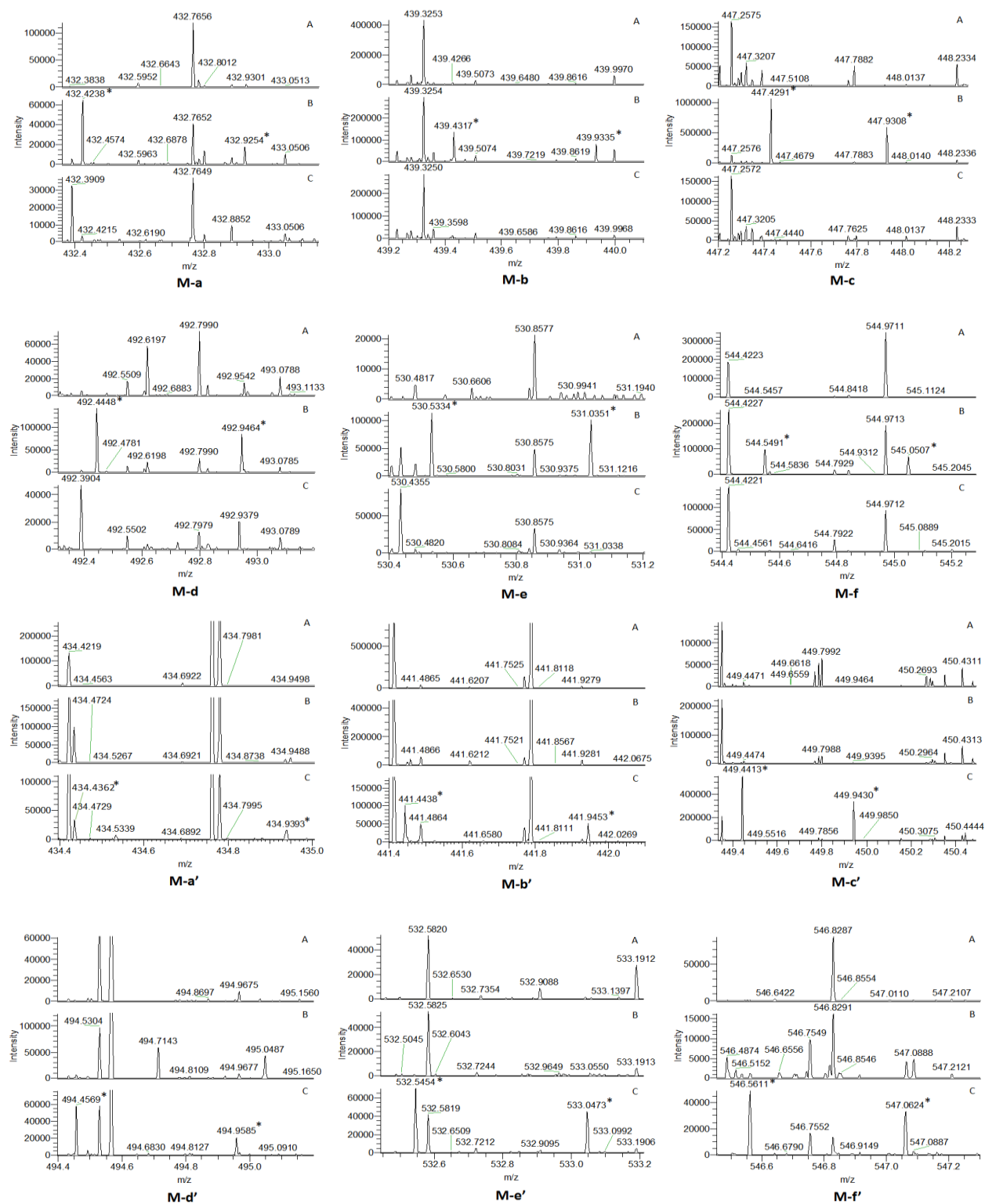


Figure S4. MS spectra of the metabolites of 16-7N(GK)-16 and 16-7N(GK)-16-D₄ based on accurate mass measurement at 5 h treatment. (* denotes the peaks of the identified metabolites. panel A- control, panel B-

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cells treated with 16-7N(GK)-16 nanoparticles, and panel C- cells treated with 16-7N(GK)-16-D₄ nanoparticles. Zoomed in to show the peaks of the metabolites).

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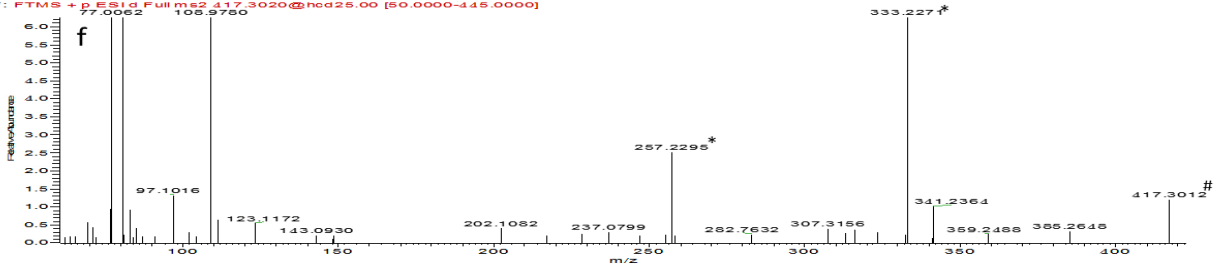
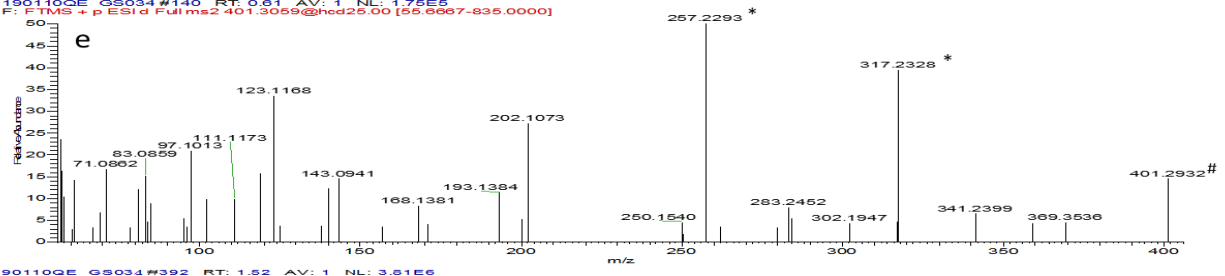
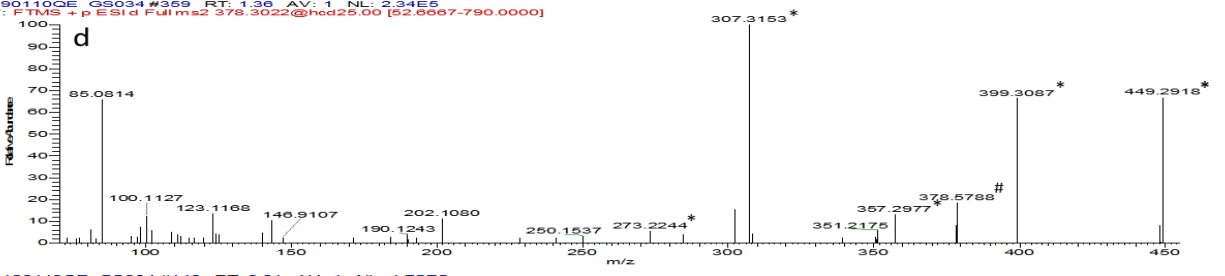
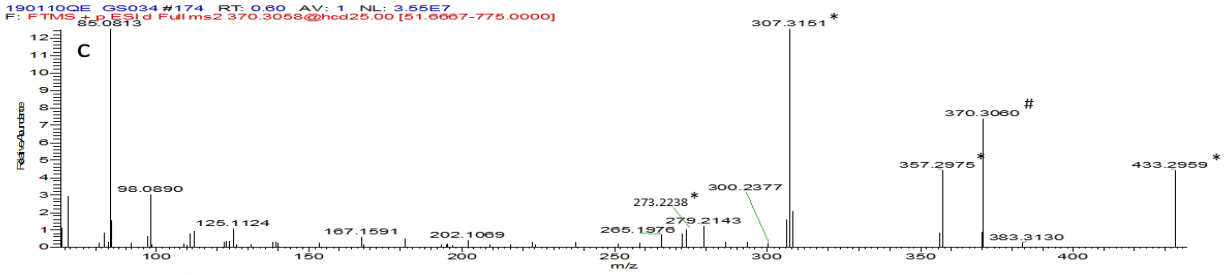
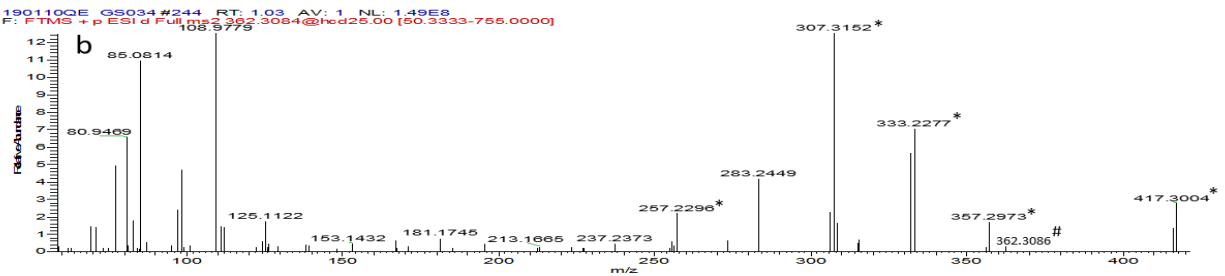
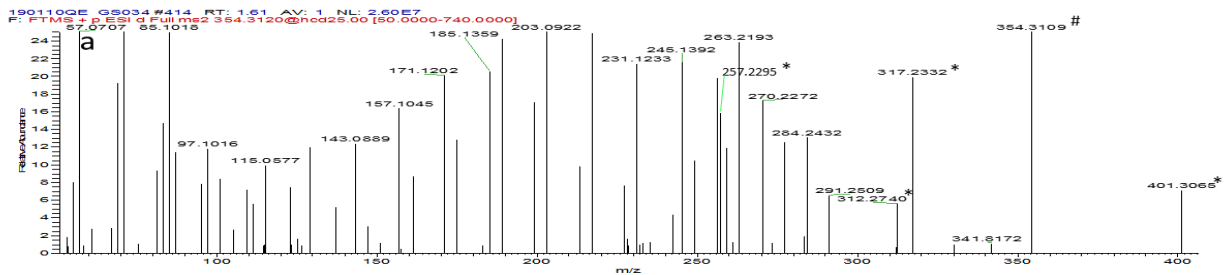


Figure S5. The MS/MS spectra of 16(Py)-S-2-S-16(Py)-D₁₀ and its metabolites: a) 16(Py)-S-2-S-16(Py)-D₁₀, b) M-1', c) M-2', d) M-3', e) M-4', and f) M-5' (# denotes the parent ion and * denotes the diagnostic product ion).

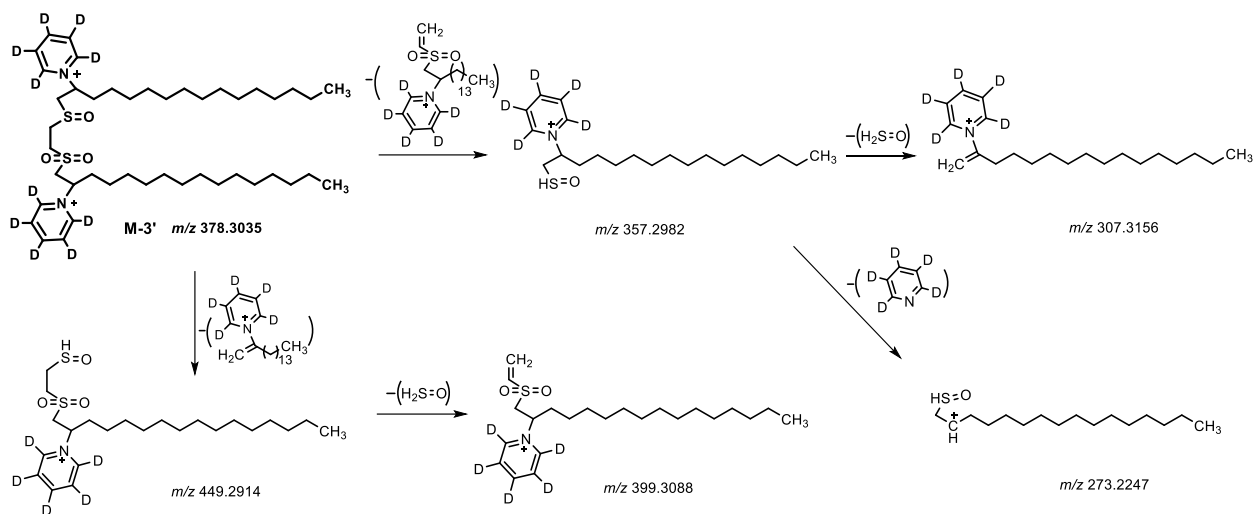


Figure S6. The proposed structures of major product ions of metabolite M-3' of 16(Py)-S-2-S-16(Py)-D₁₀,

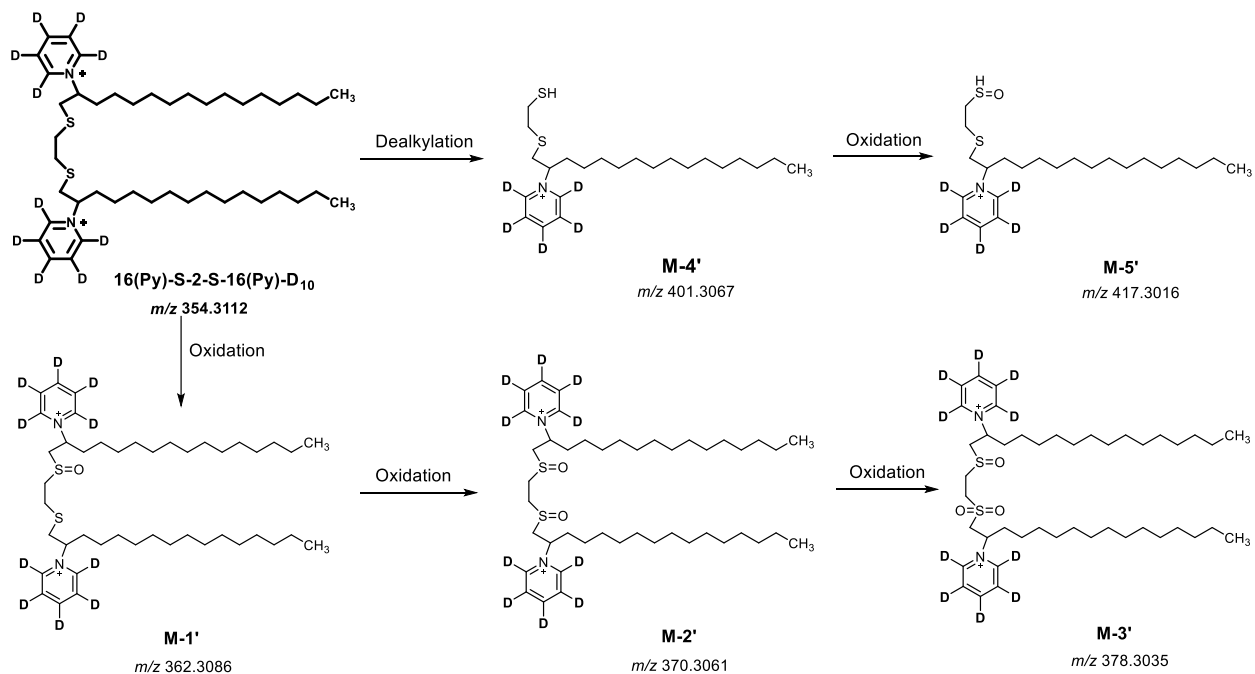


Figure S7. The metabolites of 16(Py)-S-2-S-16(Py)-D₁₀ and its proposed metabolic pathway.

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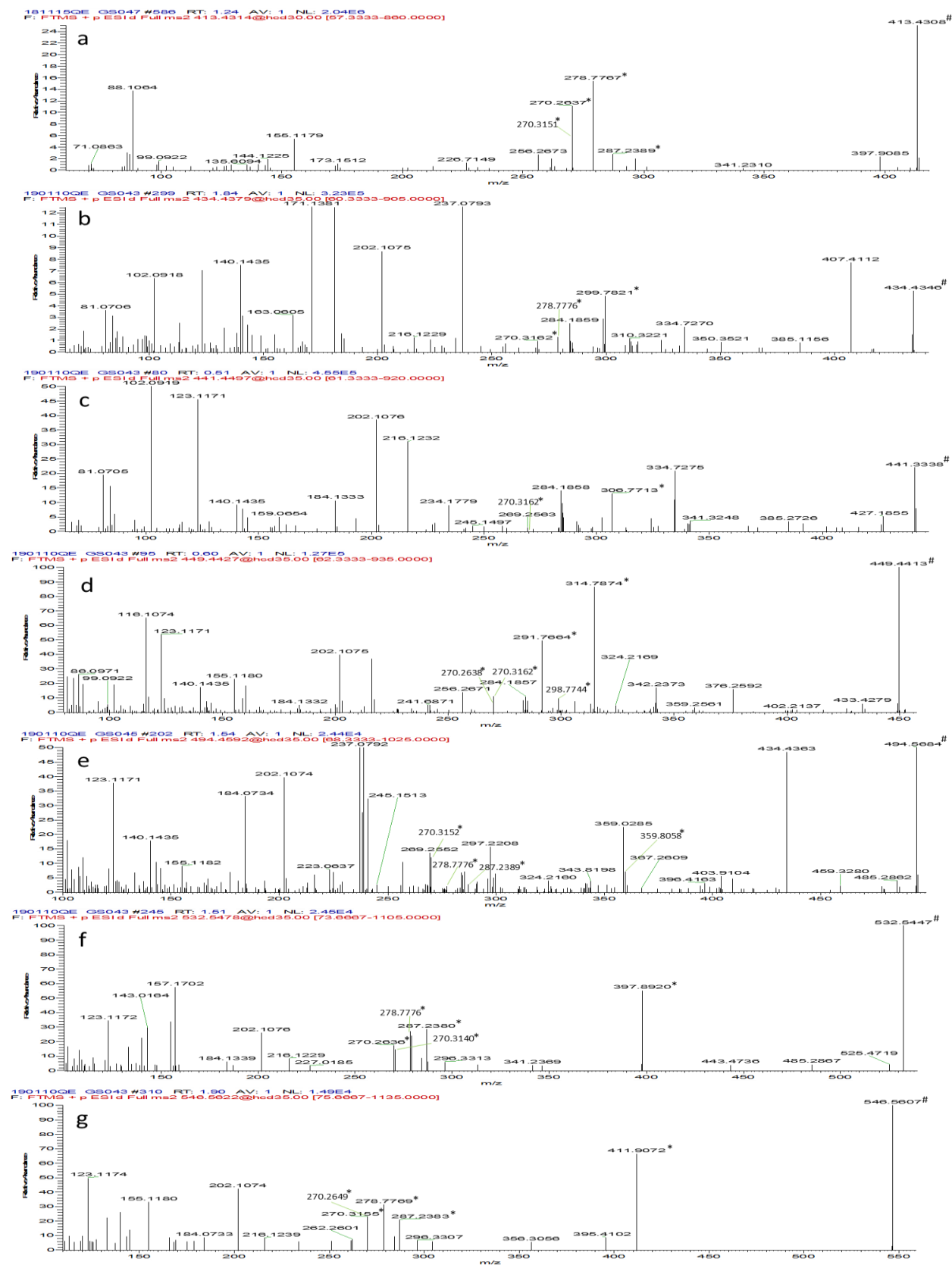


Figure S8. The MS/MS spectra of 16-7N(GK)-16-D₄ and its metabolites: a) 16-7N(GK)-16-D₄, b) M-a', c) M-b', d) M-c', e) M-d', f) M-e', and g) M-f'. (# denotes the parent ions and * denotes the diagnostic product ions).

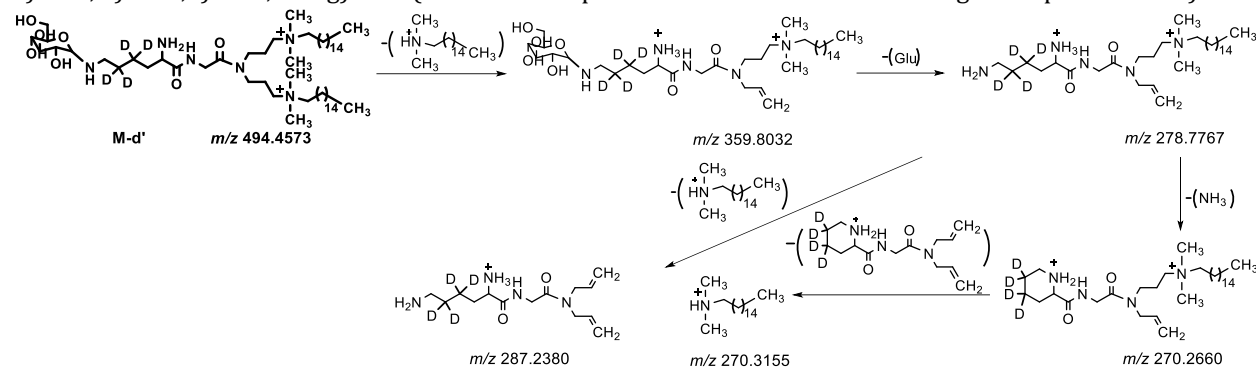
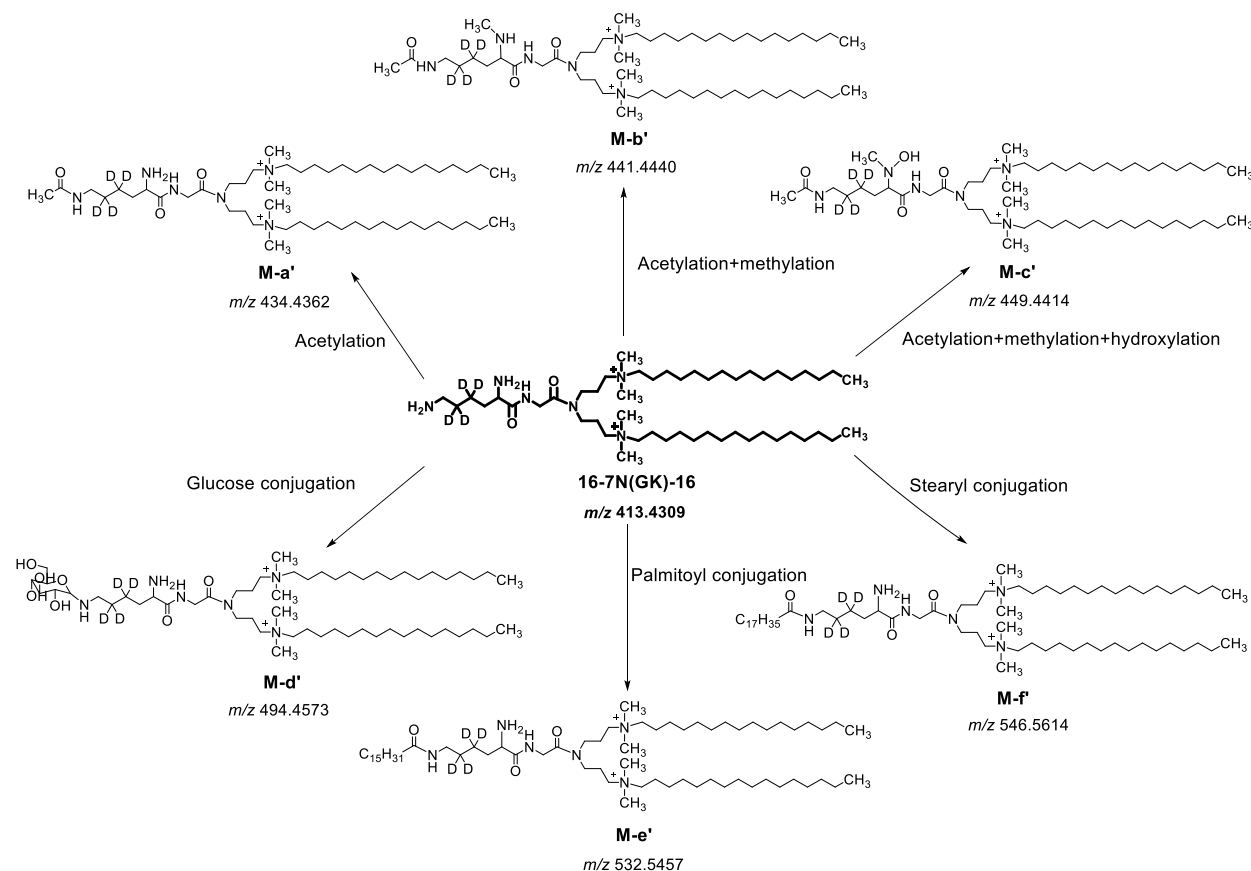


Figure S9. The proposed structures of major product ions of the metabolite M-d' of 16-7N(GK)-16-D₄



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Figure S10. The metabolites of 16-7N(GK)-16-D₄ and its proposed metabolic pathway.

Table S1. Potential metabolites based on accurate mass measurements for the gemini surfactants 16(Py)-S-2-S-16(Py)-D₁₀ and 16-7N(GK)-16-D₄.

Name	Molecular Formula	Exact Mass (m/z)	Observed Mass (m/z)	Mass Accuracy (ppm)	Metabolic Reactions
16(Py)-S-2-S-16(Py)-D ₁₀	C ₄₄ H ₆₈ D ₁₀ N ₂ S ₂ ²⁺	354.3112	354.3115	0.8	NA
M-1'	C ₄₄ H ₆₈ D ₁₀ N ₂ OS ₂ ²⁺	362.3086	362.3085	0.3	Oxidation
M-2'	C ₄₄ H ₆₈ D ₁₀ N ₂ O ₂ S ₂ ²⁺	370.3061	370.3063	0.5	Oxidation
M-3'	C ₄₄ H ₆₈ D ₁₀ N ₂ O ₃ S ₂ ²⁺	378.3035	378.3038	0.8	Oxidation
M-4'	C ₂₃ H ₃₇ D ₅ NS ₂ ⁺	401.3067	401.3070	0.7	Dealkylation
M-5'	C ₂₃ H ₃₇ D ₅ NOS ₂ ⁺	417.3016	417.3016	0.0	Dealkylation, Oxidation
16-7N(GK)-16-D ₄	C ₅₀ H ₁₀₂ D ₄ N ₆ O ₂ ²⁺	413.4309	413.4307	0.5	NA
M-a'	C ₅₂ H ₁₀₄ D ₄ N ₆ O ₃ ²⁺	434.4362	434.4362	0.0	Acetylation
M-b'	C ₅₃ H ₁₀₆ D ₄ N ₆ O ₃ ²⁺	441.4440	441.4438	0.5	Acetylation, Methylation
M-c'	C ₅₃ H ₁₀₆ D ₄ N ₆ O ₄ ²⁺	449.4414	449.4413	0.2	Acetylation, Methylation, Hydroxylation
M-d'	C ₅₆ H ₁₁₂ D ₄ N ₆ O ₇ ²⁺	494.4573	494.4569	0.8	Glucose conjugation
M-e'	C ₆₆ H ₁₃₂ D ₄ N ₆ O ₃ ²⁺	532.5457	532.5454	0.6	Palmitoyl conjugation
M-f'	C ₆₈ H ₁₃₆ D ₄ N ₆ O ₃ ²⁺	546.5614	546.5611	0.5	Stearyl conjugation