

# Supplementary Information

## Studies of Surface Preparation for the Fluorosequencing of Peptides

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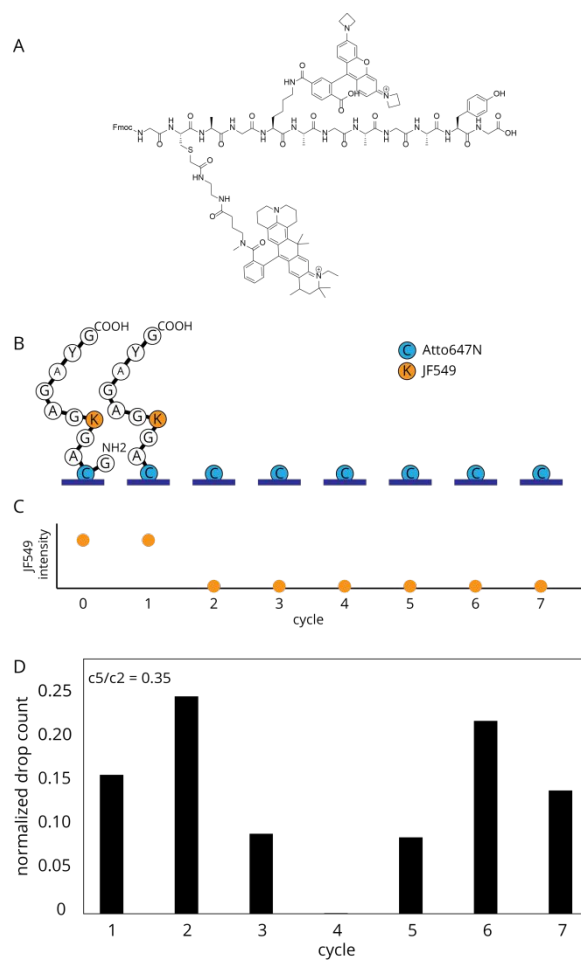
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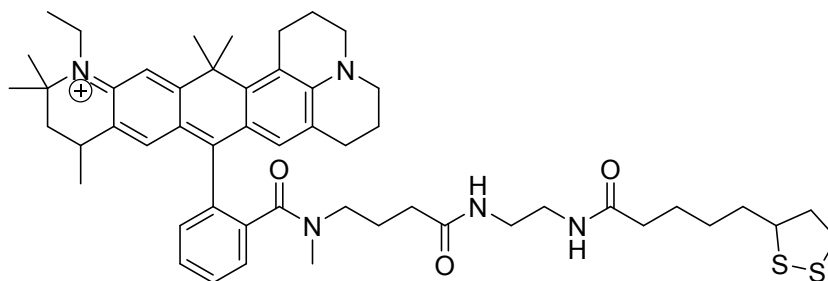
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Figure S1



**FIGURE S1.** A) Structure of peptide PS1 B) Expected sequence changes of a single PS1 peptide attached through the Atto647N after rounds of Edman degradation C) The relative fluorescence changes for Atto647N on JF549 attached peptide P2 where signal loss is correlated to liberation of the remaining peptide after Edman degradation at the labeled lysine. D) Fluorosequencing results for peptides PS1, with a peak drop count after the second Edman cycle (normalized to total counts, n=6450).

Figure S2: HiRes MS data for modified Atto647N for aldehyde labeling

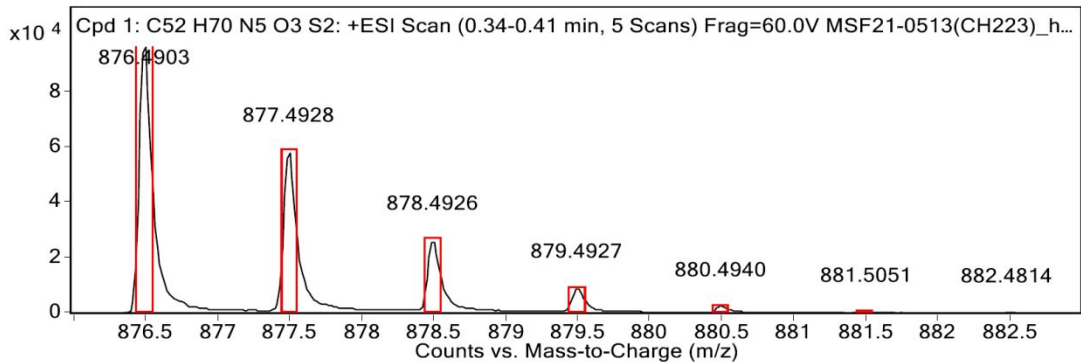


Chemical Formula:  $C_{52}H_{70}N_5O_3S_2^+$

Exact Mass: 876.49

m/z: 876.49 (100.0%), 877.50 (57.2%), 878.50 (16.7%), 878.49 (11.0%), 879.49 (5.3%), 879.50 (3.9%),  
877.49 (3.4%), 880.49 (1.6%)

MS Zoomed Spectrum

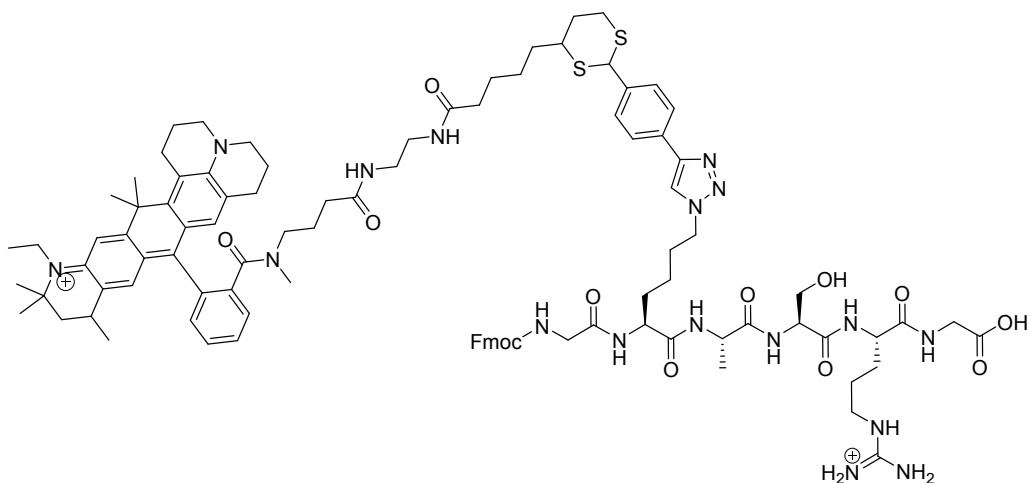


MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
876.4903	876.4915	1	97795	$C_{52}H_{70}N_5O_3S_2$	M+	1.28
877.4928	877.4946	1	58535	$C_{52}H_{70}N_5O_3S_2$	M+	2.04
878.4926	878.4942	1	26609	$C_{52}H_{70}N_5O_3S_2$	M+	1.87
879.4927	879.4947	1	9192	$C_{52}H_{70}N_5O_3S_2$	M+	2.26
880.4940	880.4951	1	2453	$C_{52}H_{70}N_5O_3S_2$	M+	1.19
881.5051	881.4956	1	455	$C_{52}H_{70}N_5O_3S_2$	M+	-10.77
882.4814	882.4963	1	229	$C_{52}H_{70}N_5O_3S_2$	M+	16.96

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Figure S3: HiRes MS data for fluorophore labeling of P1

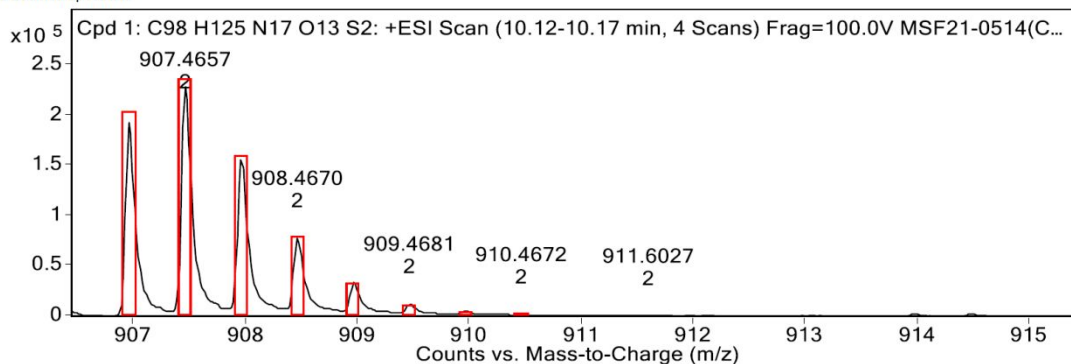


Chemical Formula:  $C_{98}H_{127}N_{17}O_{13}S_2^{2+}$

Exact Mass: 1813.92

m/z: 907.46 (100.0%), 906.96 (92.6%), 907.96 (57.6%), 908.46 (25.3%), 907.96 (14.8%), 908.46 (9.8%), 907.46 (7.3%), 908.96 (6.6%), 908.97 (6.4%), 909.46 (2.4%), 909.47 (1.6%), 908.96 (1.2%)

MS Zoomed Spectrum

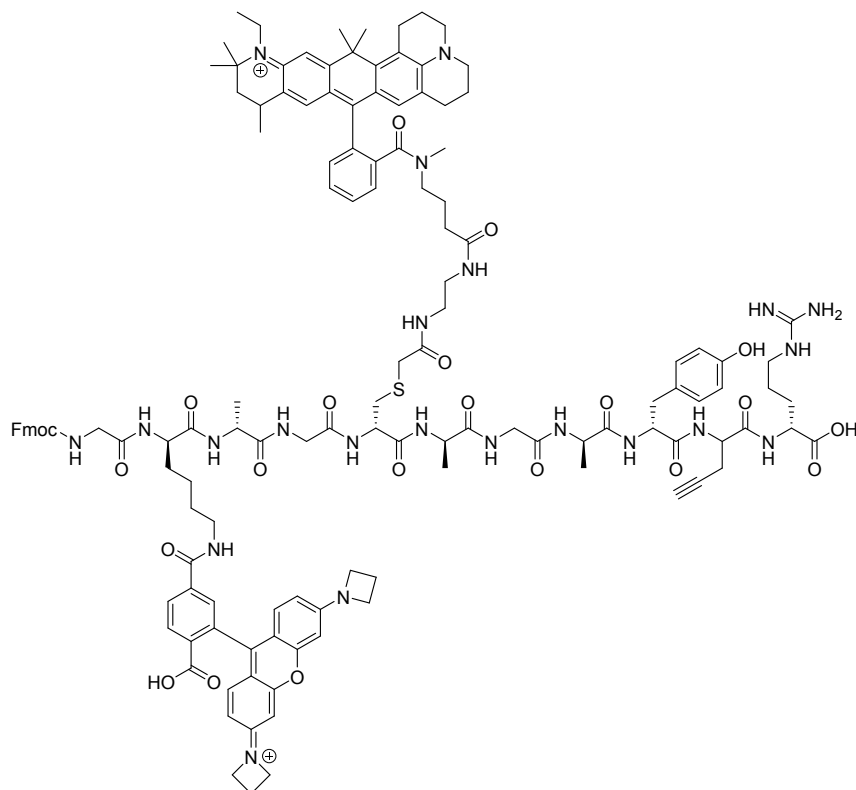


MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
605.3137			300642			
906.9647	906.9615	2	194673	C98H125N17O13S2	(M+2H)+2	-3.55
907.4657	907.4630	2	230659	C98H125N17O13S2	(M+2H)+2	-2.99
907.9664	907.9639	2	157803	C98H125N17O13S2	(M+2H)+2	-2.77
908.4670	908.4646	2	78569	C98H125N17O13S2	(M+2H)+2	-2.65
908.9686	908.9652	2	34409	C98H125N17O13S2	(M+2H)+2	-3.77
909.4681	909.4658	2	12950	C98H125N17O13S2	(M+2H)+2	-2.51
909.9609	909.9664	2	5052	C98H125N17O13S2	(M+2H)+2	6.12
910.4672	910.4671	2	2436	C98H125N17O13S2	(M+2H)+2	-0.09
911.0305	910.9678	2	1523	C98H125N17O13S2	(M+2H)+2	-68.76

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Figure S4: HiRes MS data for Fluorophore Labeling of P2



Chemical Formula:  $C_{132}H_{158}N_{22}O_{22}S^{2+}$

Exact Mass: 2435.16

m/z: 1218.08 (100.0%), 1218.58 (76.2%), 1217.58 (68.8%), 1219.08 (43.9%), 1219.59 (15.5%), 1218.58 (11.3%), 1219.58 (6.8%), 1218.08 (6.1%), 1220.09 (5.2%), 1219.08 (5.1%), 1220.08 (3.1%), 1219.09 (1.9%), 1220.59 (1.8%)

Elemental Analysis: C, 65.06; H, 6.54; N, 12.65; O, 14.44; S, 1.32

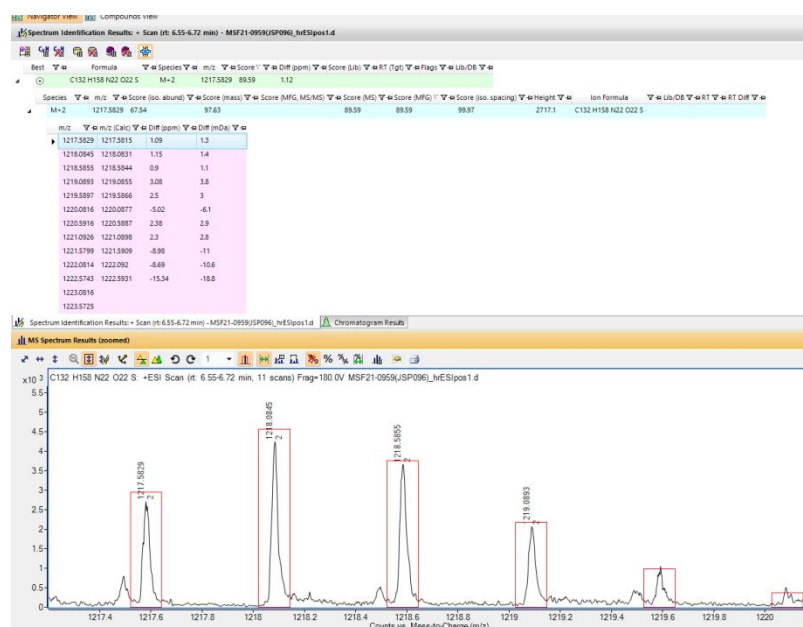
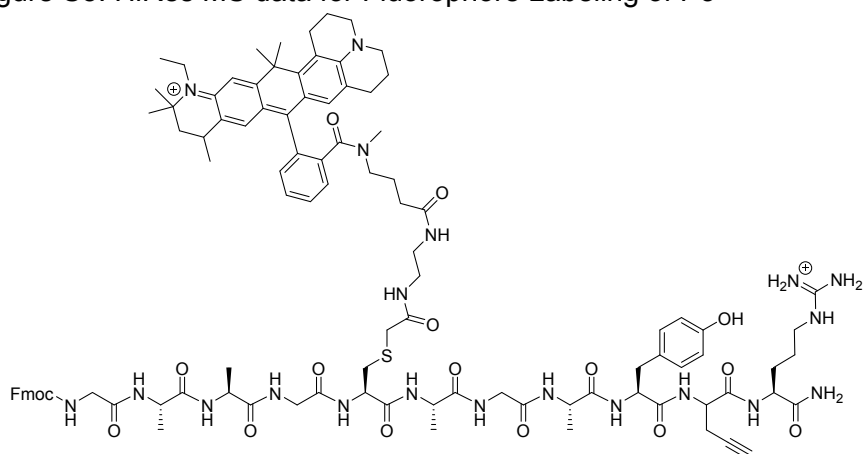


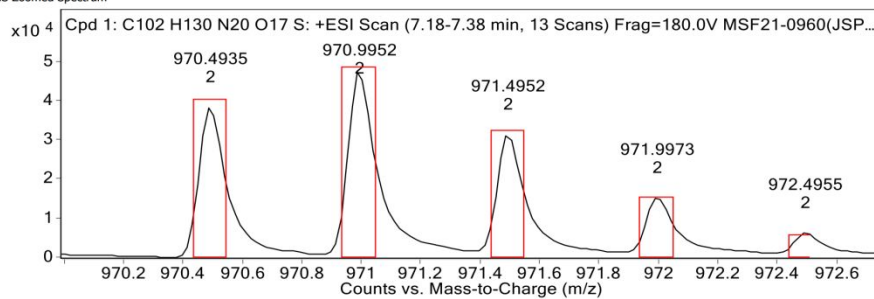
Figure S5: HiRes MS data for Fluorophore Labeling of P3



Chemical Formula:  $C_{102}H_{132}N_{20}O_{17}S^{2+}$   
Exact Mass: 1940.98

m/z: 970.99 (100.0%), 970.49 (83.9%), 971.49 (52.6%), 971.99 (23.2%), 971.49 (14.5%), 971.99 (8.6%), 972.49 (8.6%),  
972.49 (3.2%), 972.99 (2.1%), 970.99 (1.3%), 973.00 (1.2%)  
Elemental Analysis: C, 63.07; H, 6.85; N, 14.42; O, 14.00; S, 1.65

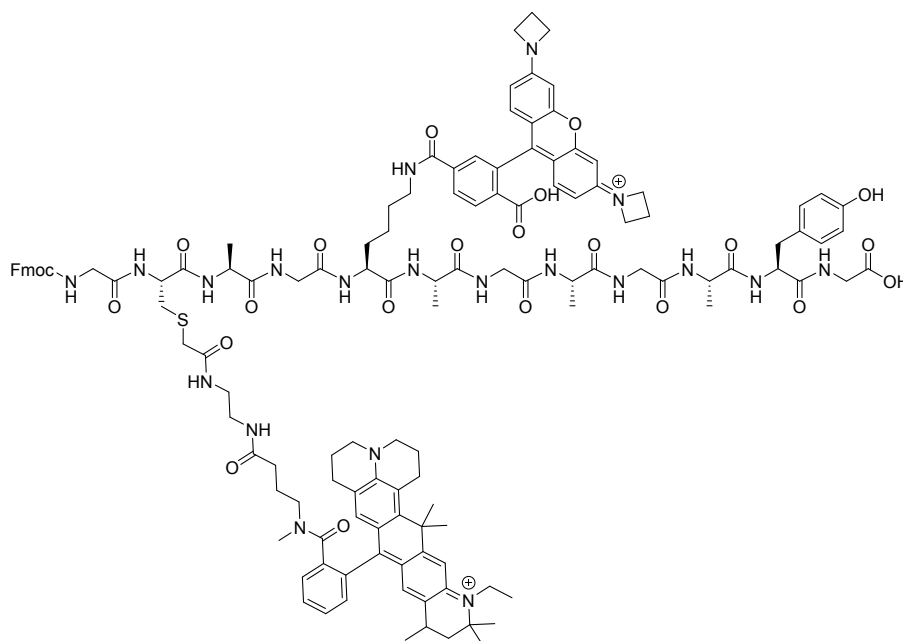
MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
265.1787			242900			
970.4935	970.4895	2	38608	C <sub>102</sub> H <sub>130</sub> N <sub>20</sub> O <sub>17</sub> S	(M+2H) <sup>2+</sup>	-4.15
970.9952	970.9909	2	47953	C <sub>102</sub> H <sub>130</sub> N <sub>20</sub> O <sub>17</sub> S	(M+2H) <sup>2+</sup>	-4.42
971.4952	971.4921	2	31633	C <sub>102</sub> H <sub>130</sub> N <sub>20</sub> O <sub>17</sub> S	(M+2H) <sup>2+</sup>	-3.12
971.9973	971.9931	2	15476	C <sub>102</sub> H <sub>130</sub> N <sub>20</sub> O <sub>17</sub> S	(M+2H) <sup>2+</sup>	-4.35
972.4955	972.4941	2	6530	C <sub>102</sub> H <sub>130</sub> N <sub>20</sub> O <sub>17</sub> S	(M+2H) <sup>2+</sup>	-1.51
973.0350	972.9950	2	2973	C <sub>102</sub> H <sub>130</sub> N <sub>20</sub> O <sub>17</sub> S	(M+2H) <sup>2+</sup>	-41.12
973.4877	973.4960	2	1248	C <sub>102</sub> H <sub>130</sub> N <sub>20</sub> O <sub>17</sub> S	(M+2H) <sup>2+</sup>	8.52

Figure S6: HiRes MS data for Fluorophore Labeling of PS1



Chemical Formula:  $C_{128}H_{152}N_{20}O_{23}S^{2+}$

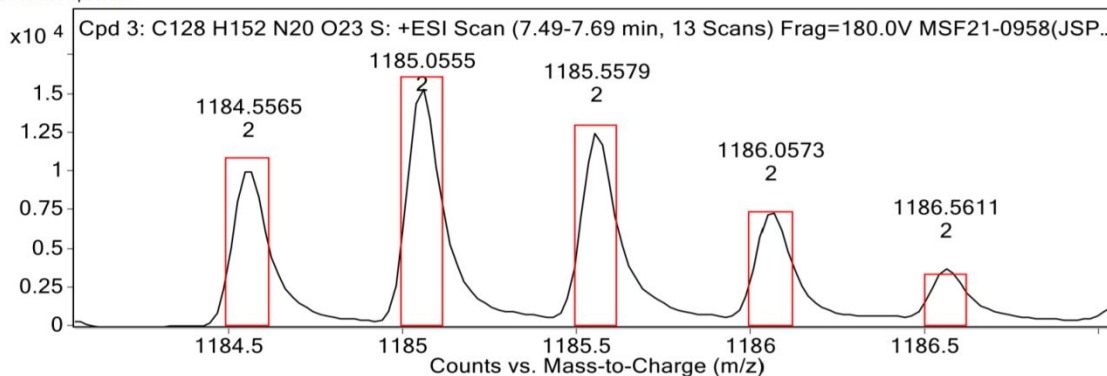
Exact Mass: 2369.10

Molecular Weight: 2370.80

m/z: 1185.05 (100.0%), 1185.55 (79.4%), 1184.55 (70.5%), 1186.06 (32.3%), 1186.05 (15.1%), 1186.56 (14.7%), 1186.55 (6.3%), 1187.06 (5.4%), 1185.05 (5.2%), 1185.55 (3.4%), 1187.05 (2.3%), 1185.56 (1.7%), 1187.56 (1.4%)

Elemental Analysis: C, 64.85; H, 6.46; N, 11.82; O, 15.52; S, 1.35

MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
797.7037			109711			
1184.5565	1184.5524	2	10248	C <sub>128</sub> H <sub>152</sub> N <sub>20</sub> O <sub>23</sub> S	M+2	-3.41
1185.0555	1185.0540	2	15393	C <sub>128</sub> H <sub>152</sub> N <sub>20</sub> O <sub>23</sub> S	M+2	-1.3
1185.5579	1185.5553	2	12527	C <sub>128</sub> H <sub>152</sub> N <sub>20</sub> O <sub>23</sub> S	M+2	-2.21
1186.0573	1186.0564	2	7438	C <sub>128</sub> H <sub>152</sub> N <sub>20</sub> O <sub>23</sub> S	M+2	-0.76
1186.5611	1186.5575	2	3739	C <sub>128</sub> H <sub>152</sub> N <sub>20</sub> O <sub>23</sub> S	M+2	-3
1187.0540	1187.0586	2	1453	C <sub>128</sub> H <sub>152</sub> N <sub>20</sub> O <sub>23</sub> S	M+2	3.89
1187.5691	1187.5596	2	724	C <sub>128</sub> H <sub>152</sub> N <sub>20</sub> O <sub>23</sub> S	M+2	-8.01
1188.0141	1188.0607	2	356	C <sub>128</sub> H <sub>152</sub> N <sub>20</sub> O <sub>23</sub> S	M+2	39.25
1188.5406	1188.5618	2	257	C <sub>128</sub> H <sub>152</sub> N <sub>20</sub> O <sub>23</sub> S	M+2	17.82



Table S1: Average difference between initial and TFA3 of samples in Figure 4.

Average difference between initial and TFA3

	EDC	Buffer only
None	1067 ± 121	357 ± 73
Hexyl	1162 ± 153	340 ± 226
n-propyl	1419 ± 67	357 ± 172
t-butyl	1480 ± 237	508 ± 154
PEG-3	1514 ± 129	185 ± 57

Table S2: Average difference between initial and TFA3 of samples in Figure 5

Average difference between initial and TFA3

	EDC	Buffer only
0 PEG-3	2123 ± 75	1432 ± 78
0.75 PEG-3	623 ± 61	132 ± 16
1.5 PEG-3	1249 ± 367	141 ± 48
2.0 PEG-3	1194 ± 251	224 ± 32
2.5 PEG-3	936 ± 245	262 ± 27
0 PEG-3	1354 ± 188	327 ± 53
0.75 PEG-3	1053 ± 155	178 ± 51
1.5 PEG-3	895 ± 101	211 ± 50
2.0 PEG-3	1287 ± 78	274 ± 59
2.5 PEG-3	1064 ± 109	345 ± 70
0 PEG-3	149 ± 12	103 ± 15
0 PEG-3	532 ± 48	516 ± 95
0.75 PEG-3	439 ± 59	128 ± 20
1.5 PEG-3	799 ± 38	300 ± 15
2.0 PEG-3	538 ± 217	280 ± 69
2.5 PEG-3	857 ± 69	170 ± 20
0 PEG-3	1034 ± 52	775 ± 73
0 PEG-3	373 ± 28	219 ± 29
0.75 PEG-3	1289 ± 166	825 ± 52
1.5 PEG-3	434 ± 52	332 ± 39
2.0 PEG-3	609 ± 82	529 ± 58

Table S3: Average difference between initial and TFA3 of samples in Figure 6

Average difference between initial and TFA3		
	EDC	Buffer only
0PEG0APTES_1	1581 ± 73	857 ± 122
0PEG0APTES_2	1110 ± 245	376 ± 61
0PEG0APTES_3	1578 ± 107	1399 ± 71
0PEG1APTES_1	619 ± 80	496 ± 43
0PEG1APTES_2	164 ± 127	892 ± 65
0PEG1APTES_3	241 ± 60	214 ± 31
2PEG0APTES_1	1234 ± 141	713 ± 80
2PEG0APTES_2	1874 ± 57	716 ± 80
2PEG0APTES_3	1262 ± 75	829 ± 67
2PEG1APTES_1	694 ± 48	732 ± 44
2PEG1APTES_2	750 ± 55	727 ± 51
2PEG1APTES_3	817 ± 74	589 ± 38