

Supplemental Table 1

NEDD8-UbcH12 Linkage	Peptide Sequence	Predicted <i>m/z</i> NEDD8-modified LysC Peptide	Predicted <i>m/z</i> GG-modified Tryptic Peptide
K8 (aa 4-11)	(K ₃)LFSLKQQK ₁₁ (K)	(2+) 1274.28 (3+) 849.86 (4+) 637.64	(2+) 553.32 (3+) 369.22 (4+) 277.16
K11 (aa 9-12)	(K ₈)QQKK ₁₂ (E)	(2+) 1044.14 (3+) 696.43 (4+) 522.58	(2+) 323.19 (3+) 215.79 (4+) 162.10
K12 (aa 12-21)	(K ₁₁)KEEESAGGTK ₂₁ (G)	(2+) 1296.23 (3+) 864.49 (4+) 648.62	(2+) 575.27 (3+) 383.85 (4+) 288.14
K21 (aa 13-25)	(K ₁₂)EEESAGGTKKGSSK ₂₅ (K)	(2+) 1411.77 (3+) 941.52 (4+) 706.39	(2+) 690.81 (3+) 460.88 (4+) 345.91
K25 (aa 22-26)	(K ₂₁)GSSKK ₂₆ (A)	(2+) 1031.63 (3+) 688.09 (4+) 516.32	(2+) 310.67 (3+) 207.45 (4+) 155.84
K26 (aa 26-36 or aa 26-33)	(K ₂₅)KASAAQLRIQK ₃₆ (D) - LysC or (K ₂₅)KASAAQLR ₃₃ (I) - Trypsin	(2+) 1385.35 (3+) 923.90 (4+) 693.18	(2+) 479.77 (3+) 320.19 (4+) 240.39

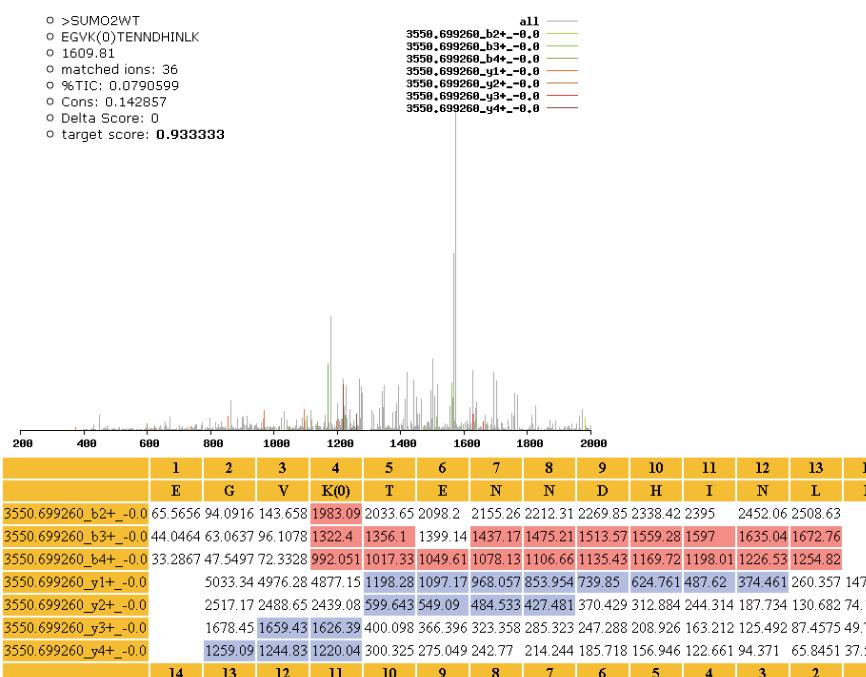
Supplemental Figure 1

A

Scan	Charge	relPos	modScore	precursor mass	matched ions	TIC	%TIC	BPI
6001	4	1	1	1609.81	131	172878	0.455145	10802.8

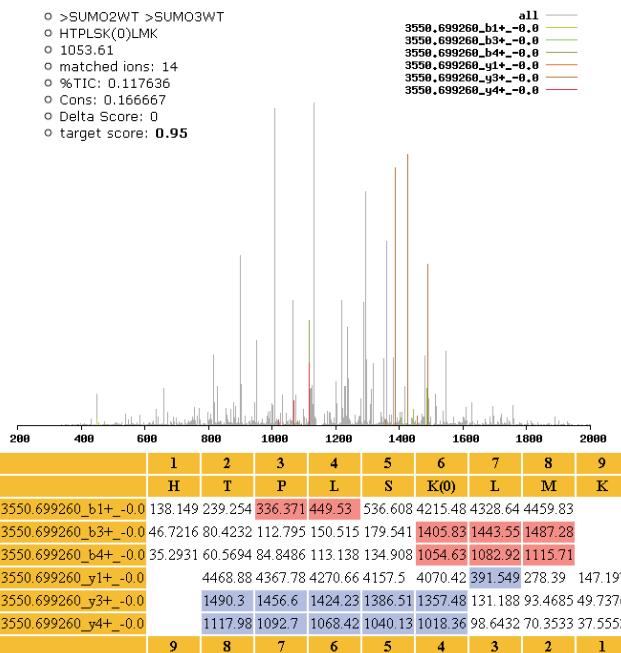
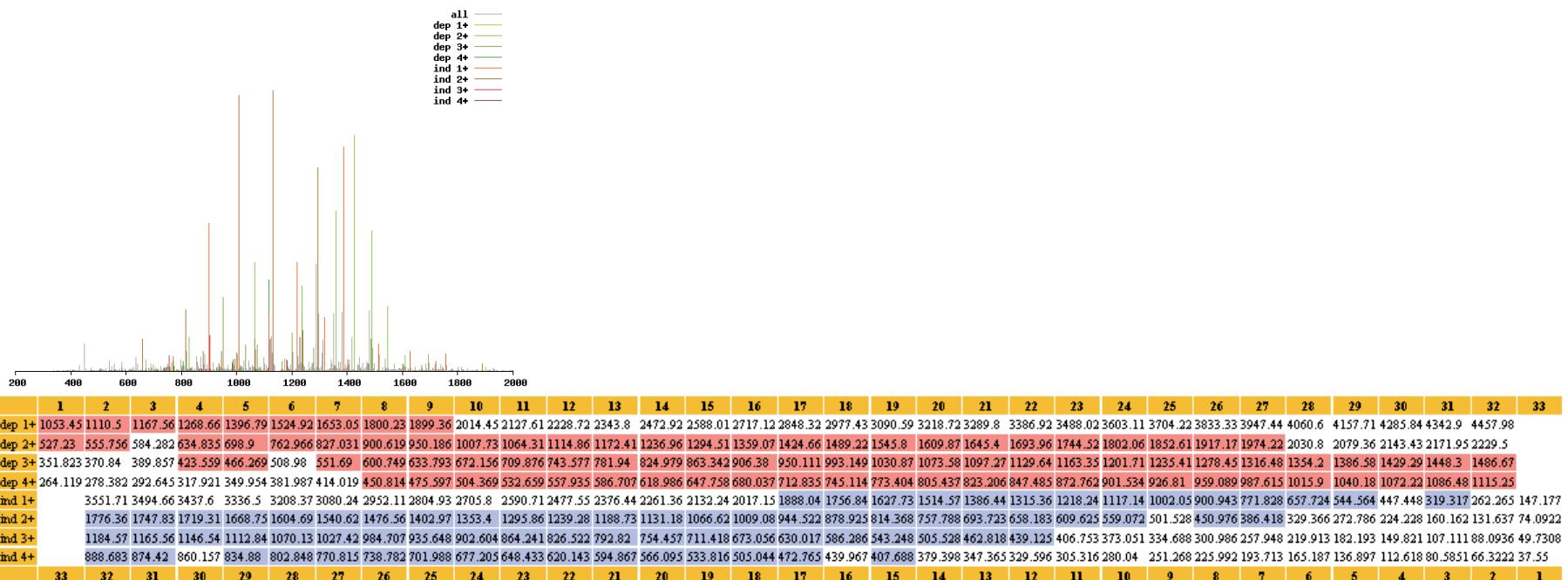


	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33
dep 1+	1609.66	1666.71	1723.76	1824.87	1953	2081.13	2209.26	2256.43	2455.57	2570.66	2683.82	2784.92	2900.01	3029.12	3144.21	3273.33	3404.52	3533.64	3646.8	3774.93	3846.01	3943.12	4044.23	4159.32	4260.42	4389.54	4503.64	4616.8	4713.92	4842.05	4899.1	5014.19	
dep 2+	805.332	833.858	862.384	912.937	977.002	1041.07	1105.13	1178.72	1228.29	1285.83	1342.41	1392.96	1450.51	1515.07	1572.61	1637.17	1702.76	1767.32	1823.9	1887.97	1923.51	1972.07	2022.62	2080.16	2130.71	2195.27	2252.32	2308.9	2357.46	2421.53	2450.05	2507.6	
dep 3+	537.224	556.242	575.259	608.961	651.671	694.381	737.091	786.15	819.194	857.557	895.277	928.979	967.342	1010.38	1048.74	1091.78	1135.51	1178.55	1216.27	1258.98	1282.67	1315.05	1348.73	1387.11	1420.81	1463.85	1501.89	1539.61	1571.98	1614.69	1633.7	1672.07	
dep 4+	403.17	417.433	431.696	456.972	489.005	521.038	533.07	589.865	614.648	643.42	671.71	696.986	725.758	758.037	786.809	819.088	851.886	884.165	912.455	944.488	962.257	986.536	1011.81	1040.58	1065.86	1098.14	1126.67	1154.96	1179.24	1211.27	1225.53	1254.3	
ind 1+	3551.71	3494.66	3437.6	3336.5	3208.37	3080.24	2952.11	2804.93	2705.8	2590.71	2477.55	2376.44	2261.36	2123.24	2017.15	1888.04	1756.84	1627.73	1514.57	1386.44	1315.30	1218.24	1117.14	1002.05	900.943	771.828	657.724	544.564	447.448	319.317	262.265	147.177	
ind 2+	1776.36	1747.83	1719.31	1668.75	1604.69	1540.62	1476.56	1402.97	1353.4	1295.86	1239.28	1188.73	1131.18	1066.62	1009.08	944.522	878.925	814.368	757.788	693.723	658.183	609.625	559.072	501.528	450.976	386.418	329.366	272.786	224.228	160.162	131.637	74.0922	
ind 3+	1184.57	1165.56	1146.54	1112.84	1070.13	1027.42	984.707	935.648	902.604	864.241	826.522	792.82	754.457	711.418	673.056	630.017	586.286	543.248	505.528	462.818	439.125	406.753	373.051	334.688	300.986	257.948	219.913	182.193	149.821	107.11188.0936	49.7308		
ind 4+	888.683	874.42	860.157	834.88	802.848	770.815	738.782	701.988	677.205	648.433	620.143	594.867	566.095	533.816	505.044	472.765	439.967	407.688	379.398	347.365	329.596	305.316	280.04	251.268	225.992	193.713	163.187	136.897	112.618	80.5851	66.3222	37.55	
	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	6	5	4	3	2	1



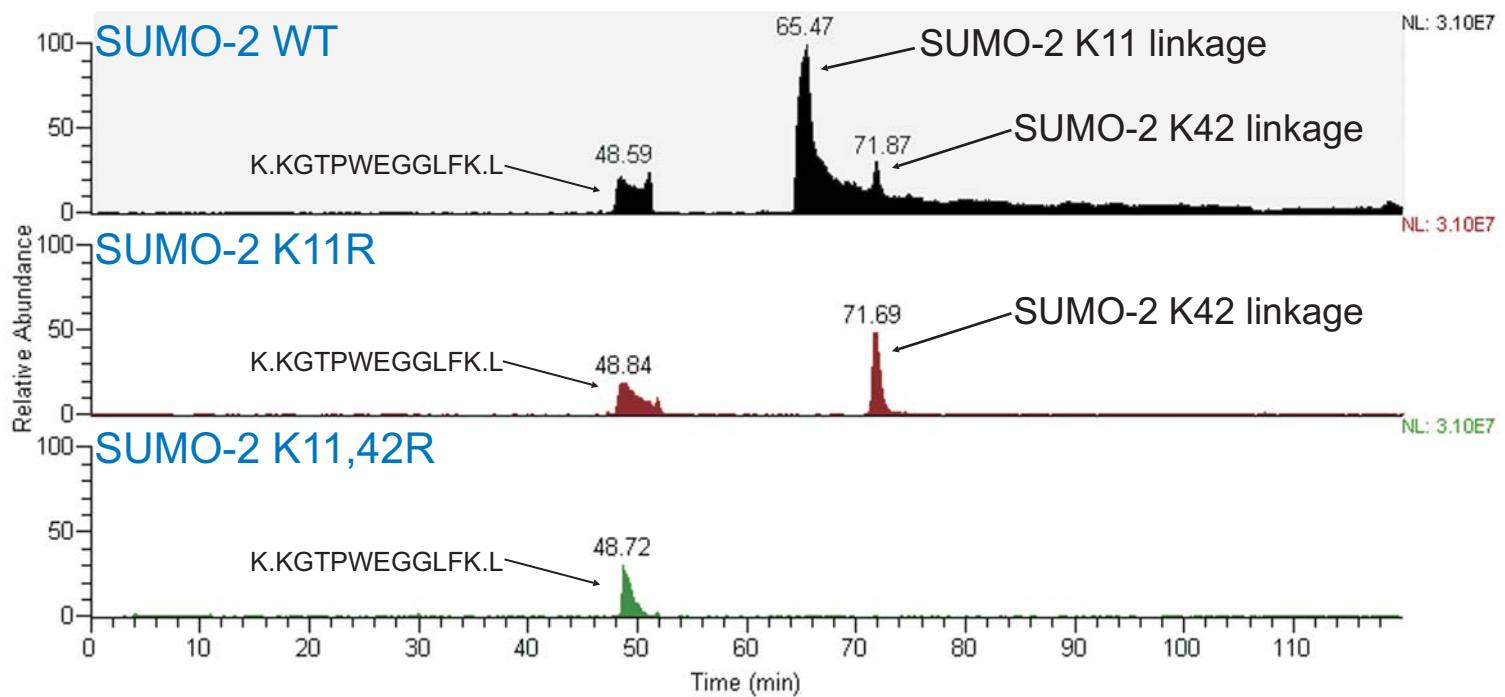
B

Scan	Charge	relPos	modScore	precursor mass	matched ions	TIC	%TIC	BPI
4873	4	1	1	1053.61	108	3.05424e+06	0.663465	120069

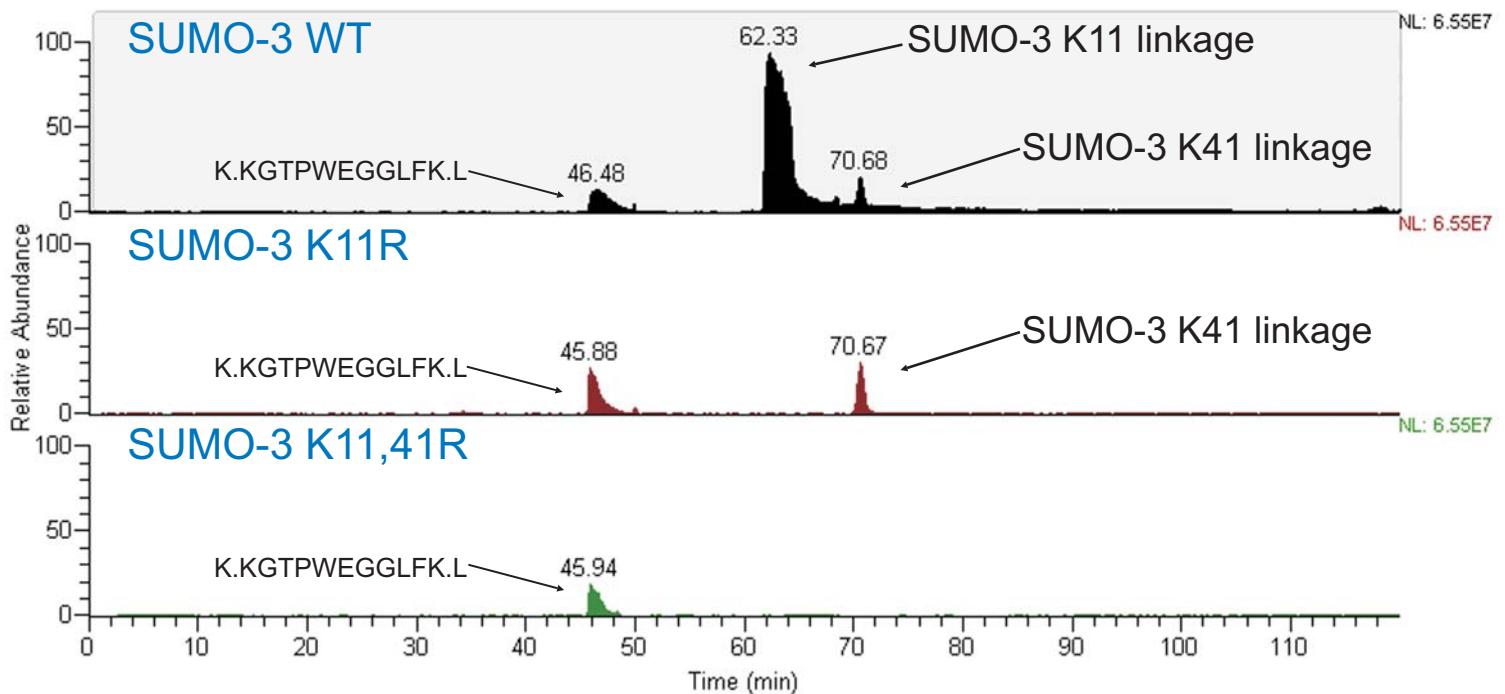


Supplemental Figure 2

A

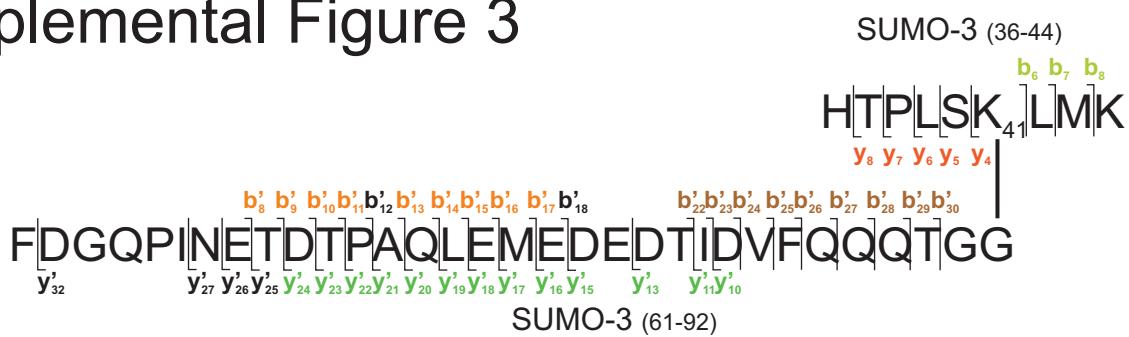


B

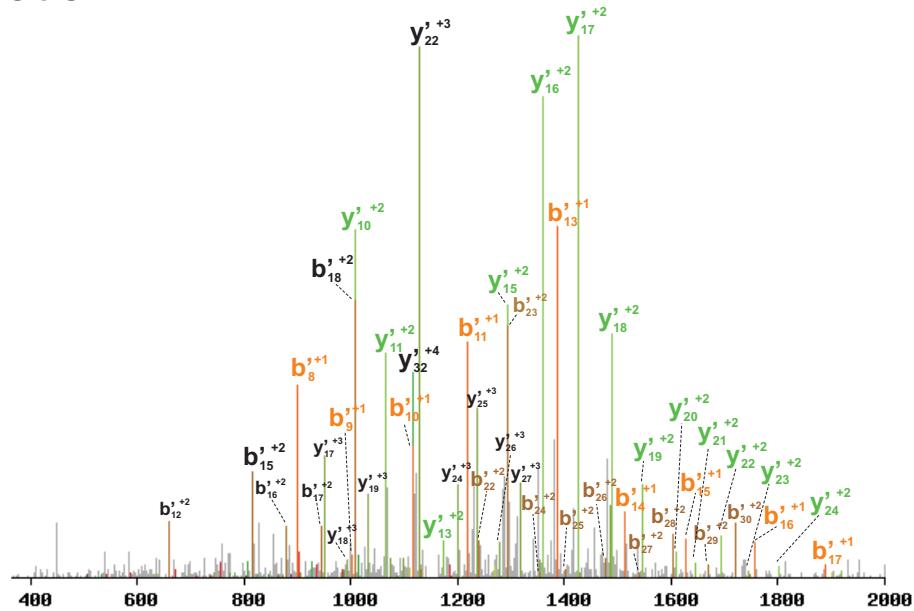


Supplemental Figure 3

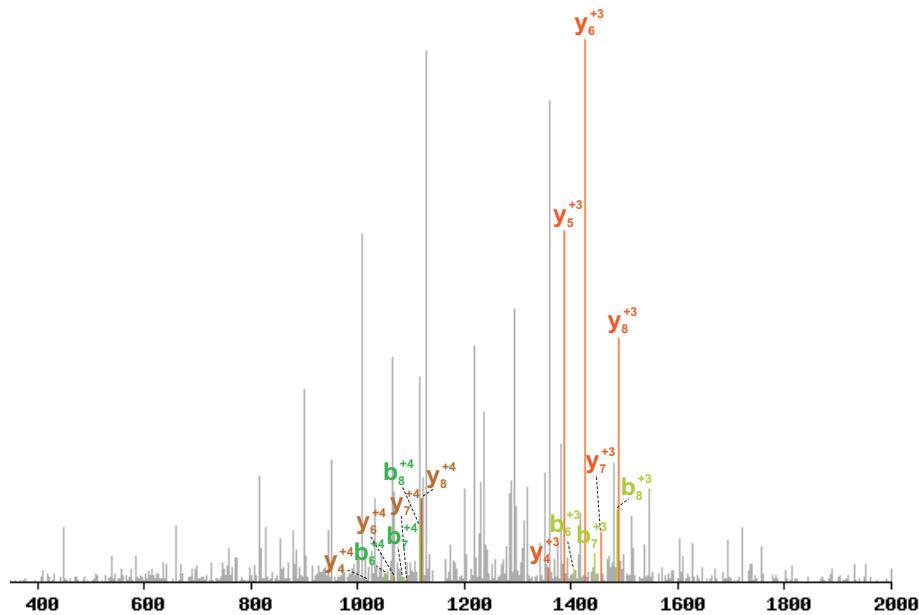
A



B Modification

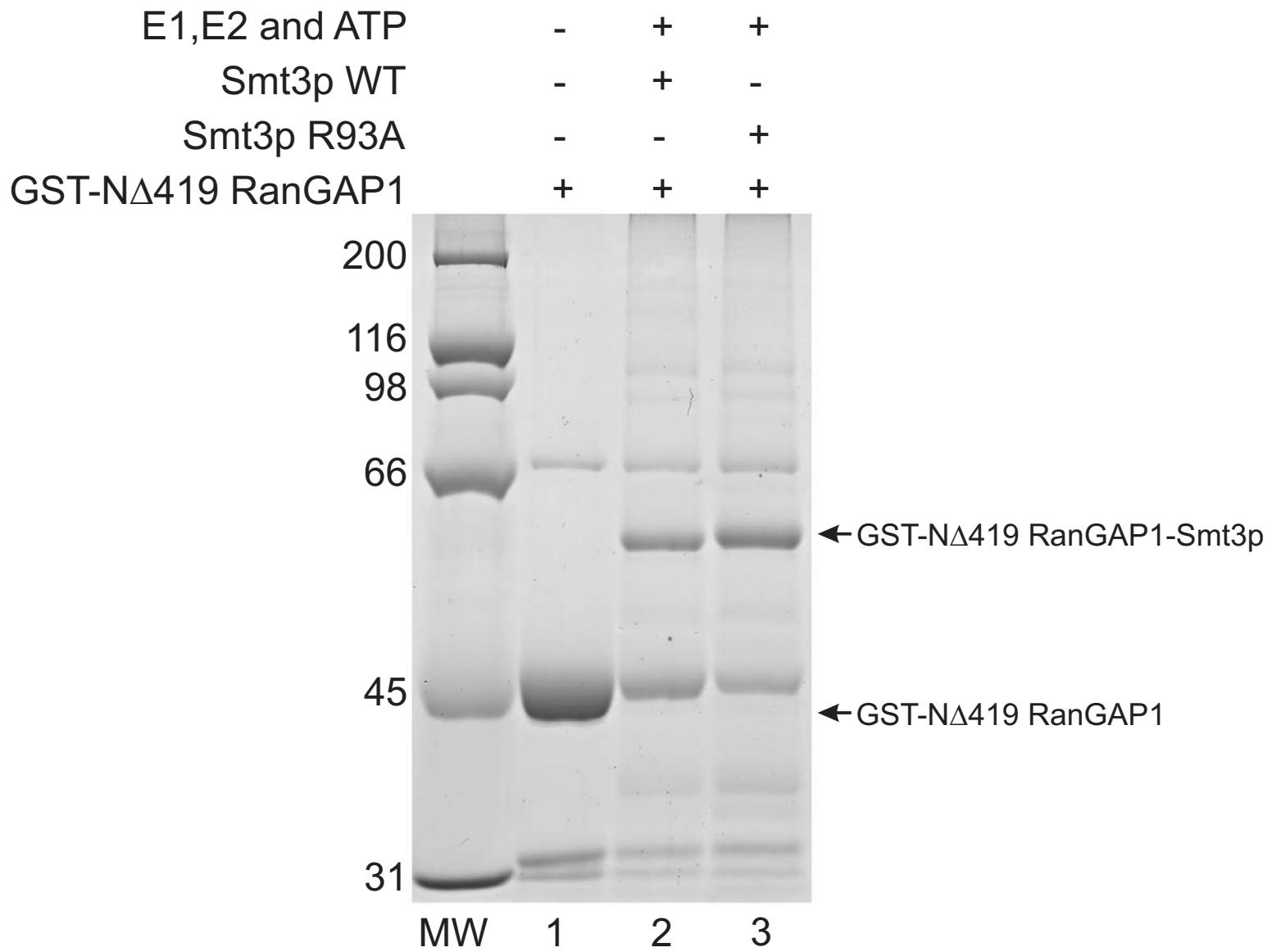


C Target



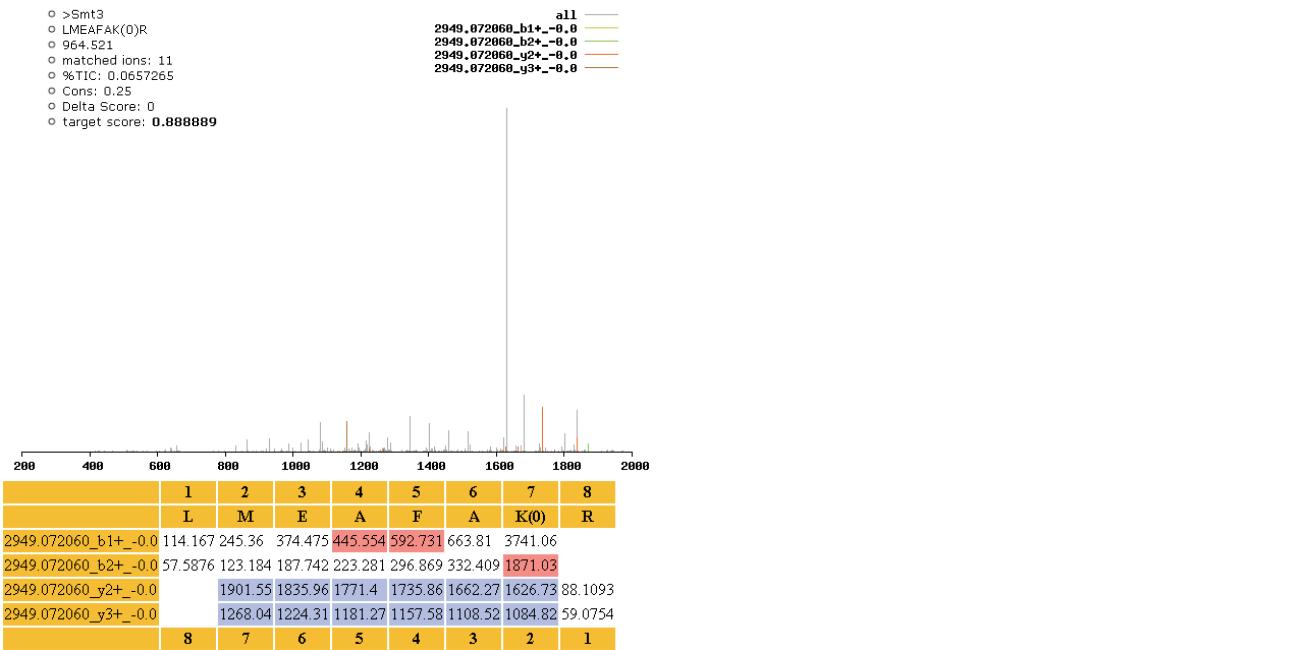
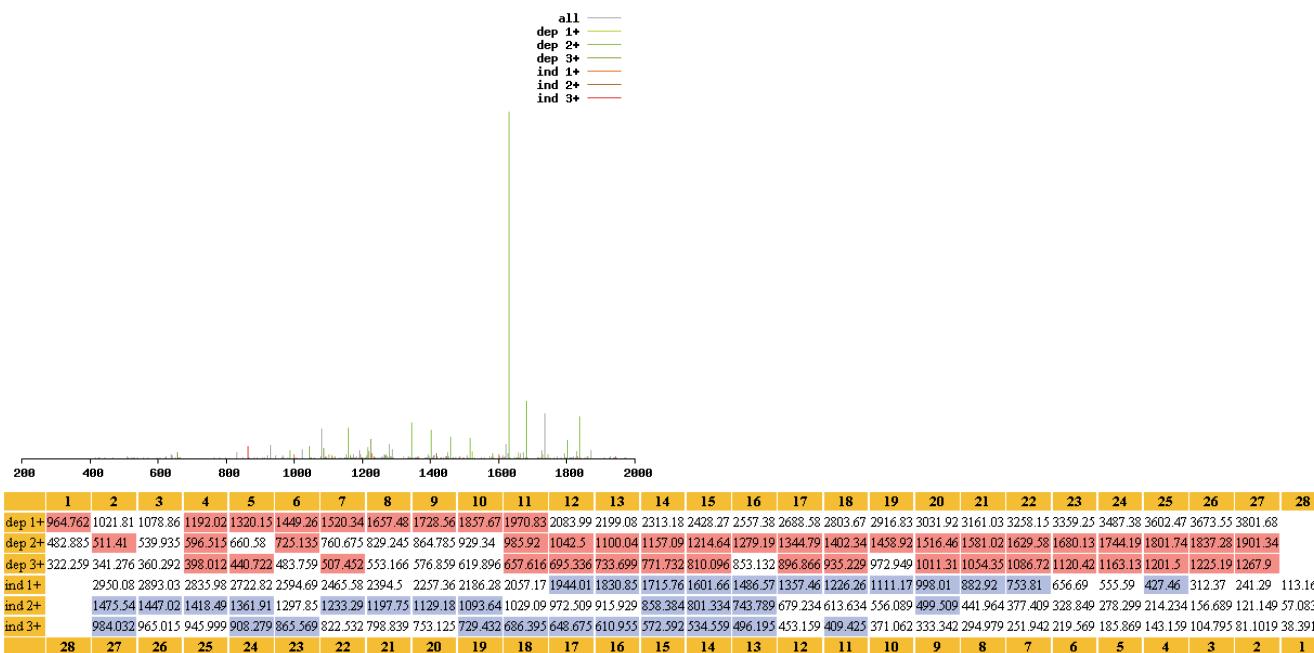
Supplemental Figure 4

A



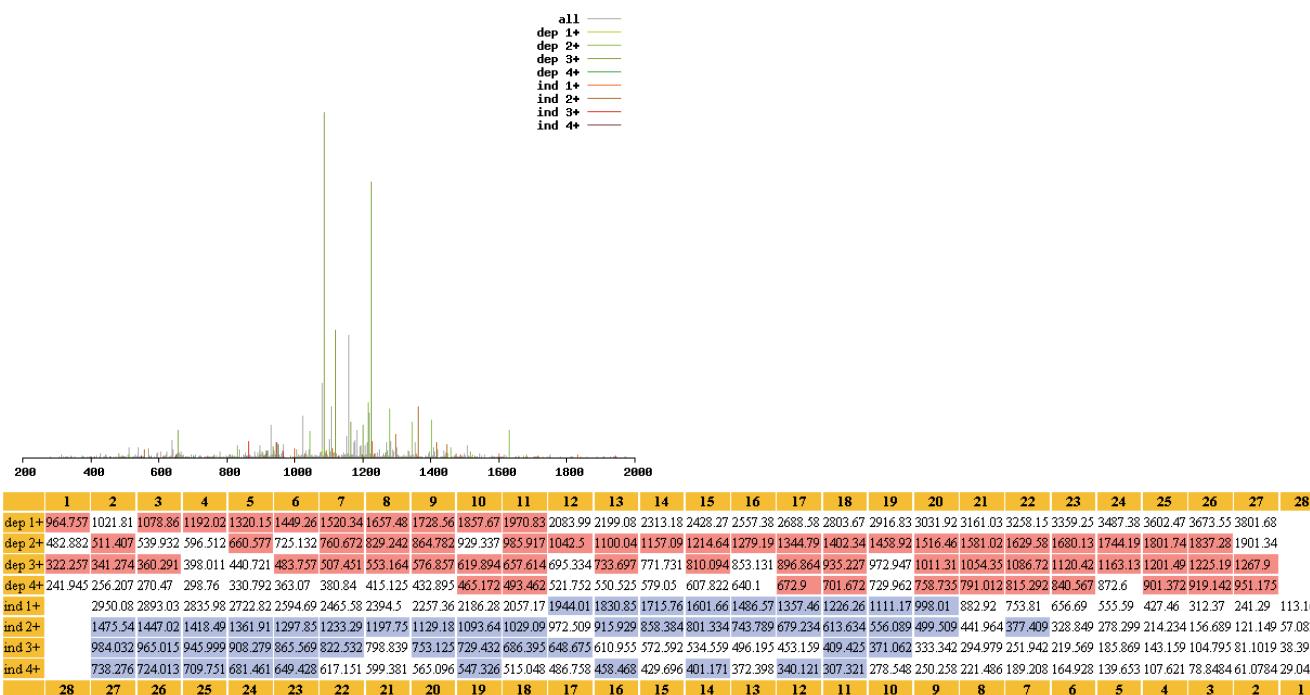
B

Scan	Charge	relPos	modScore	precursor mass	matched ions	TIC	%TIC	BPI
3989	3	1	1	964.527	70	190259	0.461746	39681.3

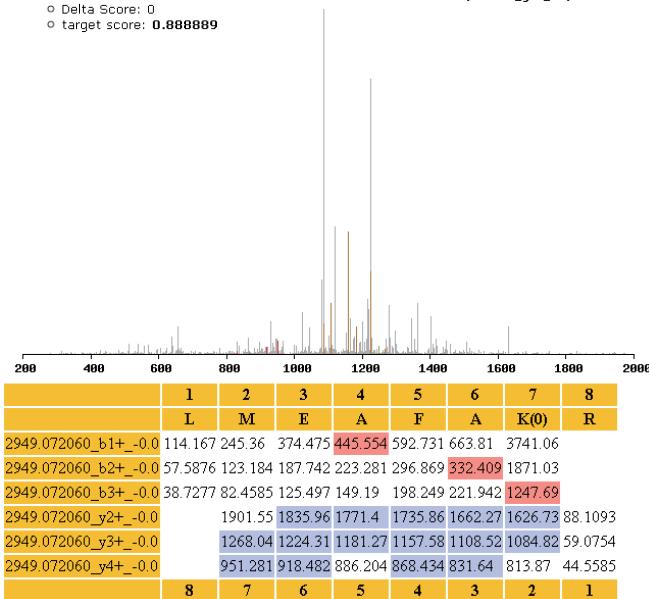


C

Scan	Charge	relPos	modScore	precursor mass	matched ions	TIC	%TIC	BPI
4035	4	1	1	964.522	97	928687	0.36897	81150.6



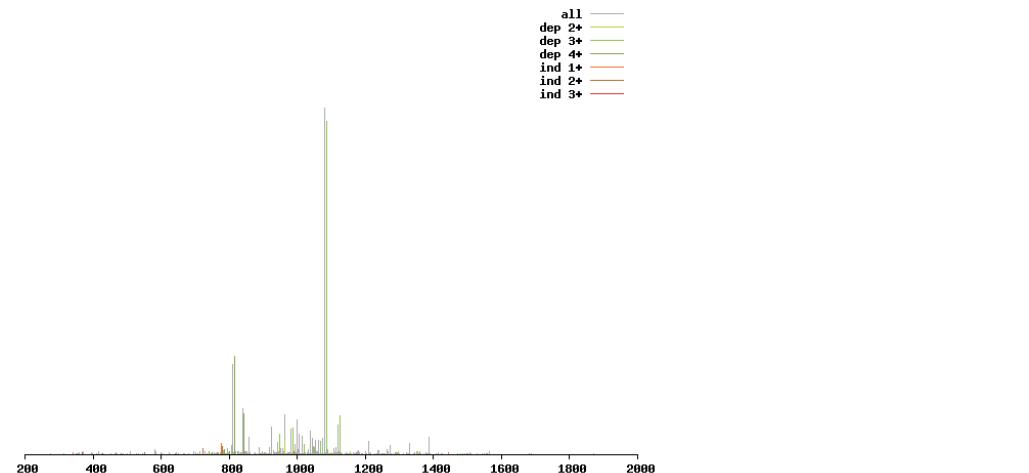
- ```
o >Smt3 all
o MLEAFAK(0)R 2949,072060_b1+-,0.0
o 964_521 2949,072060_b2+-,0.0
o matched ions: 15 2949,072060_b3+-,0.0
o %TIC: 0.088307 2949,072060_y1+-,0.0
o Cons: 0.166667 2949,072060_y3+-,0.0
o Delta Score: 0 2949,072060_y4+-,0.0
o target score: 0.8888889
```



# Supplemental Figure 5

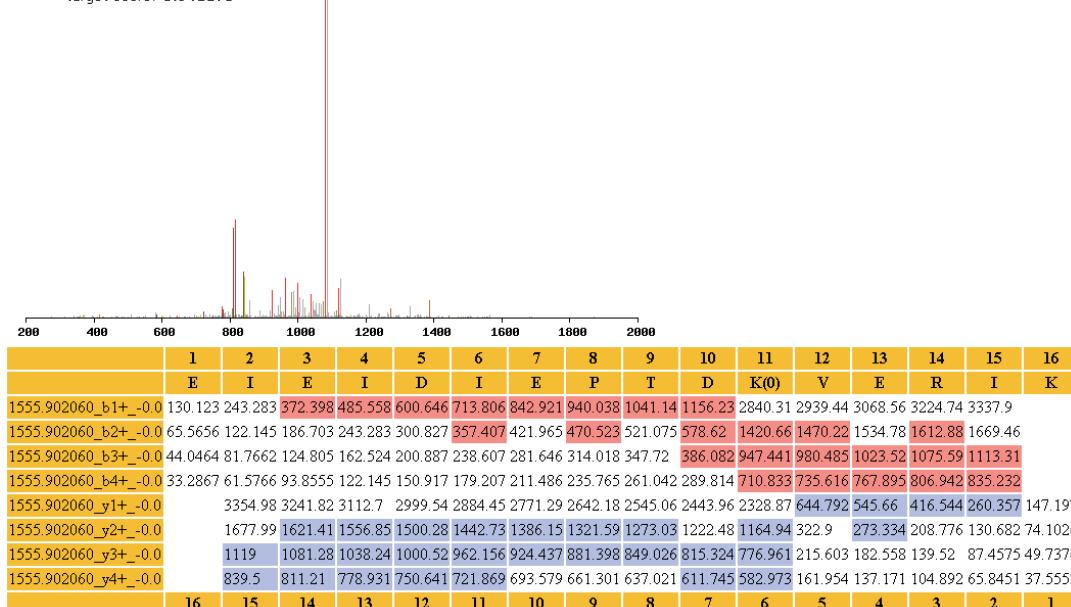
A

| Scan | Charge | relPos | modScore | precursor mass | matched ions | TIC    | %TIC     | BPI     |
|------|--------|--------|----------|----------------|--------------|--------|----------|---------|
| 8215 | 4      | 1      | 0.971888 | 1926.01        | 51           | 254378 | 0.287173 | 36952.9 |



|        | 1       | 2       | 3       | 4       | 5       | 6       | 7       | 8       | 9       | 10      | 11      | 12      | 13      | 14      | 15      | 16      | 17 |
|--------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|----|
| dep 2+ | 964.04  | 992.565 | 1021.09 | 1099.18 | 1155.76 | 1191.3  | 1247.88 | 1297.45 | 1354.03 | 1422.6  | 1479.18 | 1528.74 | 1572.28 | 1600.81 | 1629.33 | 1685.91 |    |
| dep 3+ | 643.029 | 662.046 | 681.062 | 733.126 | 770.846 | 794.539 | 832.259 | 865.302 | 903.022 | 948.736 | 986.456 | 1019.5  | 1048.53 | 1067.54 | 1086.56 | 1124.28 |    |
| dep 4+ | 482.524 | 496.786 | 511.049 | 550.096 | 578.386 | 596.156 | 624.446 | 649.229 | 677.519 | 711.804 | 740.094 | 764.876 | 786.646 | 800.909 | 815.171 | 843.461 |    |
| ind 1+ | 1556.91 | 1499.86 | 1442.81 | 1286.62 | 1173.46 | 1102.38 | 989.22  | 890.09  | 776.93  | 639.79  | 526.63  | 427.5   | 340.42  | 283.37  | 226.32  | 113.16  |    |
| ind 2+ | 778.959 | 750.434 | 721.909 | 643.814 | 587.234 | 551.694 | 495.114 | 445.549 | 388.969 | 320.399 | 263.819 | 214.254 | 170.714 | 142.189 | 113.664 | 57.0839 |    |
| ind 3+ | 519.642 | 500.625 | 481.609 | 429.545 | 391.825 | 368.132 | 330.412 | 297.369 | 259.649 | 213.935 | 176.215 | 143.172 | 114.145 | 95.1286 | 76.1119 | 38.3919 |    |
|        | 17      | 16      | 15      | 14      | 13      | 12      | 11      | 10      | 9       | 8       | 7       | 6       | 5       | 4       | 3       | 2       | 1  |

- o >Q15843|HUMAN Nedd8
- o EIEIDIEPTDK(0)VERIK
- o 1926.04
- o matched ions: 48
- o %TIC: 0.388307
- o Cons: 0.125
- o Delta Score: 0
- o target score: **0.941176**

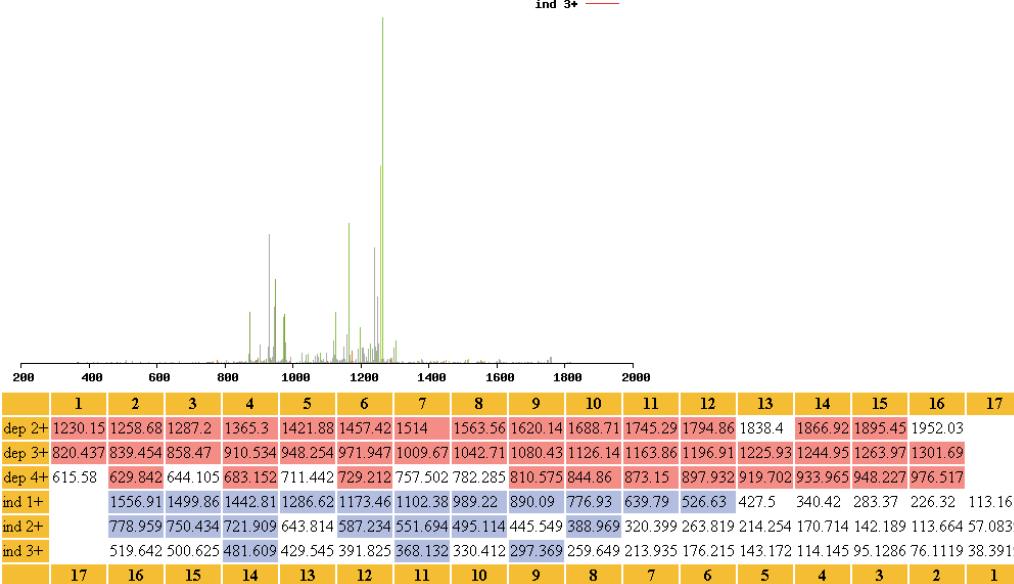


|                     | 1         | 2       | 3       | 4       | 5       | 6       | 7       | 8       | 9       | 10      | 11      | 12      | 13      | 14      | 15      | 16 |
|---------------------|-----------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|----|
| E                   | I         | E       | I       | D       | I       | E       | P       | T       | D       | K(0)    | V       | E       | R       | I       | K       |    |
| 1555.902060_b1+-0.0 | 130.123   | 243.283 | 372.398 | 485.558 | 600.646 | 713.806 | 842.921 | 940.038 | 1041.14 | 1156.23 | 2840.31 | 2939.44 | 3068.56 | 3224.74 | 3337.9  |    |
| 1555.902060_b2+-0.0 | 65.5656   | 122.145 | 186.703 | 243.283 | 300.827 | 357.407 | 421.965 | 470.523 | 521.075 | 578.62  | 1420.66 | 1470.22 | 1534.78 | 1612.88 | 1669.46 |    |
| 1555.902060_b3+-0.0 | 0.44.0464 | 81.7662 | 124.805 | 162.524 | 200.887 | 238.607 | 281.646 | 314.18  | 347.72  | 386.082 | 947.441 | 980.485 | 1023.52 | 1075.59 | 1113.31 |    |
| 1555.902060_b4+-0.0 | 33.2867   | 61.5766 | 93.8555 | 122.145 | 150.917 | 179.207 | 211.486 | 235.765 | 261.042 | 289.814 | 710.833 | 735.616 | 767.895 | 806.942 | 835.232 |    |
| 1555.902060_y1+-0.0 | 3354.98   | 3241.82 | 3112.7  | 2999.54 | 2884.45 | 2771.29 | 2642.18 | 2545.06 | 2443.96 | 2328.87 | 644.792 | 545.66  | 416.544 | 260.357 | 147.197 |    |
| 1555.902060_y2+-0.0 | 1677.99   | 1621.41 | 1556.85 | 1500.28 | 1442.73 | 1388.15 | 1321.59 | 1273.03 | 1222.48 | 1164.94 | 322.9   | 273.334 | 208.776 | 130.682 | 74.1026 |    |
| 1555.902060_y3+-0.0 | 1119      | 1081.28 | 1038.24 | 1000.52 | 962.156 | 924.437 | 881.398 | 849.026 | 815.324 | 776.961 | 215.603 | 182.558 | 139.52  | 87.4575 | 49.7376 |    |
| 1555.902060_y4+-0.0 | 839.5     | 811.21  | 778.931 | 750.641 | 721.869 | 693.579 | 661.301 | 637.021 | 611.745 | 582.973 | 161.954 | 137.171 | 104.892 | 65.8451 | 37.5552 |    |
|                     | 16        | 15      | 14      | 13      | 12      | 11      | 10      | 9       | 8       | 7       | 6       | 5       | 4       | 3       | 2       | 1  |

B

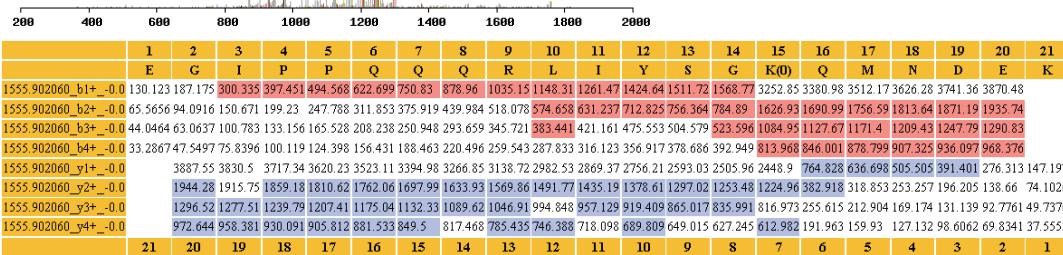
| Scan | Charge | relPos | modScore | precursor mass | matched ions | TIC    | %TIC     | BPI    |
|------|--------|--------|----------|----------------|--------------|--------|----------|--------|
| 8331 | 4      | 1      | 0.977332 | 2458.24        | 48           | 961903 | 0.418254 | 114176 |

all  
dep 2+  
dep 3+  
dep 4+  
ind 1+  
ind 2+  
ind 3+



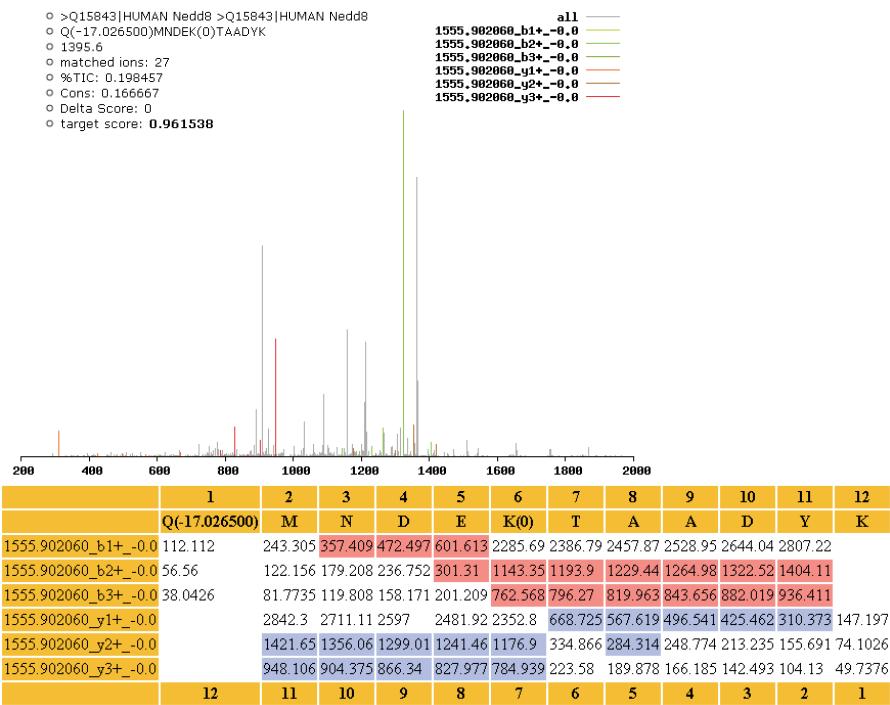
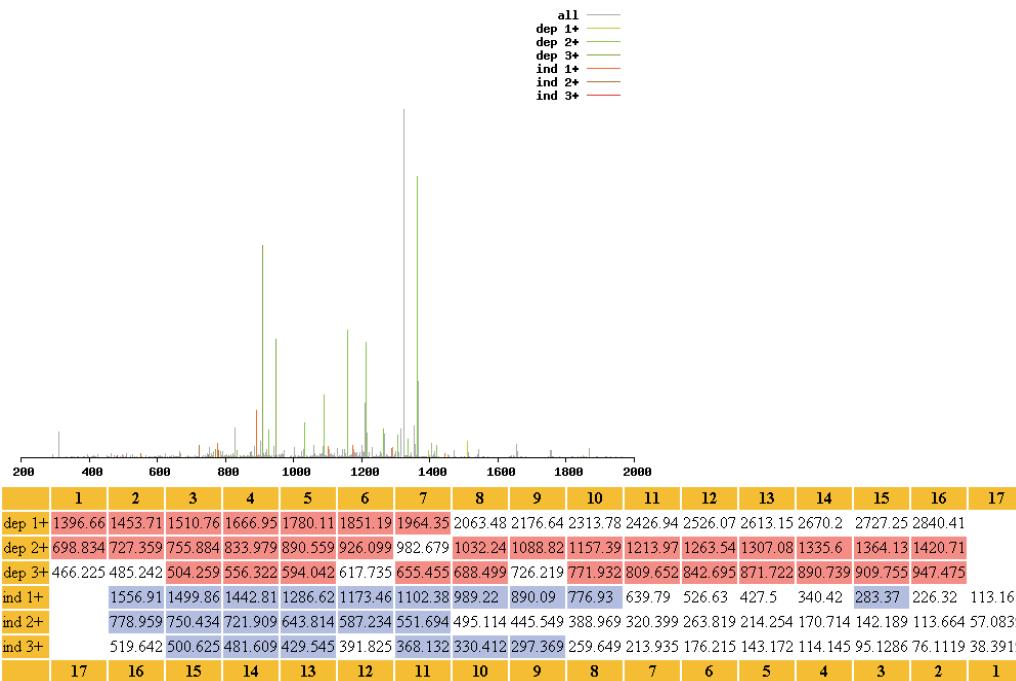
o >Q15843|HUMAN Nedd8 >Q15843|HUMAN Nedd8  
o EGIPQQQRLLYSGK(D)QMNEK  
o 2458.23  
o matched ions: 65  
o %TIC: 0.17934  
o Cons: 0.125  
o Delta Score: 0  
o target score: 0.977273

all  
1555.902060\_b1+-0.0  
1555.902060\_b2+-0.0  
1555.902060\_b3+-0.0  
1555.902060\_b4+-0.0  
1555.902060\_y1+-0.0  
1555.902060\_y2+-0.0  
1555.902060\_y3+-0.0  
1555.902060\_y4+-0.0



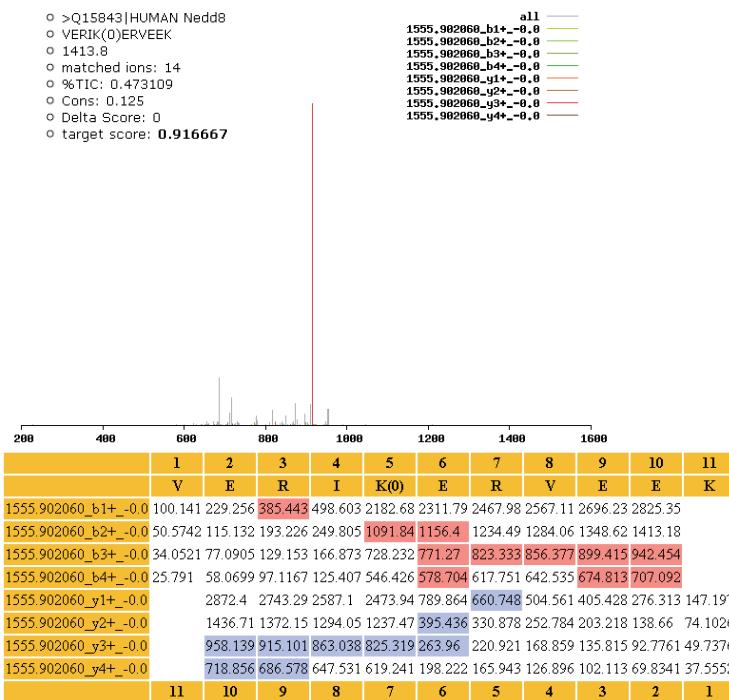
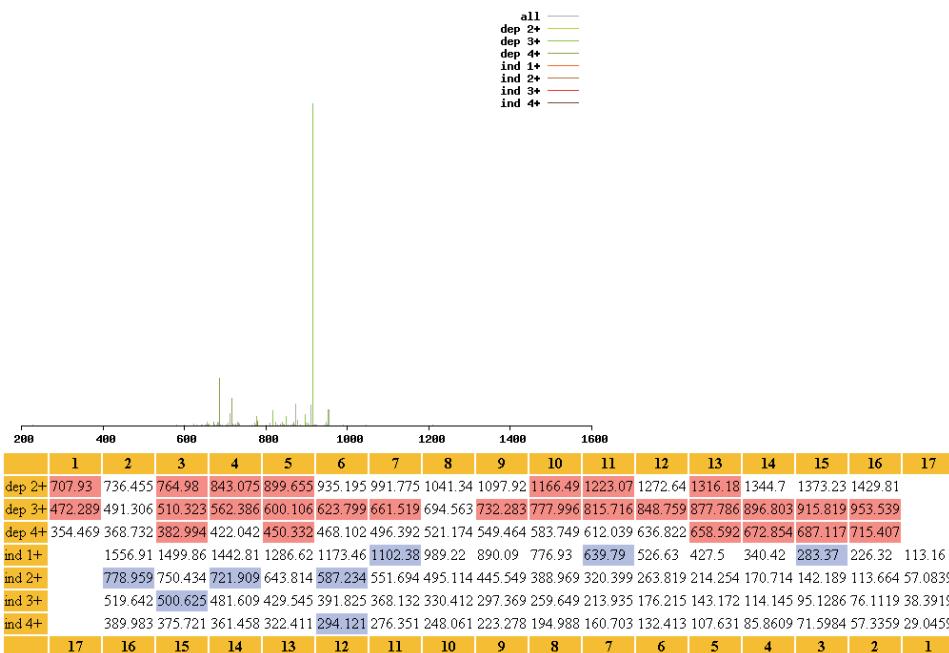
C

| Scan | Charge | relPos | modScore | precursor mass | matched ions | TIC    | %TIC     | BPI     |
|------|--------|--------|----------|----------------|--------------|--------|----------|---------|
| 8954 | 3      | 1      | 0.979072 | 1395.6         | 52           | 330173 | 0.386181 | 32915.6 |



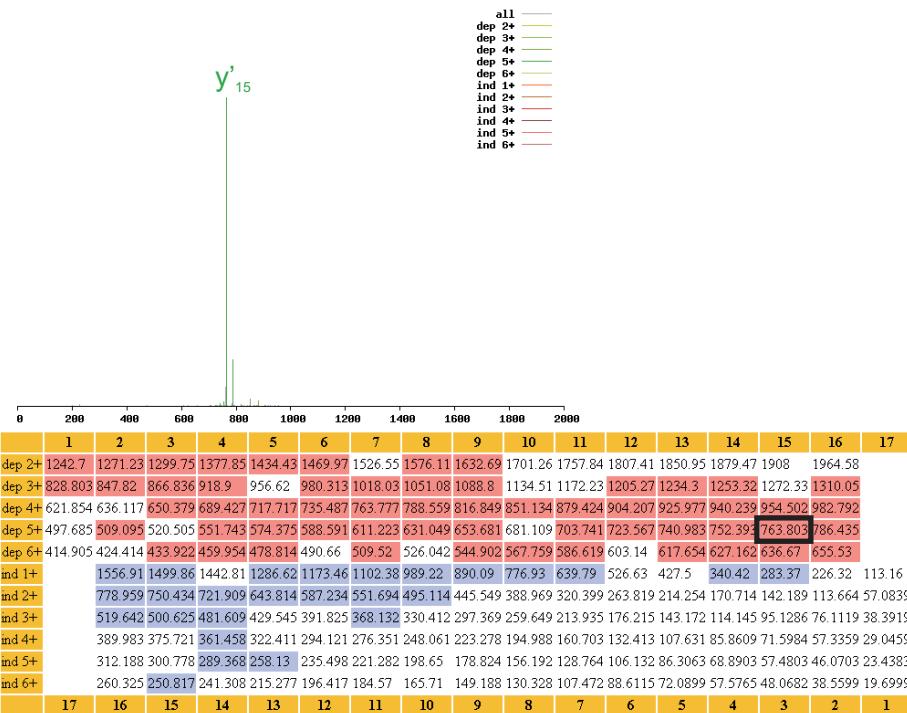
D

| Scan | Charge | relPos | modScore | precursor mass | matched ions | TIC    | %TIC     | BPI     |
|------|--------|--------|----------|----------------|--------------|--------|----------|---------|
| 6881 | 4      | 1      | 1        | 1413.79        | 31           | 186201 | 0.682315 | 83563.5 |

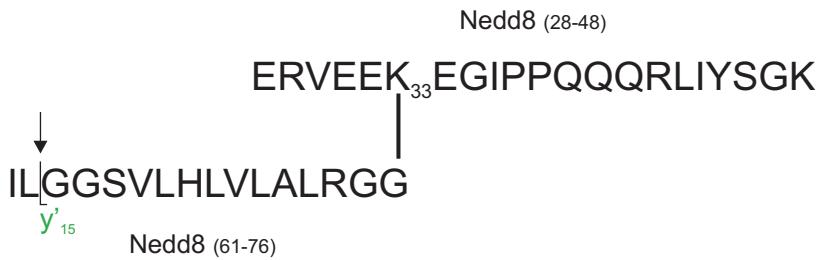


E

| Scan | Charge | relPos | modScore | precursor mass | matched ions | TIC    | %TIC    | BPI    |
|------|--------|--------|----------|----------------|--------------|--------|---------|--------|
| 7887 | 6      | 1      | 1        | 2483.33        | 56           | 614857 | 1.39409 | 385186 |

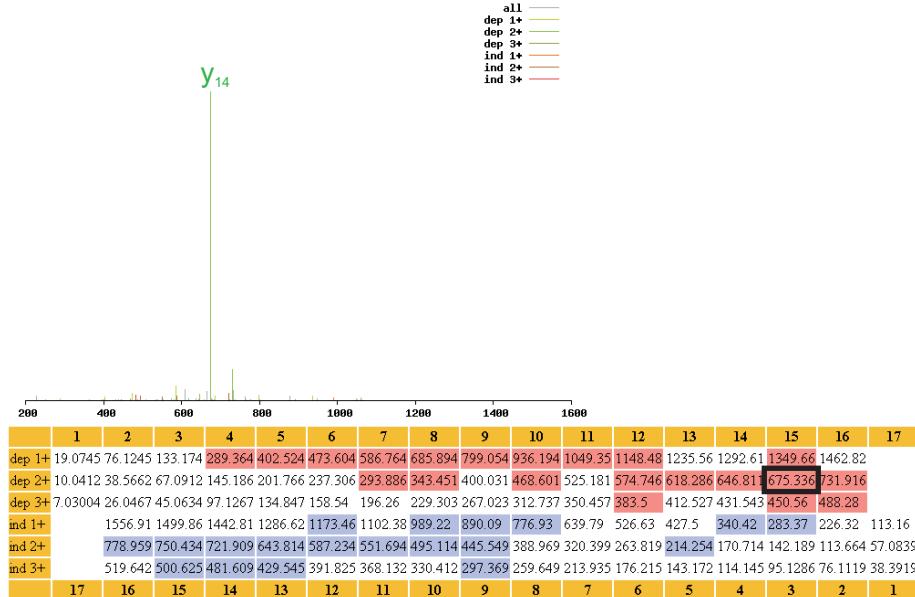


NEDD8 K33 linkage:



F

| Scan  | Charge | relPos | modScore | precursor mass | matched ions | TIC    | %TIC     | BPI    |
|-------|--------|--------|----------|----------------|--------------|--------|----------|--------|
| 13273 | 3      | 1      | 0.983017 | 18.0163        | 29           | 341325 | 0.761792 | 193355 |

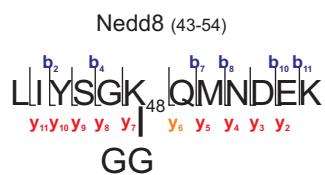


Unconjugated NEDD8 tail:

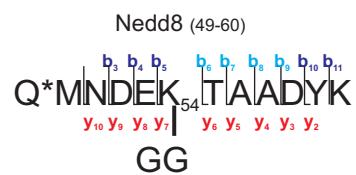


# Supplemental Figure 6

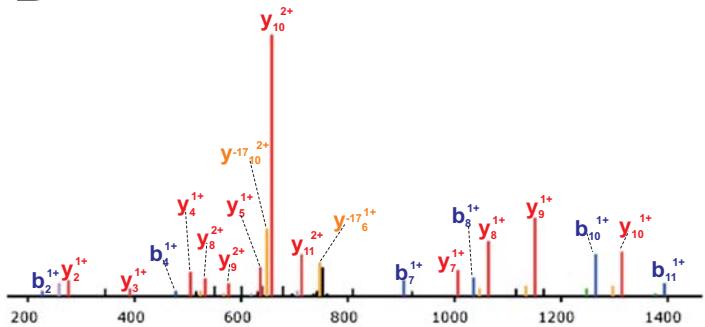
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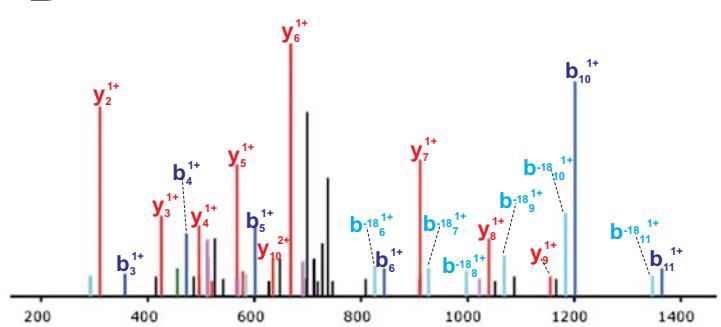
C



B

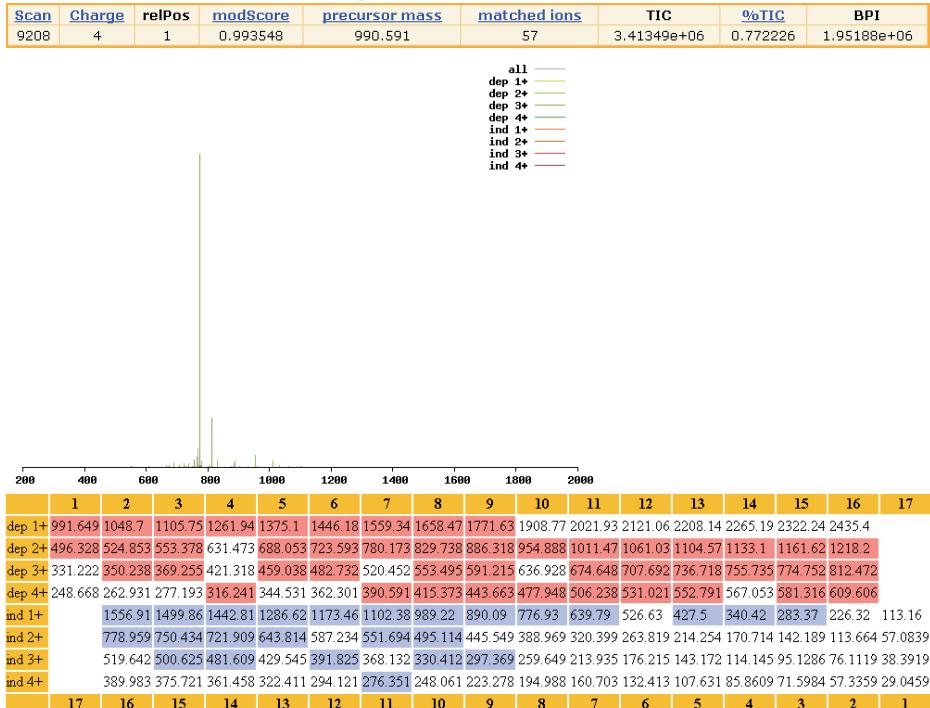


D

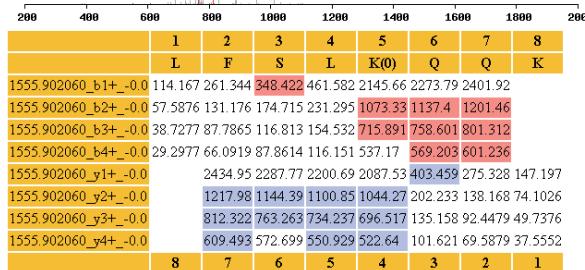


# Supplemental Figure 7

A

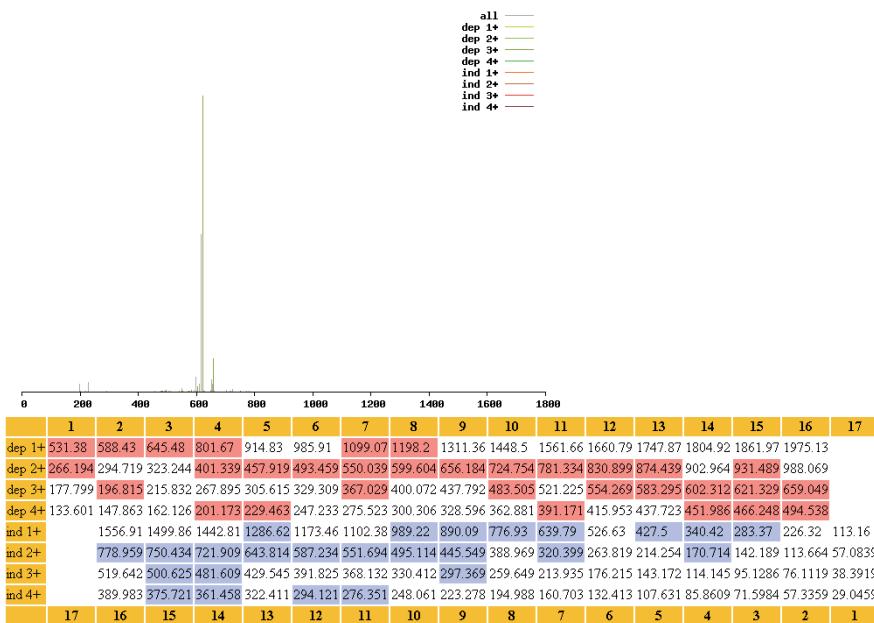


- o >P61001|UbcH12 E2M Nedd8 E2
  - o LFSLK(O)QQK
  - o 990.591
  - o matched ions: 13
  - o %TIC: 0.114867
  - o Cons: 0.125
  - o Delta Score: 0
  - o target score: 0.888889
- 
- |                     | 1       | 2       | 3       | 4       | 5       | 6       | 7       | 8 |
|---------------------|---------|---------|---------|---------|---------|---------|---------|---|
|                     | L       | F       | S       | L       | K(O)    | Q       | Q       | K |
| 1555.902060_b1+-0.0 | 114.167 | 261.344 | 348.422 | 461.582 | 2145.66 | 2273.79 | 2401.92 |   |
| 1555.902060_b2+-0.0 | 57.5876 | 131.176 | 174.715 | 231.295 | 1073.33 | 1137.4  | 1201.46 |   |
| 1555.902060_b3+-0.0 | 38.7277 | 87.7865 | 116.813 | 154.532 | 715.891 | 758.601 | 801.312 |   |
| 1555.902060_b4+-0.0 | 29.2977 | 66.0919 | 87.8614 | 116.151 | 537.17  | 569.203 | 601.236 |   |
| 1555.902060_y1+-0.0 | 2434.95 | 2287.77 | 2200.69 | 2087.53 | 403.459 | 275.328 | 147.197 |   |
| 1555.902060_y2+-0.0 | 1217.98 | 1144.39 | 1100.85 | 1044.27 | 202.233 | 138.168 | 74.1026 |   |
| 1555.902060_y3+-0.0 | 812.322 | 763.263 | 734.237 | 696.517 | 135.158 | 92.4479 | 49.7376 |   |
| 1555.902060_y4+-0.0 | 609.493 | 572.699 | 550.929 | 522.64  | 101.621 | 69.5879 | 37.5552 |   |
|                     | 8       | 7       | 6       | 5       | 4       | 3       | 2       | 1 |

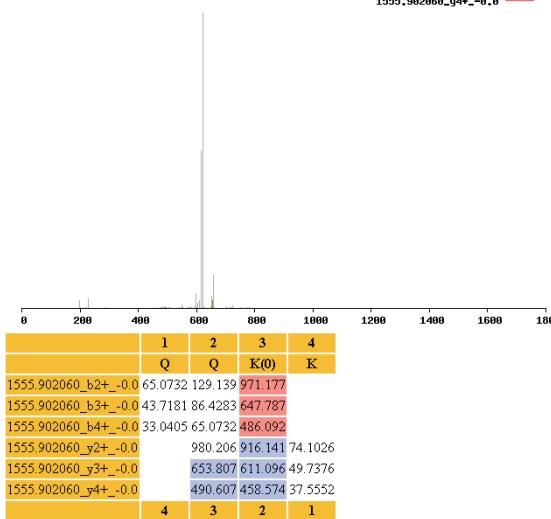


B

| Scan | Charge | relPos | modScore | precursor mass | matched ions | TIC    | %TIC     | BPI    |
|------|--------|--------|----------|----------------|--------------|--------|----------|--------|
| 7981 | 4      | 1      | 0.972805 | 530.322        | 38           | 632102 | 0.546491 | 293220 |

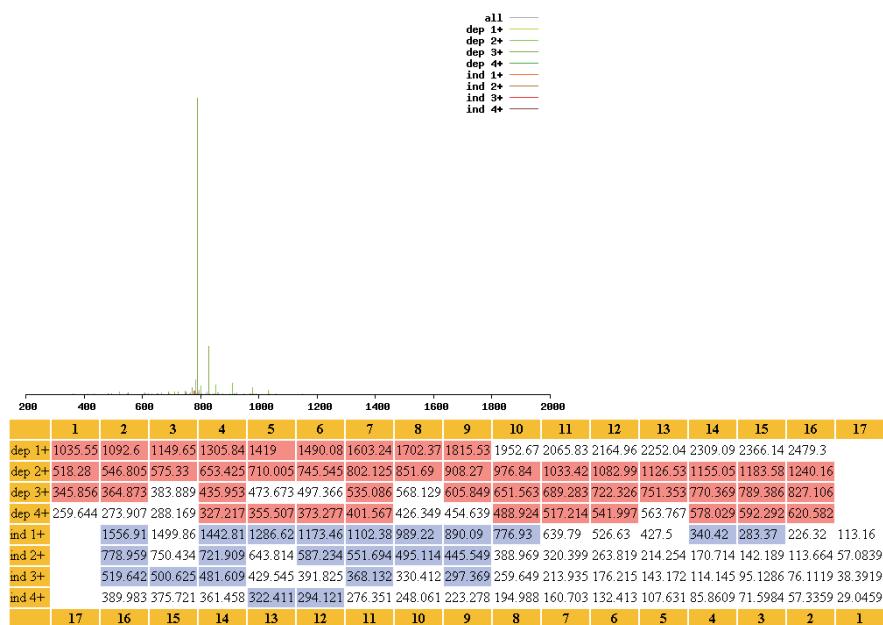


- o >P61081|UbcH12 E2M Nedd8 E2
  - o QQK(O)K
  - o 530.322
  - o matched ions: 6
  - o %TIC: 0.0136005
  - o Cons: 0.166657
  - o Delta Score: 0
  - o target score: 0.8
- all  
1555.902060\_b2+-0.0  
1555.902060\_b3+-0.0  
1555.902060\_b4+-0.0  
1555.902060\_y2+-0.0  
1555.902060\_y3+-0.0  
1555.902060\_y4+-0.0

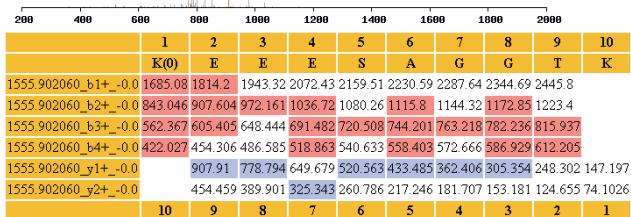


C

| Scan | Charge | relPos | modScore | precursor mass | matched ions | TIC    | %TIC     | BPI    |
|------|--------|--------|----------|----------------|--------------|--------|----------|--------|
| 7952 | 4      | 1      | 0.981308 | 1034.49        | 56           | 562157 | 0.719293 | 295007 |

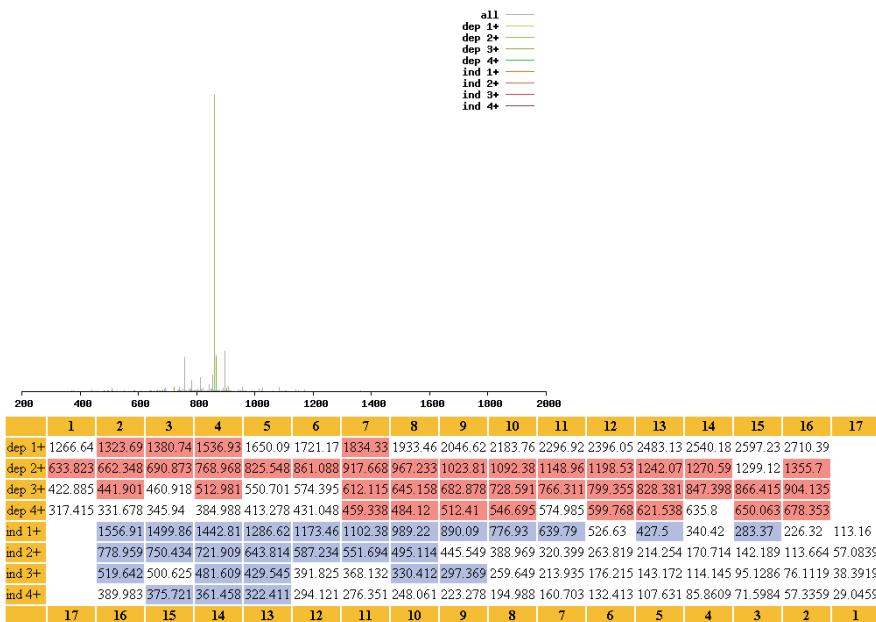


- o >P61081|UbcH12 E2M Nedd8 E2
- o K(D)EESAGGT
- o 1024.49
- o matched ions: 20
- o %TIC: 0.0536365
- o Cons: 0.166667
- o Delta Score: 0
- o target score: **0.909091**
- o 1555.902060\_b1+-0.0
- o 1555.902060\_b2+-0.0
- o 1555.902060\_b3+-0.0
- o 1555.902060\_b4+-0.0
- o 1555.902060\_y1+-0.0
- o 1555.902060\_y2+-0.0



D

| Scan | Charge | relPos | modScore | precursor mass | matched ions | TIC    | %TIC    | BPI    |
|------|--------|--------|----------|----------------|--------------|--------|---------|--------|
| 8029 | 4      | 1      | 0.922825 | 1265.58        | 37           | 449434 | 0.47149 | 176930 |

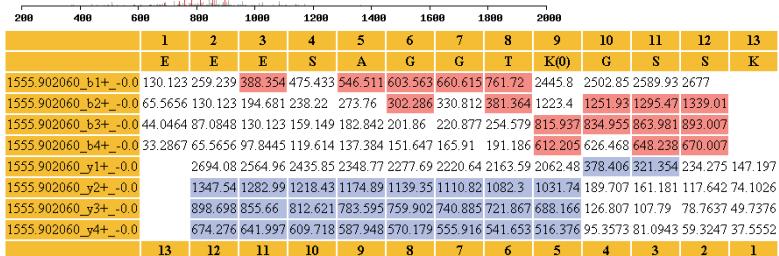


○ >P61081|Ubch12 E2M Nedd8 E2  
○ EEEESAGGTTK(0)GSSK  
○ 1265.58  
○ matched ions: 32  
○ %TIC: 0.176361  
○ Cons: 0.125  
○ Delta Score: 0  
○ target score: **0.928571**

Mass spectrum showing relative intensity versus m/z. The x-axis ranges from 200 to 2000. A prominent peak is at m/z 1265.58. The y-axis shows relative intensity from 0 to 100.

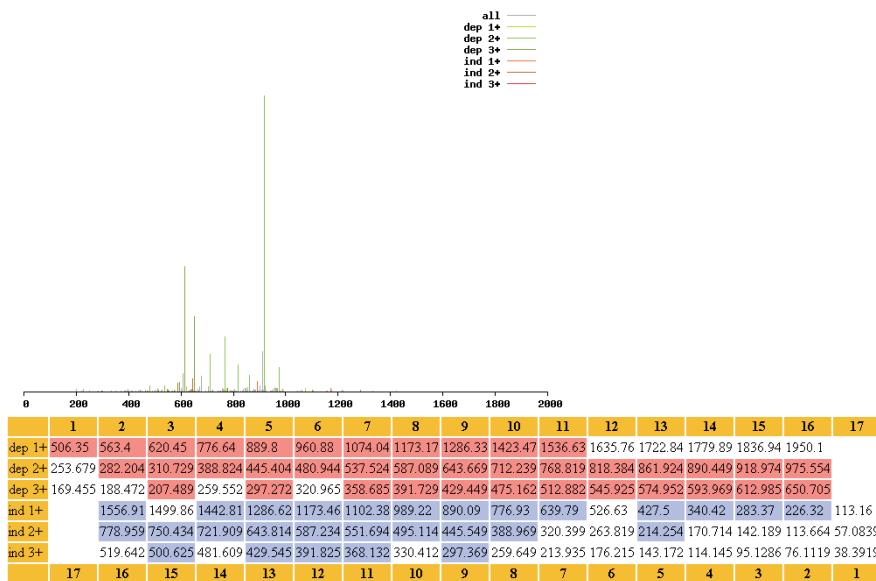
Legend for ion types:

- all
- 1555.902060\_b1+-0.0
- 1555.902060\_b2+-0.0
- 1555.902060\_b3+-0.0
- 1555.902060\_b4+-0.0
- 1555.902060\_y1+-0.0
- 1555.902060\_y2+-0.0
- 1555.902060\_y3+-0.0
- 1555.902060\_y4+-0.0

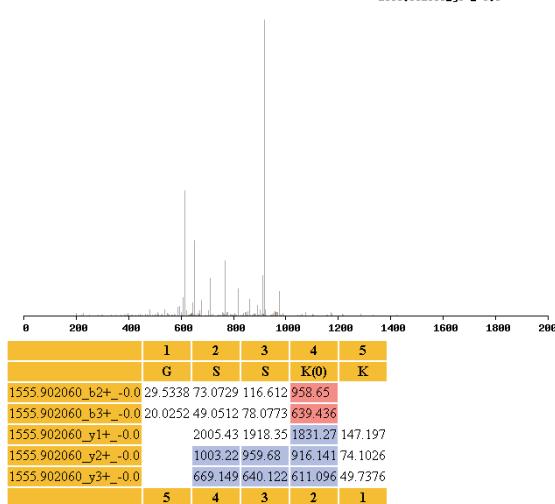


E

| Scan | Charge | relPos | modScore | precursor mass | matched ions | TIC    | %TIC     | BPI    |
|------|--------|--------|----------|----------------|--------------|--------|----------|--------|
| 7965 | 3      | 1      | 1        | 505.292        | 58           | 584666 | 0.696372 | 148405 |

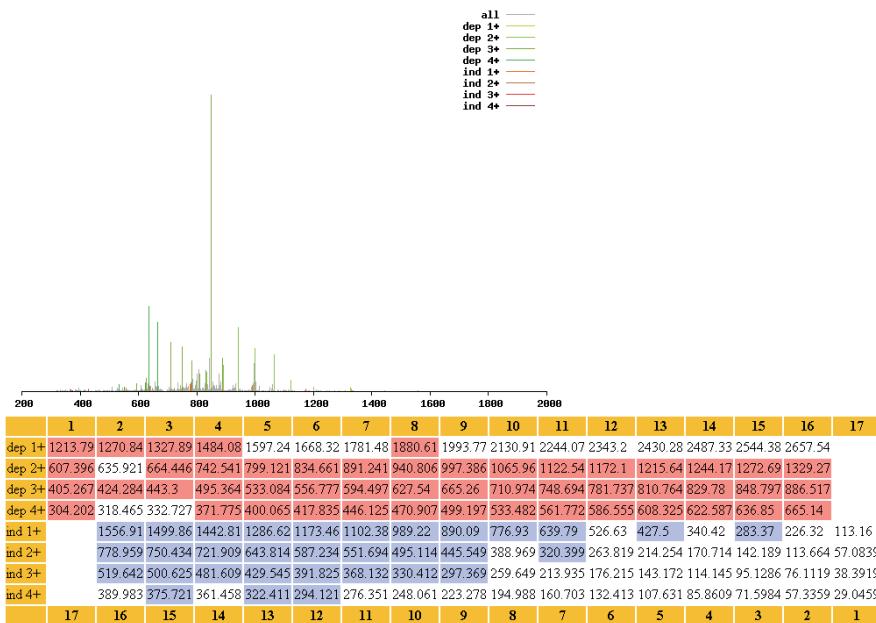


o >P61081|UbcH12 E2M Nedd8 E2  
 o GSSK(0)K  
 o S05.291  
 o matched ions: 3  
 o %TIC: 0.00238028  
 o Cons: 0.2  
 o Delta Score: 0  
 o target score: 0  
 o %TIC: 0.00238028  
 o Cons: 0.2  
 o Delta Score: 0  
 o target score: 0

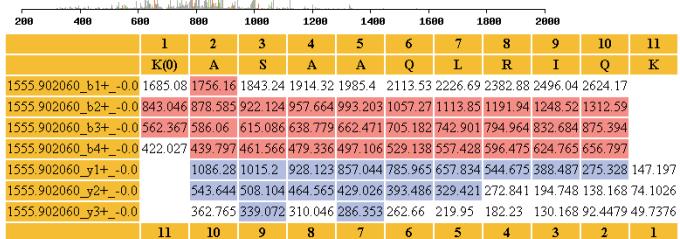


F

| Scan | Charge | relPos | modScore | precursor mass | matched ions | TIC    | %TIC     | BPI     |
|------|--------|--------|----------|----------------|--------------|--------|----------|---------|
| 7534 | 4      | 1      | 1        | 1212.73        | 75           | 432025 | 0.507157 | 61270.5 |



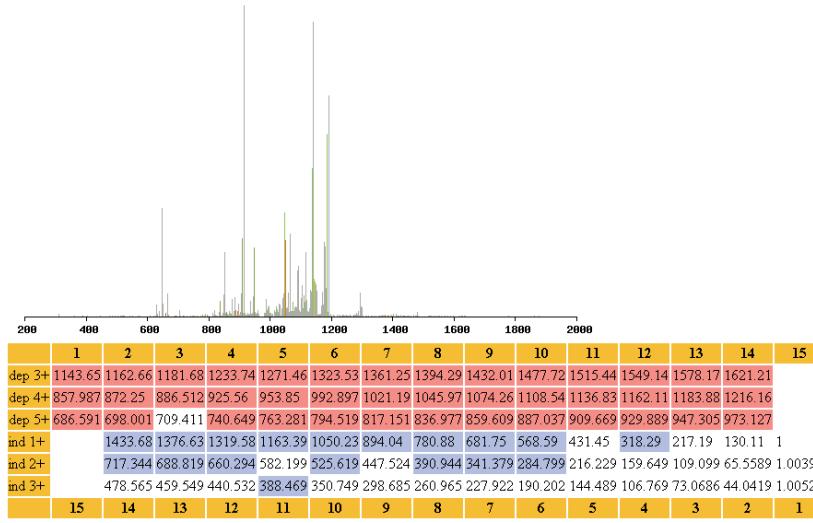
o >P61081|UbchH12 E2M Nedd8 E2  
 o K(0)ASAAQLRIQK  
 o 1212.73  
 o matched ions: 44  
 o %TIC: 0.0819807  
 o Cons: 0.142857  
 o Delta Score: 0  
 o target score: **0.916667**



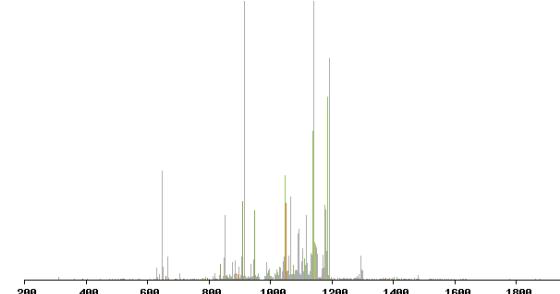
# Supplemental Figure 8

A

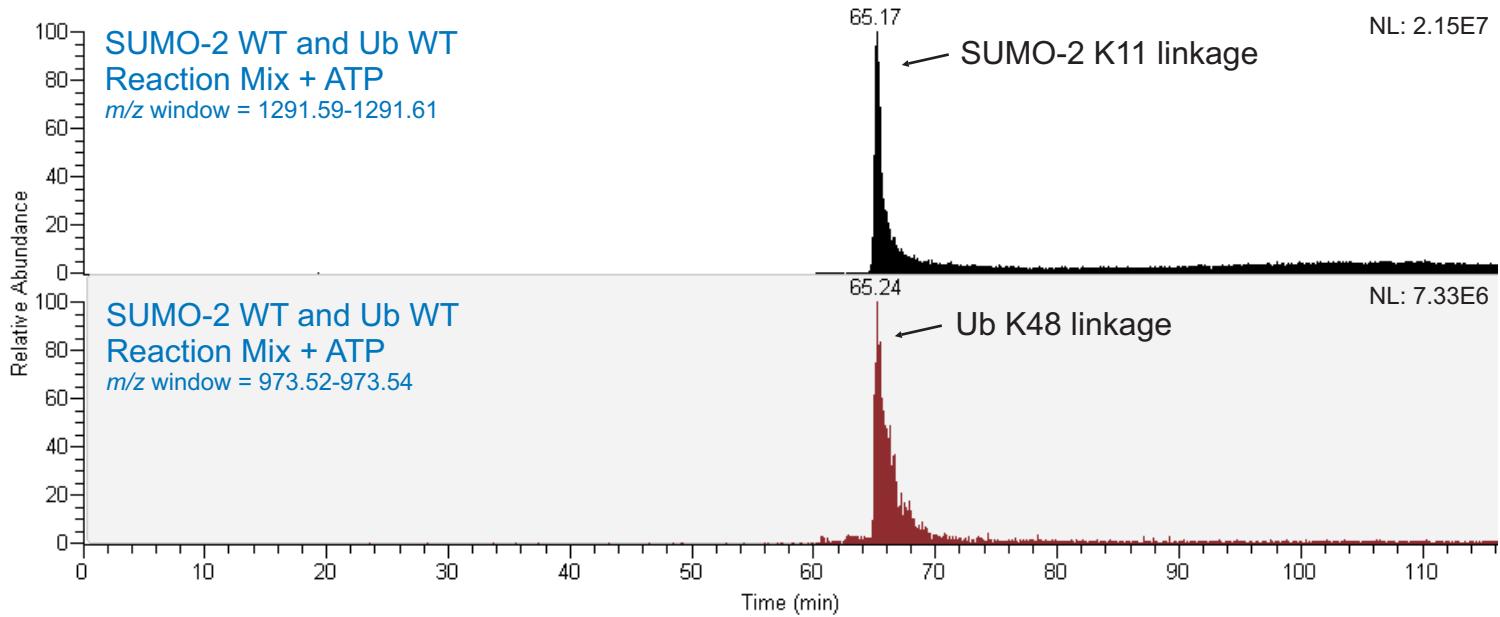
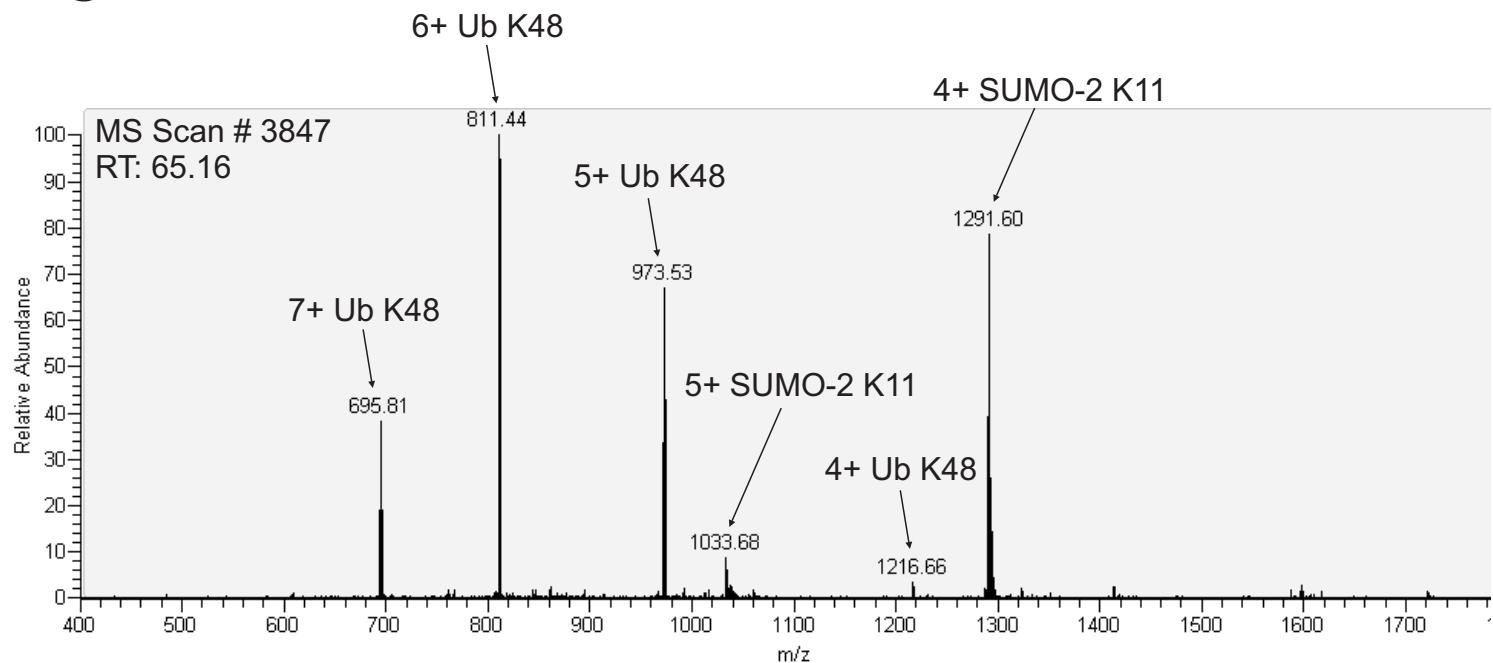
| Scan | Charge | relPos | modScore | precursor mass | matched ions | TIC    | %TIC     | BPI     |
|------|--------|--------|----------|----------------|--------------|--------|----------|---------|
| 6451 | 5      | 1      | 0.914141 | 3428.76        | 40           | 943305 | 0.193573 | 70218.2 |



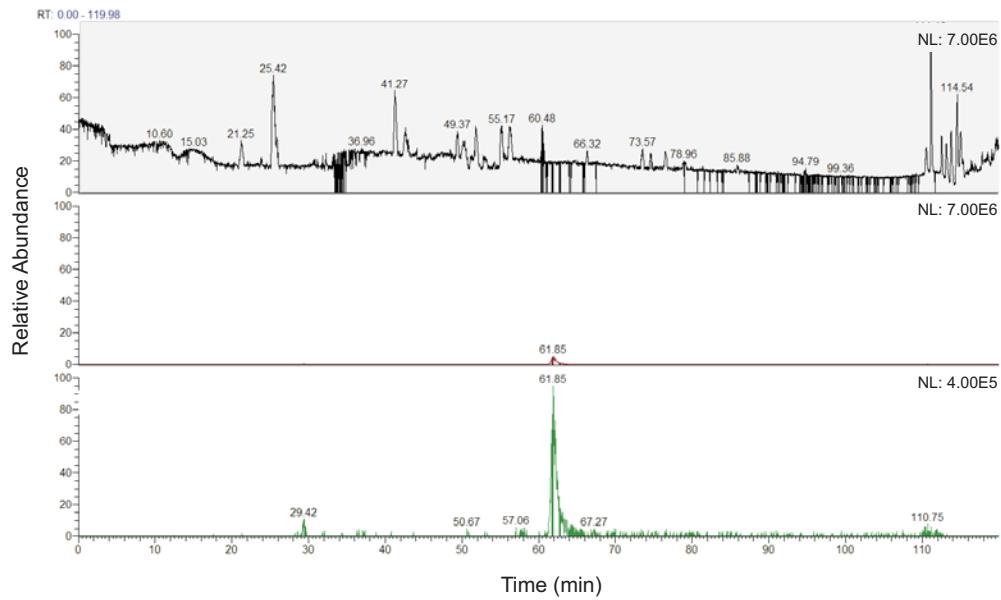
- >P62988|HUMAN Ubiquitin
- EGIPPDQQRIFAGK(0)QLEDGRTLSNDNIQK
- 3428.77
- matched ions: 100
- modScore: 0.508359
- Cons: 0.1
- Delta Score: 0
- target score: **0.967742**



|                     | 1       | 2       | 3       | 4       | 5       | 6       | 7       | 8       | 9       | 10      | 11      | 12      | 13      | 14      | 15      | 16      | 17      | 18      | 19      | 20      | 21      | 22      | 23      | 24      | 25      | 26      | 27      | 28      | 29      | 30      |
|---------------------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|
| E                   | G       | I       | P       | P       | D       | Q       | Q       | R       | L       | I       | F       | A       | G       | K(0)    | Q       | L       | E       | D       | G       | R       | T       | L       | S       | D       | Y       | N       | I       | Q       | K       |         |
| 1432.672060_b1+-0.0 | 130.123 | 187.175 | 300.335 | 397.451 | 494.568 | 609.657 | 737.787 | 865.918 | 1022.11 | 1135.27 | 1248.42 | 1395.6  | 1466.68 | 1523.73 | 3084.58 | 3212.71 | 3325.87 | 3454.98 | 3570.07 | 3627.12 | 3783.31 | 3884.42 | 3997.58 | 4084.65 | 4199.74 | 4362.92 | 4477.02 | 4590.18 | 4718.31 |         |
| 1432.672060_b2+-0.0 | 65.5656 | 94.0916 | 150.671 | 199.23  | 247.788 | 305.332 | 369.398 | 433.463 | 511.557 | 568.136 | 624.716 | 699.304 | 733.844 | 762.37  | 1542.79 | 1606.86 | 1663.44 | 1728    | 1785.54 | 1814.07 | 1892.16 | 1942.71 | 1999.29 | 2042.83 | 2100.38 | 2181.96 | 2239.02 | 2295.59 | 2359.66 |         |
| 1432.672060_b3+-0.0 | 0.044   | 0.0464  | 63.0637 | 100.783 | 133.156 | 165.528 | 203.891 | 246.601 | 289.311 | 341.374 | 379.094 | 416.813 | 465.872 | 489.565 | 508.582 | 1028.86 | 1071.57 | 1109.29 | 1152.33 | 1190.7  | 1209.71 | 1261.78 | 1295.48 | 1333.2  | 1362.22 | 1400.59 | 1454.98 | 1493.01 | 1530.73 | 1573.44 |
| 1432.672060_b4+-0.0 | 33.2867 | 47.5497 | 75.8396 | 100.119 | 124.398 | 153.17  | 185.203 | 217.235 | 256.282 | 284.572 | 312.862 | 349.656 | 367.426 | 381.689 | 771.9   | 803.933 | 832.223 | 864.502 | 893.274 | 907.537 | 946.584 | 971.86  | 1000.15 | 1021.92 | 1050.69 | 1091.49 | 1120.01 | 1148.3  | 1180.33 |         |
| 1432.672060_b5+-0.0 | 26.8309 | 38.2413 | 60.8732 | 80.2966 | 99.7199 | 122.738 | 148.364 | 173.99  | 205.237 | 227.859 | 250.491 | 279.926 | 294.142 | 305.553 | 617.722 | 643.348 | 665.98  | 691.803 | 714.821 | 726.231 | 757.469 | 777.66  | 800.321 | 817.737 | 840.755 | 873.39  | 896.211 | 918.843 | 944.469 |         |
| 1432.672060_y1+-0.0 | 4735.39 | 4678.33 | 4565.18 | 4468.06 | 4370.94 | 4255.85 | 4127.72 | 3999.59 | 3843.4  | 3730.25 | 3617.09 | 3469.91 | 3398.83 | 3341.78 | 1780.93 | 1652.8  | 1539.64 | 1410.53 | 1295.44 | 1238.39 | 1082.2  | 981.094 | 867.934 | 780.856 | 665.767 | 502.591 | 388.487 | 275.328 | 147.197 |         |
| 1432.672060_y2+-0.0 | 2368.2  | 2339.67 | 2283.09 | 2234.53 | 2185.97 | 2128.43 | 2064.37 | 2000.3  | 1922.21 | 1865.63 | 1809.05 | 1735.46 | 1699.92 | 1671.39 | 890.97  | 826.905 | 770.325 | 705.767 | 648.223 | 619.697 | 541.603 | 491.051 | 434.471 | 390.932 | 333.388 | 251.8   | 194.748 | 138.168 | 74.1026 |         |
| 1432.672060_y3+-0.0 | 1579.13 | 1560.12 | 1522.4  | 1490.02 | 1457.65 | 1419.29 | 1376.58 | 1333.87 | 1281.81 | 1244.09 | 1206.37 | 1157.31 | 1133.62 | 1114.6  | 594.316 | 551.606 | 513.886 | 470.847 | 432.485 | 413.467 | 361.405 | 327.703 | 289.983 | 260.957 | 222.594 | 168.202 | 130.168 | 92.4479 | 49.7376 |         |
| 1432.672060_y4+-0.0 | 1184.6  | 1170.34 | 1142.05 | 1117.77 | 1093.49 | 1064.72 | 1032.69 | 1000.65 | 961.607 | 933.317 | 905.027 | 868.233 | 850.463 | 836.2   | 445.989 | 413.956 | 385.666 | 353.388 | 324.615 | 310.352 | 271.306 | 246.029 | 217.739 | 195.97  | 167.198 | 126.404 | 97.8777 | 69.5879 | 37.5552 |         |
| 1432.672060_y5+-0.0 | 947.884 | 936.473 | 913.841 | 894.418 | 874.995 | 851.977 | 826.351 | 800.725 | 769.487 | 746.855 | 724.223 | 694.788 | 680.572 | 669.162 | 356.993 | 331.367 | 308.735 | 282.912 | 259.894 | 248.483 | 217.246 | 197.025 | 174.393 | 156.977 | 133.96  | 101.325 | 78.5038 | 55.8719 | 30.475  |         |
|                     | 30      | 29      | 28      | 27      | 26      | 25      | 24      | 23      | 22      | 21      | 20      | 19      | 18      | 17      | 16      | 15      | 14      | 13      | 12      | 11      | 10      | 9       | 8       | 7       | 6       | 5       | 4       | 3       | 2       | 1       |

**B****C**

# Supplemental Figure 9



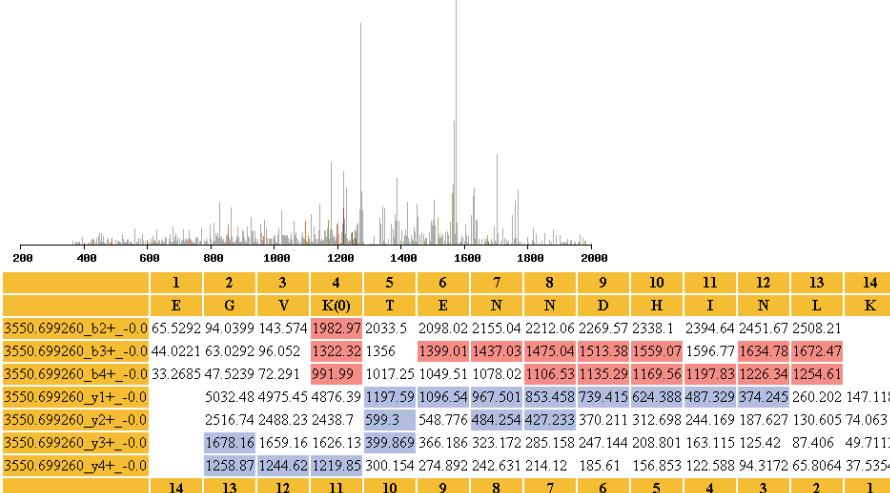
# Supplemental Figure 10

| Scan | Charge | relPos | modScore | precursor mass | matched ions | TIC     | %TIC     | BPI     |
|------|--------|--------|----------|----------------|--------------|---------|----------|---------|
| 2354 | 4      | 1      | 0.882353 | 1609.79        | 132          | 48020.8 | 0.379819 | 2348.23 |



|        | 1       | 2       | 3       | 4       | 5       | 6       | 7       | 8       | 9       | 10      | 11      | 12      | 13      | 14      | 15      | 16      | 17      | 18      | 19      | 20      | 21      | 22      | 23      | 24      | 25      | 26      | 27      | 28      | 29      | 30      | 31      | 32      | 33 |
|--------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|----|
| dep 1+ | 1609.64 | 1666.69 | 1723.74 | 1824.85 | 1952.98 | 2081.11 | 2209.24 | 2356.42 | 2455.55 | 2570.64 | 2683.8  | 2784.9  | 2899.99 | 3029.11 | 3144.2  | 3273.31 | 3404.5  | 3533.62 | 3646.78 | 3774.91 | 3845.99 | 3943.11 | 4044.21 | 4159.3  | 4260.4  | 4389.52 | 4503.62 | 4616.78 | 4713.9  | 4842.03 | 4899.08 | 5014.17 |    |
| dep 2+ | 805.324 | 833.85  | 862.376 | 912.929 | 976.994 | 1041.06 | 1105.12 | 1178.71 | 1228.28 | 1285.82 | 1342.4  | 1392.96 | 1450.5  | 1515.06 | 1572.6  | 1637.16 | 1702.76 | 1767.31 | 1823.89 | 1887.96 | 1923.5  | 1972.06 | 2022.61 | 2080.15 | 2130.71 | 2195.26 | 2252.32 | 2308.9  | 2357.45 | 2421.52 | 2450.05 | 2507.59 |    |
| dep 3+ | 537.219 | 556.236 | 575.253 | 608.955 | 651.665 | 694.375 | 737.086 | 786.145 | 819.188 | 857.552 | 895.271 | 928.973 | 967.336 | 1010.31 | 1048.74 | 1091.78 | 1135.51 | 1178.55 | 1216.27 | 1258.98 | 1282.67 | 1315.04 | 1348.74 | 1387.11 | 1420.81 | 1463.85 | 1501.88 | 1539.6  | 1571.97 | 1614.68 | 1633.7  | 1672.06 |    |
| dep 4+ | 403.166 | 417.429 | 431.692 | 456.968 | 489.001 | 521.034 | 553.066 | 589.86  | 614.644 | 643.416 | 671.706 | 696.982 | 725.734 | 758.033 | 786.805 | 819.084 | 851.882 | 884.161 | 912.451 | 944.483 | 962.253 | 986.532 | 1011.81 | 1040.58 | 1065.86 | 1098.14 | 1126.66 | 1154.95 | 1179.23 | 1211.26 | 1225.53 | 1254.3  |    |
| ind 1+ | 3551.71 | 3494.66 | 3437.6  | 3336.5  | 3208.37 | 3080.24 | 2952.93 | 2705.8  | 2590.71 | 2477.55 | 2376.44 | 2261.36 | 2132.24 | 2017.15 | 1988.04 | 1756.81 | 1627.73 | 1514.57 | 1386.44 | 1315.36 | 1218.21 | 1117.14 | 1002.05 | 900.943 | 771.828 | 657.724 | 544.564 | 447.448 | 319.317 | 262.265 | 147.177 |         |    |
| ind 2+ | 1776.36 | 1747.83 | 1719.31 | 1668.75 | 1604.69 | 1540.62 | 1476.56 | 1402.97 | 1353.4  | 1295.86 | 1239.28 | 1188.73 | 1131.18 | 1066.62 | 1009.08 | 944.522 | 878.925 | 814.368 | 757.788 | 693.723 | 658.183 | 609.625 | 559.072 | 501.528 | 450.976 | 386.418 | 329.366 | 272.786 | 224.228 | 160.162 | 131.637 | 74.0922 |    |
| ind 3+ | 1184.57 | 1165.56 | 1146.54 | 1112.84 | 1070.13 | 1027.42 | 984.707 | 935.648 | 902.604 | 864.241 | 826.522 | 792.82  | 754.457 | 711.418 | 673.056 | 630.017 | 586.286 | 543.248 | 505.528 | 462.818 | 439.125 | 406.753 | 373.051 | 334.688 | 300.986 | 257.948 | 219.913 | 182.193 | 149.821 | 107.111 | 88.0936 | 49.7308 |    |
| ind 4+ | 888.683 | 874.42  | 860.157 | 834.88  | 802.848 | 770.815 | 738.782 | 701.988 | 677.205 | 648.433 | 620.143 | 594.867 | 566.095 | 533.816 | 505.044 | 472.765 | 439.967 | 407.688 | 379.398 | 347.365 | 329.596 | 305.316 | 280.04  | 251.268 | 225.992 | 193.713 | 165.187 | 136.897 | 112.618 | 80.5851 | 66.3222 | 37.55   |    |
|        | 33      | 32      | 31      | 30      | 29      | 28      | 27      | 26      | 25      | 24      | 23      | 22      | 21      | 20      | 19      | 18      | 17      | 16      | 15      | 14      | 13      | 12      | 11      | 10      | 9       | 8       | 7       | 6       | 5       | 4       | 3       | 2       | 1  |

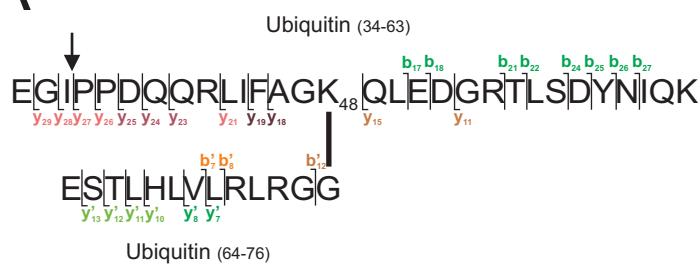
- o >SUMO2WT
- o EGVK(D)TENNNDHINLK
- o 1609.81
- o matched ions: 30
- o %TIC: 0.05396999
- o Cons: 0.142857
- o Delta Score: 0
- o target score: 0.966667



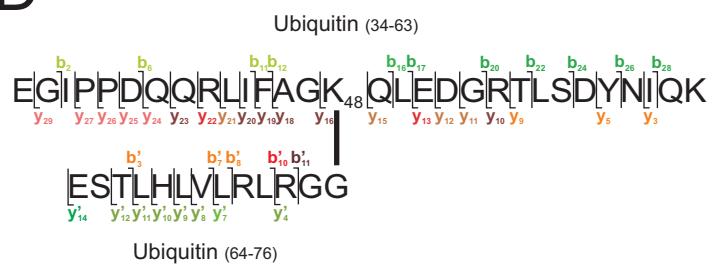
|                      | 1       | 2       | 3       | 4       | 5       | 6       | 7       | 8       | 9       | 10      | 11      | 12      | 13      | 14 |
|----------------------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|----|
| E                    | G       | V       | K(0)    | T       | E       | N       | N       | D       | H       | I       | N       | L       | K       |    |
| 3550.699260_b2+_-0.0 | 65.5292 | 94.0399 | 143.574 | 1982.97 | 2033.5  | 2098.02 | 2155.04 | 2212.06 | 2269.57 | 2338.1  | 2394.64 | 2451.67 | 2508.21 |    |
| 3550.699260_b3+_-0.0 | 44.0221 | 63.0292 | 96.052  | 1322.32 | 1356    | 1399.01 | 1437.03 | 1475.04 | 1513.38 | 1559.07 | 1596.77 | 1634.78 | 1672.47 |    |
| 3550.699260_b4+_-0.0 | 33.2685 | 47.5239 | 72.291  | 991.99  | 1017.25 | 1049.51 | 1078.02 | 1106.52 | 1135.29 | 1169.56 | 1197.83 | 1226.34 | 1254.61 |    |
| 3550.699260_y1+_-0.0 | 503.48  | 4975.45 | 4876.39 | 1197.59 | 1096.54 | 967.501 | 853.458 | 739.415 | 624.388 | 487.329 | 374.245 | 260.202 | 147.118 |    |
| 3550.699260_y2+_-0.0 | 2516.74 | 2488.23 | 2438.7  | 599.3   | 548.776 | 484.254 | 427.233 | 370.211 | 312.698 | 244.169 | 187.627 | 130.605 | 74.063  |    |
| 3550.699260_y3+_-0.0 | 1678.16 | 1659.16 | 1626.13 | 399.869 | 366.186 | 323.172 | 285.158 | 247.144 | 208.801 | 163.115 | 125.42  | 87.406  | 49.7113 |    |
| 3550.699260_y4+_-0.0 | 1258.87 | 1244.62 | 1219.85 | 300.154 | 274.892 | 242.631 | 214.12  | 185.61  | 156.853 | 122.588 | 94.3172 | 65.8064 | 37.5354 |    |
|                      | 14      | 13      | 12      | 11      | 10      | 9       | 8       | 7       | 6       | 5       | 4       | 3       | 2       | 1  |

# Supplemental Figure 11

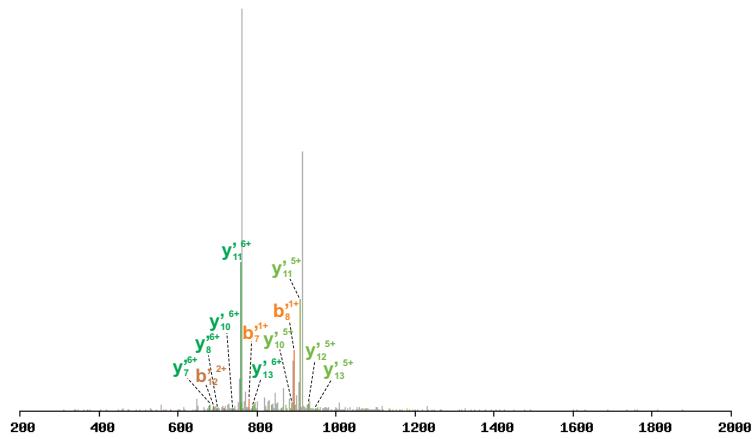
**A**



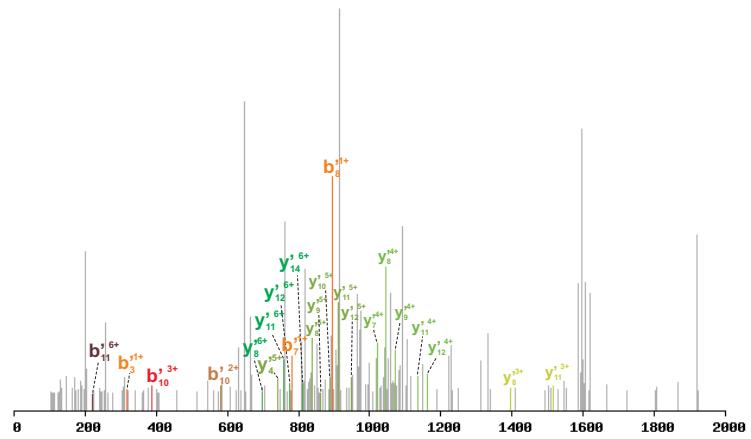
**D**



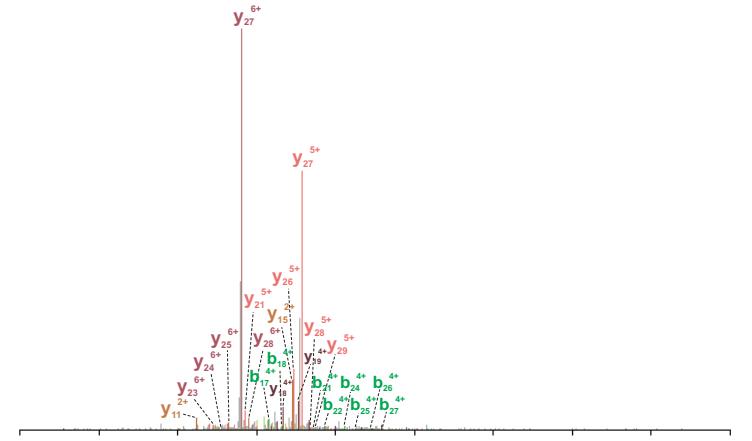
**B** Modification CID



**E** Modification HCD



**C** Target CID



**F** Target HCD

