SUPPLEMENTARY INFORMATION

Structural Motifs for CTD Kinase Specificity on RNA Polymerase II during Eukaryotic Transcription

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Supplementary Figure 1: Sequence logos of the conservation of Yeast and Human CTD. (LogOdds Logo NCBI: Yu, Y.-K.; Capra, J. A.; Stojmirović, A.; Landsman, D.; Altschul, S. F. Log-Odds Sequence Logos. Bioinformatics 2015, 31 (3), 324–331. https://doi.org/10.1093/bioinformatics/btu634.)

			Mass				Mass				Mass
Identified	Theoretical	Observed	Difference	Identified	d Theoretical	Observed [oifference	Identified	Theoretical	Observed I	Differer
lons	Mass	Mass	(ppm)	lons	Mass	Mass	(ppm)	lons	Mass	Mass	(ppm
a12	1361.554	1361.561	5.0	a12+1	1282.596	1282.593	-2.1	a ¹⁵	1628.736	1628.733	-
P.1	1302.517	1302.527	7.6	a°	645.312	645.314	2.1	a1/	1812.821	1812.816	
b ³²	1389.549	1389.554	4.0	a	746.360	746.357	-3.5	a	645.312	645.314	
b ¹⁶	1823.729	1823.725	-2.5	a°	909.423	909.421	-2.0	a ⁸	909.423	909.424	
b ¹⁹	2094.846	2094.856	4.6	b11	1222.551	1222.559	7.1	b11	1222.551	1222.552	
P,	489.222	489.224	2.6	b12	1309.583	1309.586	2.6	b12	1309.583	1309.586	
P ₂	576.254	576.255	1.7	b14	1493.667	1493.666	-1.1	b14	1493.667	1493.667	-
b ²	774.355	774.358	4.0	b ²²	2441.994	2441.999	1.9	b18	1927.848	1927.848	
b ⁸	937.418	937.420	2.2	b ⁴	489.222	489.222	0.1	b ²²	2362.028	2362.026	-
b ⁹	1038.466	1038.469	3.2	b ⁸	937.418	937.420	2.2	b ²³	2449.060	2449.039	-
C14	1590.660	1590.663	2.1	b ⁹	1038.466	1038.467	0.7	b ⁴	489.222	489.223	
C21	2295.957	2295.956	-0.5	C14	1510.694	1510.694	-0.2	be	673.307	673.310	
c'	791.381	791.383	2.3	c ²¹	2295.957	2295.955	-1.1	b ²	774.355	774.357	
c ⁸	954.445	954.439	-5.7	c ⁶	690.333	690.334	0.1	b ⁸	937.418	937.420	
x11+1	1163.522	1163.523	1.0	c'	791.381	791.382	1.6	b ⁹	1038.466	1038.466	
x13+1	1347.607	1347.597	-7.1	x ⁸ +1	892.405	892.402	-3.5	C14	1510.694	1510.695	
x8+1	892.405	892.405	-0.5	V ¹⁰	1129,469	1129,467	-2.4	c6	690,333	690,333	
V10	1049.503	1049.506	3.0	Y11	1216.501	1216.491	-8.7	c?	791.381	791.382	
Y11	1136.535	1136.539	3.4	Y12	1313.554	1313.556	1.3	c ⁸	954 445	954 444	
V12	1233.588	1233.590	2.1	V13	1400.586	1400.588	1.3	x11+1	1243,488	1243,491	
V13	1320.620	1320.622	1.9	v13-1	1399.578	1399.582	2.5	V10	1129,469	1129,469	
V14	1483.683	1483.674	-6.0	V14	1563,649	1563.650	0.1	V11	1216 501	1216.501	
V16	1667.768	1667.769	0.7	V16	1747.734	1747.736	1.2	V12	1313.554	1313.556	
V17	1754,800	1754,804	2.3	V17	1834,766	1834,769	1.7	v12-1	1312 546	1312 547	
V19	2018.851	2018.858	3.3	V ¹⁹	2018.851	2018.856	2.3	V13	1400.586	1400.587	
V20	2119.899	2119,900	0.7	V20	2119,899	2119,900	0.7	¥14	1563 649	1563 652	
v ²³	2481.062	2481.072	3.8	V21	2282.962	2282.968	2.7	V15	1650,681	1650,692	
V24	2568.095	2568.097	1.0	V ²³	2481.062	2481.069	2.5	¥16	1747 734	1747 736	
V26	2793.206	2793.207	0.6	V24	2568.095	2568,100	2.1	V17	1834 766	1834 771	
V ³	330.190	330.190	0.2	y26-1	2792.198	2792.194	-1.5	V19	2018.851	2018.856	
VS	528,291	528,291	1.3	V ³	330,190	330,191	0.9	¥20	2119 899	2119 900	
v5-1	527.283	527.281	-3.0	vs	528.291	528.292	1.8	V21	2282.962	2282.968	
V ⁶	615.323	615.323	0.8	V ⁶	615.323	615.324	1.1	V23	2481.052	2481.061	
V2	778.386	778.392	7.4	V2	778.386	778,391	5.9	v24	2568.095	2568.098	
VS	865 418	865 420	2.5	V ⁹	962 471	962 472	0.7	126-1	2792 198	2792 199	1.1
V ⁹	962,471	962,472	1.3	z ¹³	1384,567	1384.571	2.6	VS -	608 257	608 257	
Z13	1304.601	1304.594	-5.4	Z ⁶	599,304	599,305	1.0	x5-1	607 249	607 249	
74	415 219	415 219	-0.1	-8	849 399	849 400	0.5	y -1	607.249	695 290	
76	599.304	599.303	-1.4		0.0.000	212.100		22	858 352	858 352	
-	222.204	555.505						*	046 205	0/E 204	
								22	1042 427	1042 429	
								-10	1112 /54	1112 457	
								24	1113.451	1113.45/	
								-6	435.186	435.166	
								2	6/9.270	6/9.268	

Supplementary Figure 2: Lists of fragment ions for peptides analyzed in Figure 1B by UVPD-MS for repeats 33,34 and 35. In each case, the 3+ charge state was selected, and UVPD was performed using 2 pulses (1.5 mJ per pulse). The identified site of phosphorylation is shaded in blue in the sequence map above each table.

Mass Identified Theoretical Observed Difference Mass lons Mass (ppm) 0.9 a°+1 685.260 685.261 a⁹+1 1036.403 1036.399 -3.5 a¹⁰ 1132.448 1132.447 -0.9 a¹² 1334.543 1334.544 0.3 a¹³ 1431.596 1431.593 -2.1 a⁸ 948.363 948.358 -4.9 b12 1362.538 1362.541 2.0 b? 813.295 813.295 0.4 b⁸ 976.358 976.359 1.5 **c**¹¹ 1278.517 1278.517 0.2 c² 267.122 267.122 2.1 c7 830.321 830.321 -0.5 **c**⁸ 993.384 993.387 2.7 x13+1 1469.620 1469.618 -1.2 x8+1 920.437 920.435 -1.3 x11 1284.528 1284.531 2.4 x4 471.233 471.233 0.9 x7 818.381 818.380 -1.3 y10 1157.501 1157.504 2.9 1258.548 1258.548 y11 -0.3 y12 1355.601 1355.602 0.9 y¹³ 1442.633 1442.634 0.9 $\gamma^{2} = \gamma^{3} = \gamma^{5} = \gamma^{6} = \gamma^{7} = \gamma^{8} = \gamma^{9} = z^{11}$ 243.158 243.159 1.4 344.206 344.206 0.9 542.306 542.307 0.5 629.338 629.340 1.9 792.402 792.403 1.7 893.449 893.450 1.1 990.502 990.504 1.9 1242.529 1242.530 0.1 z¹³ 1426.614 1426.616 1.1 z4 429.235 429.235 0.8 z 613.320 613.320 1.1 z⁸ 877.431 877.431 0.3

Identified	Theoretical	Observed	Mass
Identified	Mass	Mass	Difference
ions	IVIdSS	IVIdSS	(ppm)
a ¹² +1	1335.551	1335.554	2.0
a ⁶ +1	605.294	605.293	-1.6
a ⁸ +1	869.405	869.403	-1.8
a²	222.100	222.101	1.3
b12	1362.538	1362.539	1.0
b ⁴	448.196	448.196	1.1
P ₈	896.392	896.393	2.0
c ²	267.122	267.122	2.0
C ⁴	465.222	465.223	1.3
C ⁶	649.307	649.306	-2.1
x13+1	1469.620	1469.620	0.1
x6+1	736.292	736.288	-4.9
x7+1	899.355	899.354	-1.7
x ⁸ +1	1000.403	1000.400	-2.9
x ¹⁰	1183.480	1183.482	1.4
x ¹¹	1284.528	1284.531	2.6
y ¹⁰	1157.501	1157.503	2.2
y ¹¹	1258.548	1258.551	2.3
y ¹²	1355.601	1355.603	1.4
y ¹³	1442.633	1442.634	0.9
y ²	243.158	243.159	1.5
Y ²	872.368	872.370	2.0
Y ⁸	973.416	973.418	2.5
Y ²	1070.469	1070.461	-7.5
z ¹¹	1242.529	1242.533	2.5
z ¹³	1426.614	1426.619	3.6
z ⁸	957.397	957.392	-5.2

Supplementary Figure 3: Lists of fragment ions for peptides analyzed in Figure 1B by UVPD-MS for repeats 41 and 42. In each case, the 3+ charge state was selected, and UVPD was performed using 2 pulses (1.5 mJ per pulse). The identified site of phosphorylation is shaded in blue in the sequence map above each table.

Identifie Ions	d Theoretical Mass	Observed I	Mass Difference (ppm)	Identified	d Theoretical Mass	Observed Mass	Mass Difference	Identifie Ions	d Theoretical Mass	Observed Mass	Mass Difference (ppm)
a ¹⁴ +1	1517.657	1517.659	2.0	a13+1	1338.622	1338.621	-0.6	a14+1	1517.657	1517.660	2.
a ⁹ +1	956.437	956.440	3.6	a ⁹ +1	956,437	956,427	-9.6	b11	1261.490	1261.492	1.
a ¹⁵	1679.712	1679.713	0.5	a ¹²	1240.561	1240.553	-6.7	b12	1348.522	1348.524	1.
a ⁷	705.333	705.334	0.5	a ⁸	868.397	868.397	0.7	b14	1544.644	1544.646	1.
a ⁸	868.397	868.393	-3.7	b11	1181.524	1181.522	-1.7	b15	1707.707	1707.709	1.
b11	1181.524	1181.521	-3.0	b12	1268,556	1268.557	0.9	b16	1808.755	1808.757	1.
b14	1544.644	1544.649	3.3	b14	1464.677	1464.681	2.8	b ⁴	448.196	448.196	0.
b15	1707.707	1707.706	-0.3	b15	1627,741	1627.743	1.3	b ²	813.295	813.295	0
b ¹⁶	1808.755	1808.759	2.6	b16	1728.788	1728.790	0.7	b ⁸	976.358	976.359	0
b ²	250.095	250.096	1.2	b ²	250.095	250.096	0.6	b ⁹	1063.390	1063.391	1
b4	448.196	448.196	0.5	b ⁴	448.196	448.196	-0.1	C14	1561.670	1561.672	1
b ⁷	733.328	733.329	1.0	b ⁷	733.328	733.330	2.6	c ²	267.122	267.122	1
b ⁸	896.392	896.393	1.6	b ⁸	896.392	896.393	1.4	c?	830.321	830.322	0
b ⁹	983.424	983.425	1.8	b ⁹	983.424	983.425	1.3	C ⁹	1080.416	1080.415	-0
c14	1561.670	1561.677	4.6	c14	1481.704	1481.706	1.8	x11+1	1203.590	1203.592	2
c ²	267.122	267.122	2.0	C ²	267.122	267.122	1.6	x2+1	270.145	270.145	C
c ²	750.355	750.357	2.8	c7	750.355	750.356	2.3	x7+1	819.389	819.389	C
x11+1	1283.556	1283.555	-0.7	C ⁸	913.418	913,420	2.0	x ⁸	917.449	917.453	3
x4+1	458.225	458.225	0.1	c ⁹	1000.450	1000.450	0.4	Y10	1075.555	1075.557	1
¢.	457.217	457.218	0.8	V10	1155.521	1155.523	1.7	y11	1176.603	1176.606	3
2	818.381	818.383	2.5	Y11	1256.569	1256.570	0.7	y12	1273.655	1273.657	1
8	917.449	917.451	1.3	V12	1353.622	1353.623	0.8	Y13	1360.687	1360.690	2
10	1155.521	1155.524	2.4	V13	1440.654	1440.656	1.8	y14	1523.751	1523.750	-0
V11	1256.569	1256.565	-3.5	V14	1603.717	1603.722	2.8	y16	1721.851	1721.853	1
y12	1353.622	1353.623	1.3	Y16	1801.818	1801.820	1.3	y17	1888.850	1888.843	-3
y ¹³	1440.654	1440.657	2.1	Y19	2086.950	2086.950	0.2	y19	2086.950	2086.952	C
y14	1603.717	1603.722	3.0	V ²	243.158	243.159	0.7	y ²	243.158	243.159	1
y15	1704.765	1704.752	-7.5	y ^s	608.257	608.257	0.2	Y3	330.190	330.191	1
y ¹⁶	1801.818	1801.821	1.9	V ⁶	709.305	709.305	0.6	y ^s	528.291	528.291	1
y ¹⁹	2086.950	2086.954	2.0	Y?	872.368	872.369	0.9	Y°	629.338	629.339	1
y ²	243.158	243.159	1.2	V ⁸	971.437	971.437	0.1	y ²	792.402	792.403	1
V ³	330.190	330.191	1.3	v	1068.489	1068.491	1.4	Y ⁸	891.470	891.473	2
y ^s	528.291	528.291	1.2	Z11	1240.550	1240.553	2.1	Y ²	988.523	988.525	1
Y ⁶	629.338	629.339	1.4	z ⁶	693.286	693.286	-0.5	Z11	1160.584	1160.583	-0
y?	792.402	792.403	1.3					z4	415.219	415.219	-0
y ⁸	891.470	891.476	6.4					z ⁶	613.320	613.320	0
V ²	988.523	988.524	1.5					z*	875.451	875.453	2
z ¹¹	1240.550	1240.549	-0.8								
z4	415.219	415.219	-0.5								
z ⁶	613.320	613.320	0.6								
8	875.451	875,453	2.1								

Supplementary Figure 4: Lists of fragment ions for peptides analyzed in Figure 1B by UVPD-MS for repeats 43,44 and 45. In each case, the 3+ charge state was selected, and UVPD was performed using 2 pulses (1.5 mJ per pulse). The identified site of phosphorylation is shaded in blue in the sequence map above each table.

JK

Identified	Theoretical	Observed	Mass
long	Mass	Marr	Difference
IONS	IVIdSS	IVIdSS	(ppm)
a°+1	685.260	685.260	-0.2
a ⁹ +1	1036.403	1036.401	-1.5
a ¹⁰	1132.448	1132.450	1.5
a ¹²	1320.528	1320.526	-0.9
a ¹³	1417.580	1417.577	-2.3
a ⁸	948.363	948.361	-2.5
b12	1348.522	1348.523	0.4
b ²	813.295	813.295	0.1
b ⁸	976.358	976.358	0.5
c11	1278.517	1278.517	-0.1
C ⁶	729.273	729.274	1.3
c ⁸	993.384	993.386	1.5
x13+1	1455.604	1455.604	-0.7
x*+1	458.225	458.225	0.4
x ⁶ +1	642.310	642.309	-1.8
x ¹⁰	1169.464	1169.463	-1.3
x11	1270.512	1270.514	1.9
x4	457.217	457.217	-0.1
x'	804.365	804.366	0.8
x ⁸	905.413	905.414	1.1
y ¹⁰	1143.485	1143.489	3.4
y ¹¹	1244.533	1244.532	-0.9
y ¹²	1341.585	1341.586	0.8
y ¹³	1428.617	1428.619	1.4
γ ²	243.158	243.159	1.5
Y ³	330.190	330.191	1.1
y ^s	528.291	528.291	1.1
y ⁶	615.323	615.324	1.5
y?	778.386	778.386	0.3
y ^s	879.434	879.436	2.7
Y ⁹	976.487	976.489	2.2
z ¹¹	1228.514	1228.516	1.7
z ¹³	1412.599	1412.600	1.3
Z ⁴	415.219	415.219	0.1
z ⁶	599.304	599.304	0.5
z ⁸	863.415	863.416	1.3

46/48 Nylslpltislpltslpltslplt Sp к 47/49

Identified	Theoretical	Observed	Mass
lons	Mass	Mass	Difference
	111035	111255	(ppm)
a12+1	1321.535	1321.536	0.8
a ⁹ +1	956.437	956.437	-0.1
a ⁸	868.397	868.399	2.5
b ⁴	448.196	448.196	1.2
b ^s	535.228	535.228	1.1
P8	896.392	896.390	-1.4
C ¹	180.090	180.090	3.0
C ²	267.122	267.122	2.3
C ²	1000.450	1000.453	3.0
x12+1	1368.572	1368.583	7.9
x13+1	1455.604	1455.608	2.6
x ⁵ +1	635.244	635.245	0.7
x7+1	885.340	885.340	0.8
x ⁸ +1	986.387	986.384	-3.2
x ¹¹	1270.512	1270.515	2.7
x ⁶	721.268	721.269	1.2
y ¹⁰	1143.485	1143.488	2.5
y ¹¹	1244.533	1244.535	1.5
y12	1341.585	1341.587	1.4
y ²	243.158	243.159	1.6
Y ⁴	511.204	511.206	2.6
YS	608.257	608.258	0.8
y?	858.352	858.354	1.2
y ⁸	959.400	959.403	3.4
y ⁹	1056.453	1056.454	1.4
z ¹¹	1228.514	1228.516	1.9
z ¹³	1412.599	1412.601	1.5
z4	495.186	495.186	0.4
z ⁶	679.270	679.270	0.1
z ⁸	943.381	943.383	2.0

Supplementary Figure 5: Lists of fragment ions for peptides analyzed in Figure 1B by UVPD-MS for repeats 46/48 to 47/49. In each case, the 3+ charge state was selected, and UVPD was performed using 2 pulses (1.5 mJ per pulse). The identified site of phosphorylation is shaded in blue in the sequence map above each table.



Supplementary Figure 6: LC-MS/MS trace obtained for doubly phosphorylated peptides of T4Q variation in human distal CTD. The peak numbers correspond to the indicated sites (blue highlights) of phosphorylation identified by MS/MS.



Supplementary Figure 7: The modeling of CTD with Ser2 in the active site places Pro6 in a hydrophilic environment.

2	GIPLG S G MISIPIKIYISIPITISIPIKIYISIP TISIPIKIYIS	
Z	24 PT C	

			Mass
Identified	Theoretical	Observed	Difference
lons	mass	mass	(ppm)
a*12	1118.530	1118.531	0.8
a ⁺¹⁵	1403.663	1403.670	5.2
a ⁺¹⁸	1781.853	1781.851	-1.4
a ⁺²⁶	2719.248	2719.249	0.6
a**	643.287	643.288	0.7
a ¹³	1218.570	1218.576	4.6
a ¹⁷	1693.813	1693.816	1.8
a ²⁰	1978.946	1978.954	4.1
a ²²	2163.031	2163.028	-1.0
a ²⁴	2454.189	2454.190	0.6
a ⁹	770.375	770.377	3.2
b ¹³	1246.565	1246.568	2.3
b14	1333.597	1333.596	-0.9
b ¹⁶	1558.745	1558.757	8.0
b ¹⁸	1808.840	1808.825	-8.6
b ²⁰	2006.941	2006.939	-0.7
b ²¹	2093.973	2093.972	-0.6
b ²³	2319.121	2319.118	-1.1
b ⁹	798.369	798.365	-6.0
C ¹⁰	978.459	978.459	-0.1
C12	1162.544	1162.545	1.3
C ¹³	1263.592	1263.591	-0.5
C ¹⁵	1447.676	1447.677	0.6
c ¹⁶	1575.771	1575.772	0.7
c ¹⁷	1738.835	1738.834	-0.3
c ²⁶	2763.261	2763.257	-1.7
C ₆	503.216	503.217	1.8
C ⁹	815.396	815.392	-4.4
x ¹⁷	1929.840	1929.839	-0.4
x ¹⁹	2220.998	2221.007	4.2
Y ¹⁰	1143.485	1143.480	-4.1
Y ¹¹	1306.548	1306.561	9.5
y ¹³ -1	1530.688	1530.682	-4.1
Y ¹⁴	1618.728	1618.722	-3.8
y ¹⁶ -1	1815.821	1815.820	-0.4
¥ ²⁰	2292.072	2292.072	0.2
¥ ²¹	2379.104	2379.108	1.9
٧ ²⁵	2711.219	2711.232	4.8
y ⁹ -1	1055.445	1055.444	-0.6
Z ²⁵	2695.200	2695.223	8.5
z ²⁶	2792.253	2792.274	7.6

3	NGPGSGM ≈)P)TC	A S PLKIYLSLPITI	SÌP K Y <mark>S</mark> PÌT	[S] P]K]Y S ≥5
	Identified	Theoretical	Observed	Mass Difference
	lons	mass	mass	(ppm)
	a	770.375	770.370	-5.3
	a ²⁶	2718.240	2718.253	4.8
	a ¹⁷	1693.813	1693.815	1.2
	a ⁸ +1	643.287	643.286	-1.9
	a ²⁵ +1	2622.195	2622.199	1.3
	a ²² +1	2244.005	2244.004	-0.2
	a ¹⁹ +1	1958.872	1958.874	0.8
	a ¹⁸ +1	1861.820	1861.818	-0.9
	a ²¹ +1	2146.952	2146.963	5.3
	a11+1	1021.478	1021.468	-9.9
	a ¹² +1	1118.530	1118.531	0.1
	b ¹⁴	1333.597	1333.595	-1.9
	b ¹¹	1048.465	1048.464	-0.7
	b ²¹	2173.939	2173.934	-2.3
	b ²³	2399.087	2399.076	-4.4
	c ¹³	1263.592	1263.585	-4.9
	c ¹²	1162.544	1162.544	0.4
	c ¹⁹	2002.886	2002.879	-3.2
	x ¹⁷	1929.840	1929.839	-0.4
	x ¹⁹	2220.998	2221.005	3.0
	y ⁶	691.354	691.354	0.3
	y ⁷	778.386	778.388	2.1
	y ¹⁰	1143.485	1143.482	-2.8
	y ¹³	1531.696	1531.693	-1.9
	y ¹⁶ -1	1815.821	1815.819	-0.8
	z ⁸	863.415	863.412	-4.0
	z ¹⁷	1887.842	1887.831	-5.9

4	NG <mark>(</mark> PG(SGI ∞]P]TC	MLS PLKIY <mark>s</mark> pit	slp κ γ]s]p]	τ ις] ρ]κ Υ]ς 25
	Identified Ions	Theoretical mass	Observed mass	Mass Difference (ppm)
	a ²⁶	2718.240	2718.252	4.2
	a ¹⁷	1773.780	1773.781	0.9
	a ²⁴	2534.155	2534.160	2.0
	a ²²	2242.997	2243.000	1.5
	a ¹² +1	1198.497	1198.497	-0.1
	a ¹⁸ +1	1861.820	1861.824	2.6
	a ¹⁹ +1	1958.872	1958.878	2.9
	a ²¹ +1	2146.952	2146.954	1.0
	b ²⁵	2649.182	2649.191	3.4
	b9	798.369	798.364	-6.7
	b ²¹	2173.939	2173.946	3.4
	C19	2002.886	2002.875	-5.6
	C ¹⁰	978.459	978.460	1.1
	X ¹⁹	2220.998	2221.012	6.4
	x ⁸ +1	906.421	906.415	-6.8
	y ⁶	691.354	691.354	-0.8
	y ⁷	778.386	778.388	2.6
	y ⁹	976.487	976.486	-1.1
	y ¹³	1451.730	1451.727	-1.5
	y ²¹	2379.104	2379.105	0.7
	y ²⁴ -1	2653.190	2653.182	-2.9
	y ⁶ -1	690.346	690.342	-6.7
	y ¹⁶ -1	1735.854	1735.852	-1.2
	z ¹⁷	1887.842	1887.843	0.5
	z ²⁶	2792.253	2792.267	5.1

Supplementary Figure 8: Lists of fragment ions for mono-phosphorylated peptides (m/z of 956.11) analyzed in Figure 6A by UVPD-MS. In each case, the 3+ charge state was selected, and UVPD was performed using 2 pulses (1.5 mJ per pulse). The identified site of phosphorylation is shaded in blue in the sequence map above each table.

	MLSLPIELYLSLPIT	lslp]elv <mark>s</mark> p	TLSLPLELYLS ??		MISLP ELYISLP	τζεζριεζγιεζρι	ד אין צער אין אין אין די	4	© G P G S[G 20]P]T ⊂	M <mark>ls[p Elvls]p</mark>]1	r <mark>s</mark> p elvisipi	TĮSĮP]EĮYĮS ≈	5 🖓	iPGS[GN]⊺	Se civisieit	lsletelvlslet	,T lsl P El¥ls ∷
Identified Ions	Theoretical mass	Observed mass	Mass Difference (ppm)	ldentified lons	I Theoretical mass	Observed mass	Mass Difference (ppm)		Identified Ions	Theoretical mass	Observed mass	Mass Difference (ppm)	id	lentified Ions	Theoretical mass	Observed mass	Mass Difference (ppm)
a11+1	1022.425	1022.416	-9.5	a11+1	1022.425	1022.42	·5.2		a10+1	935.393	935.384	-9.6		a12+1	1199.444	1199.452	5.9
a12+1	1119.478	1119.486	6.8	a12+1	1119.478	1119.482	3.5		a12+1	1119.478	1119.483	4.6		a14+1	1387.524	1387.518	-4.7
a15+1	1404.611	1404.615	2.8	a15+1	1404.611	1404.615	3.2		a16+1	1613.619	1613.628	5.5		a ¹⁸ +1	1863.715	1863.715	0.2
a23+1	2374.943	2374.953	4.5	a ²³ +1	2374.943	2374.953	4.4		a19+1	1960.768	1960.780	6.3		a ²¹ +1	2148.847	2148.859	5.5
a8+1	643.287	643.286	-1.6	a ²⁶ +1	2722.091	2722.11	7.1		a22+1	2245.900	2245.908	3.8		a ²⁶ +1	2722.091	2722,112	7.7
a ¹⁰	934.385	934.383	-3.1	a ¹⁰	934.385	934.383	-3		a23+1	2374.943	2374.950	3.2		a6+1	459.203	459.205	4.5
a13	1219.518	1219.509	-7.4	a17	1695.709	1695.717	5.1		a ¹³	1219.518	1219.507	-8.6		a ¹⁰	1014.352	1014.356	4.4
a15	1403.603	1403.61	5.2	a ¹⁸	1782.741	1782.746	3		a ²¹	2147.839	2147.840	0.4		a11	1101.384	1101.385	0.7
a17	1695.709	1695.72	6.5	a ²¹	2147.839	2147.837	-1.2		a ²⁵	2624.030	2624.044	5.1		a17	1775.675	1775.681	3.4
a ²¹	2147.839	2147.839	-0.2	a ⁹	771.322	771.318	-5.9		a ²⁶	2721.083	2721.095	4.6		b10	1042.347	1042.350	3.5
a ²⁵	2624.03	2624.057	10	b ¹⁰	962.38	962.383	2.5		a ⁹	771.322	771.321	-1.4		b11	1129.379	1129.382	2.5
a ²⁶	2721.083	2721.107	8.8	b11	1049.412	1049.416	3.8		b10	962.380	962.383	2.3		b13	1327.479	1327.480	0.9
b10	962 38	962 383	3	b13	1247.513	1247.517	3.4		b11	1049.412	1049.416	3.8		b14	1414,511	1414.517	3.8
b ¹¹	1049.412	1049.417	4.5	b14	1334.545	1334.55	3.6		b13	1247.513	1247.515	1.9		b16	1640.607	1640.612	3.4
b ¹³	1247.513	1247.517	3.6	b ¹⁶	1560.64	1560.648	5.1		b16	1640.607	1640.612	3.2		b17	1803.670	1803.678	4.6
b14	1334,545	1334.551	4.3	b17	1723.704	1723.708	2.5		b17	1803.670	1803.678	4.3		b18	1890.702	1890.712	5.4
h ¹⁶	1560.64	1560 645	2.9	b18	1810.736	1810.743	4.1		b18	1890.702	1890.710	4.4		b ²⁰	2088.802	2088.819	8
h17	1723 704	1723 708	23	b ²⁰	2008.836	2008.851	7.4		b ²⁰	2088.802	2088.805	1.2		b ²¹	2175.834	2175.841	3.2
- h ²⁰	2088 802	2088 804	0.8	b ²¹	2175.834	2175.842	3.7		b21	2175.834	2175.841	3.1		b ²³	2401.930	2401.947	7.2
	2175 834	2175 843	4.1	b24	2564,993	2565.003	4		b23	2401.930	2401.948	7.6		b24	2564.993	2565.003	3.9
	2564 993	2565.005	4.5	h ²⁵	2652 025	2652 035	3.8		b ²⁴	2564.993	2565.003	4		b ²⁵	2652.025	2652.036	4.1
h	486.19	486 191	23	b ⁶	486.19	486 191	2.1		b€	486.190	486.191	2		b6	486.190	486.191	1.9
b7	573 222	573 223	2.5	C ¹⁶	1577 667	1577 673	43		b9	799.317	799.320	3.8		b9	879.283	879.284	1.1
ь»	799 317	799 322	5.8	C ²³	2418 956	2418 965	3.9		C16	1657.633	1657.637	2.5		C10	1059.373	1059.367	-6
r13	1264 539	1264 543	2.9	c9	816.343	816.346	2.8		C ⁹	816.343	816.346	3		c13	1344,505	1344,515	7.2
C ¹⁶	1577 667	1577 673	4.7	x4+1	493.193	493.193	0.1		x ¹¹ +1	1254.517	1254.524	5.9		c ¹⁶	1657.633	1657.637	2.5
c23	7/18 956	2/18 962	73	v ¹¹	1307 496	1307 499	2.4		x9+1	1004 421	1004 429	73		C9	896 310	896 310	0.7
-9	816 343	816 346	3.4	v13	1533 591	1533 597	4		¥10	1064 466	1064 470	3.8		x ¹¹ +1	1254 517	1254 522	4.1
v4+1	493 193	493 197	-2	v14	1620 623	1620.633	6.2		v13	1453 625	1453 634	6		x ⁸	906 361	906 359	-1.4
x9+1	1004 421	1004 427	5.6	v ¹⁶ -1	1817 716	1817 719	1.6		v ¹³ -1	1452.617	1452.609	-5.4		v ¹⁰	1064.466	1064.470	4
v11	1307 496	1307 501	4	v17	1905 756	1905 764	4.2		v ¹³ -1	1452.617	1452.621	2.6		v ¹¹	1227.530	1227.536	5.6
v13	1533 591	1533 598	4.5	vis	2068 819	2068 821	1		v14	1620 623	1620 636	7.6		y12	1356 572	1356 562	-73
y 13_1	1532 583	1532 585	0.8	v ²⁰ -1	2293 907	2293 917	4.4		v17	1905 756	1905 764	4.2		v13	1453 625	1453 631	4.4
y 14	1620 623	1670.63	4.2	v ²¹	2381 946	2381.95	13		v18	2068.819	2068.828	4.3		v14	1540.657	1540.663	3.7
1 v16	1919 724	1919 73	37	v23_1	2569.001	2569.004	1 2		v ² 0	2294 914	2294 924	43		v ¹⁶	1738 757	1738 763	3
y v16_1	1817 716	1817 77	2.5	, - u ³	303 143	303 144	23		v ²¹	2381 946	2381 951	1.8		v ¹⁶ -1	1737 750	1737 752	1.2
y .1	1905 756	1905 765	4.7	v ⁶	692 302	692 304	2.9		v ²³ -1	2569.001	2568 998	-1.1		y17	1825 789	1825 795	2.8
7	2069.010	2000 010	4.7	1	859.3	859 301	0.6		, - v ³	303 143	303 144	7.5		v20_1	2213 940	2213 935	-7.7
¥	2008.819	2008.818	-0.6	7 10	960 348	960 345	-7.4		y 104	466 206	466 207	13		y 1 y21	2381 946	2215.555	0.2
y1	2293.907	2255.510	1.5	10	1057 401	1057 404	2.4		y v6	692 302	692 304	3.2		v23_1	2569.001	2569.007	2.4
y	2301.240	2301.33	1.0	y.	1057.401	1007.404	2.2		y7	779 334	779 337	4		, . v ³	303 143	303 144	27
y1	2021/2	202 144	1.0						, v9	977 434	977 437	27		7 V ⁶	692 302	692 304	3.1
γ. 14	466 206	466 207	2.0						,	554	5	2.7		, v?	779 334	779 337	4.5
y.	400.200	400.207	1.2											7	977 434	977 437	29
¥- 112	770 224	770 220	5.2											,	211.424	577.757	2.0
¥.	779.334	779.338	5.1														
Ζ'	5/9.23	3/9.233	4./														

Supplementary Figure 9: Lists of fragment ions for mono-phosphorylated peptides (m/z of 957.05) analyzed in Figure 6B by UVPD-MS. In each case, the 3+ charge state was selected, and UVPD was performed using 2 pulses (1.5 mJ per pulse). The identified site of phosphorylation is shaded in blue in the sequence map above each table.

dentified	Theoretical	Observed	Mass Difference
01111	1022.425	1022 426	
d***1	1022.425	1022.420	0.0
-19+1	1404.011	1010 707	2.0
a***1 a21+1	1918.757	1918.762	2.0
+1	2100.857	2100.640	1.5
-26+1	2203.669	2203.697	3.2
9.0+1	2680.080	2080.093	4.9
a°+1	459.203	459.204	3.4
a°+1	643.287	643.290	4.4
a ² +1	//2.330	1110 461	2.9
a12	1118.470	1118.461	-8.4
a13	1403.603	1403.608	3.5
a''	1653.698	1653.703	3.0
a ^y	//1.322	//1.322	-0.2
DIO	962.380	962.382	2.1
D''	1049.412	1049.415	2.5
b13	1247.513	1247.516	2.8
D14	1334.545	1334.551	4.4
D10	1518.630	1518.637	5.0
b1/	1681.693	1681.699	3.6
b ²⁰	2046.792	2046.810	8.8
b ²¹	2133.824	2133.831	3.2
b ²³	2359.919	2359.930	4.7
b ²⁴	2522.983	2522.994	4.4
b ²⁶	2707.067	2707.086	7.0
b ₆	486.190	486.191	3.4
C13	1264.539	1264.531	-6.6
C ¹⁶	1535.656	1535.662	3.9
C ²³	2376.945	2376.948	1.0
C ⁶	503.216	503.217	2.8
C ⁹	816.343	816.346	3.8
x ¹⁹ +1	2182.838	2182.840	0.8
x ²³ +1	2554.985	2554.981	-1.5
x4+1	493.193	493.195	2.7
Y ¹⁰	1144.433	1144.437	4.1
y ¹²	1394.528	1394.539	7.8
y ¹³	1491.581	1491.585	2.8
y13-1	1490.573	1490.568	-3.2
y ¹⁴	1578.613	1578.618	3.1
y ¹⁶	1776.713	1776.719	3.5
y ¹⁶ -1	1775.705	1775.710	2.5
y ¹⁷	1863.745	1863.753	4.3
¥ ¹⁸	2026.808	2026.809	0.2
y ²⁰ -1	2251.896	2251.904	3.6
y ²¹	2339.936	2339.947	5.0
y ³	303.143	303.144	2.8
y ⁴	466.206	466.208	3.3
y 5	595.249	595.250	2.2
, У ⁶	692.302	692.304	3.4
¥7	779.334	779.337	4.0
y ⁹	977.434	977.437	3.0
Z ¹⁸	2010.790	2010.807	8.6

2010.790 2010.807

8.6

3	NGPGSÌG 20]P]T⊂	Μ <mark>ί</mark> ςίρ ε <mark>ί</mark> γίςἰρ]τ	SPLS YLS P	T <mark>\S\P\E</mark> \Y\S 25
	Identified	Theoretical	Observed	Mass Difference
	lonsx	Mass	Mass	(ppm)
	a11+1	1022.425	1022.426	0.6
	a12+1	1119.478	1119.479	0.5
	a ¹⁸ +1	1821.704	1821.706	0.8
	a ²¹ +1	2106.837	2106.847	4.9
	a ²² +1	2203.889	2203.895	2.4
	a ²⁶ +1	2680.080	2680.087	2.6
	a ¹⁰	934.385	934.391	5.8
	a ¹³	1219.518	1219.520	2.0
	a ¹⁴	1386.516	1386.520	2.8
	a ²⁵	2582.020	2582.030	3.9
	b ¹⁰	962.380	962.383	3.2
	b ¹¹	1049.412	1049.416	3.7
	b13	1247.513	1247.517	3.5
	b ¹⁶	1598.596	1598.586	-6.2
	b ¹⁷	1761.659	1761.668	5.2
	b ¹⁸	1848.691	1848.700	4.6
	b ²⁰	2046.792	2046.805	6.3
	b ²¹	2133.824	2133.830	3.1
	b ²⁴	2522.983	2522.995	4.8
	b4	298.128	298.129	5.3
	b₀	486.190	486.191	3.3
	C ¹⁶	1615.622	1615.630	4.5
	C ²³	2376.945	2376.951	2.3
	C ₆	503.216	503.219	5.4
	C ⁹	816.343	816.347	4.3
	x ¹⁶ +1	1803.700	1803.695	-3.0
	x ²³ +1	2554.985	2554.981	-1.6
	V ¹⁰	1064.466	1064.471	4.7
	v ¹¹	1227.530	1227.533	2.9
	v ¹²	1314.562	1314.560	-0.8
	v ¹³	1411.614	1411.621	4.4
	v ¹⁴	1578.613	1578.621	5.0
	v ¹⁶	1776.713	1776.720	3.8
	v ¹⁷	1863 745	1863,754	4.8
	v ²⁰ -1	2251.896	2251.903	3.3
	V ²¹	2339 936	2339 949	5.4
	v ²³ -1	2526.990	2527.006	6.3
	v ³	303.143	303.144	3.0
	v4	466.206	466.208	4.4
	v ⁵	595,249	595,251	3.5
	v ⁶	692.302	692.304	3.5
	v ⁷	779.334	779.337	4.6
	v ⁹	977.434	977.438	3.4
	z ¹⁸	2010.790	2010.808	9.2
	-			

dentified	Theoretical	Observed	Mass
lonsx	Mass	Mass	(nnm)
a ¹⁸ +1	1821 704	1821 692	-6.7
a ²² +1	2203 889	2203 900	47
a ²³ +1	2332 932	2332 950	7.5
a ¹⁷	1733 664	1733 672	4.2
a ²¹	2105 829	2105 831	11
a ²⁵	2582 020	2582 030	3.9
h ¹⁰	1042 347	1042 349	21
h11	1179 379	1179 381	2.1
h13	1327 479	1327 488	6.6
h14	1414 511	1414 514	2.0
h17	1761 659	1761 663	2.0
h18	1848 691	1848 701	5.0
h20	2046 792	2046 802	5.0
h ²¹	2133 824	2133 829	23
b ²³	2359 919	2359 940	87
	2522 983	2522 992	3.6
h ²⁵	2610 015	2610 026	43
b ⁶	486 190	486 191	3.4
C ¹³	1344.505	1344.505	-0.3
C ¹⁶	1615.622	1615.629	4.0
C ²⁴	2540.009	2540.011	0.8
C ⁹	896.310	896.312	3.1
V ¹⁰	1064.466	1064.470	3.6
y ¹¹	1227.530	1227.533	2.9
y ¹²	1314.562	1314.549	-9.4
y ¹³	1411.614	1411.621	4.7
y ¹³ -1	1410.607	1410.615	5.9
y ¹⁴	1498.646	1498.654	5.0
y ¹⁶	1696.747	1696.743	-2.0
Y ¹⁷	1783.779	1783.785	3.7
y ²⁰ -1	2171.930	2171.936	3.0
y ²¹	2339.936	2339.949	5.7
y ³	303.143	303.144	2.9
Уз	303.143	303.144	2.9
У ⁴	466.206	466.207	2.2
y ⁴	466.206	466.207	2.2
Y ⁶	692.302	692.304	3.7
y ⁷	779.334	779.337	3.7
Y ⁹	977.434	977.437	3.4

4

Supplementary Figure 10: Lists of fragment ions for mono-phosphorylated peptides (m/z of 943.40) analyzed in Figure 6C by UVPD-MS. In each case, the 3+ charge state was selected, and UVPD was performed using 2 pulses (1.5 mJ per pulse). The identified site of phosphorylation is shaded in blue in the sequence map above each table.

r	N G P G S G MISIPISIYISIPITISIPIEIYISIP TISIP SIYIS
/	20 PT

N G P G S G I 20]P]T ⊂	M <mark>]S\P\S</mark> Y <mark>\S\P</mark> I	rlslp]elvlslp	τ ζεζΡ εζΥ<mark>δ</mark>≃	3	N G P G S G ∞]P]T ⊂	MÌSĮPÌSÌY <mark>S</mark> ĮP I	lslplelylslpl	,τ <mark>ί</mark> ς]Ρ ς γ
Identified Ions	Theoretical mass	Observed mass	Mass Difference (ppm)	_	Identified	Theoretical	Observed	Mas Differe
212+1	1077 469	1077 475	6.9	-	011+1	1060 291	1060 292	(ppn
a+1 a15±1	1362 600	1362 606	0.8		a+1	1245 514	1245 506	0.5
a+1	2628.000	2620 077	4.0		a-*+1	1345.514	1442 574	-5.0
a=-+1	2038.070	642 200	2.5		a+1	1442.500	1442.574	5.0
a°+1	043.207	045.290	4.0		a10+1	1571.609	15/1.611	1.2
d	979.407	3/3.403	1.7		a10+1	1821.704	1821.707	1.2
a17	1072.000	1052 705	-6.0		a13+1	1918.757	1918.766	4.4
a19	1053.098	1740 720	4.3		a21+1	2106.837	2106.839	1.0
a ¹⁰	1740.730	1/40./39	5.4		a20+1	2638.070	2638.079	3.6
a23	2540.009	2540.023	5.4		a°+1	459.203	459.206	1.1
D10	920.370	920.374	4.1		a*+1	643.287	643.289	2.5
D11	1007.402	1007.406	4.3		a ¹³	1257.474	1257.482	6.6
D13	1205.502	1205.507	4.0		a17	1733.664	1733.678	7.7
b ¹⁴	1292.534	1292.540	4.3		a ²⁵	2540.009	2540.017	3.2
b15	1389.587	1389.591	3.0		b ¹⁰	920.370	920.373	3.3
b16	1518.630	1518.637	4.5		b ¹⁴	1372.501	1372.506	3.8
b17	1681.693	1681.699	3.8		b ¹⁶	1598.596	1598.598	1.0
b ¹⁸	1768.725	1768.733	4.3		b17	1761.659	1761.666	3.6
b ²⁰	1966.825	1966.835	4.7		b ¹⁸	1848.691	1848.697	3.3
b ²¹	2053.857	2053.866	3.9		b ²¹	2133.824	2133.830	3.0
b ²³	2237.942	2237.958	6.9		b ²⁴	2480.972	2480.979	2.7
b ²⁴	2401.006	2401.016	4.1		b ²⁵	2568.004	2568.015	4.1
b ²⁵	2568.004	2568.017	5.2		b ²⁶	2665.057	2665.076	7.0
be	486.190	486.191	3.6		b ⁶	486.190	486.191	3.0
b7	573.222	573.224	3.6		b ⁹	757.306	757.308	2.6
b9	757.306	757.311	5.6		C13	1302.495	1302.500	4.1
C10	937.396	937.399	3.0		C ¹⁶	1615.622	1615.628	3.7
C13	1222.529	1222.531	1.7		C ⁸	687.301	687.306	8.0
C ¹⁶	1535.656	1535.663	4.5		C ⁹	774.333	774.335	3.3
C ⁶	503.216	503.218	3.1		x9+1	962.411	962.417	6.3
C ⁹	774.333	774.336	4.3		X ⁸	864.350	864.351	0.6
x ¹⁹ +1	2140.828	2140.834	3.0		V ¹⁰	1022.456	1022.460	4.0
y ¹⁰	1102.422	1102.427	4.4		V ¹¹	1185.519	1185.525	5.3
y ¹¹	1265.485	1265.494	7.2		V ¹³	1411.614	1411.619	3.4
y ¹³	1491.581	1491.586	3.8		v ¹⁴	1498.646	1498.650	2.5
Y13-1	1490.573	1490.565	-5.3		Y ¹⁶ -1	1695.739	1695.741	1.2
y ¹⁴	1578.613	1578.617	2.9		V ¹⁷	1863.745	1863.749	2.0
y ¹⁶	1776.713	1776.719	3.5		v ²⁰	2210.893	2210.898	2.1
V ¹⁷	1863.745	1863.749	2.3		v ²¹	2297,925	2297,935	4.3
V ²⁰	2210.893	2210.898	2.1		y3	303.143	303.144	2.6
V ⁴	546.173	546.175	4.3		V ⁶	650.291	650.293	3.4
V ⁶	730.257	730.260	3.1		v7	737.323	737.326	3.4
v ⁷	817.290	817.292	3.5		, v ⁸	838 371	838 373	2.0
y9	1015.390	1015.394	3.7		7 V ⁹	935 424	935 427	2.0
,	_010.000	_010.004	5		y y9-1	934 416	934 420	4.6
					y - 1 719	2097 822	2007 816	-3.0

Supplementary Figure 11: Lists of fragment ions for mono-phosphorylated peptides (m/z of 929.39) analyzed in Figure 6D by UVPD-MS. In each case, the 3+ charge state was selected, and UVPD was performed using 2 pulses (1.5 mJ per pulse). The identified site of phosphorylation is shaded in blue in the sequence map above each table.

dentified	Theoretical	Observed	Mass Difference
ions	mass	mass	(ppm)
a**+1	893.383	893.384	1.1
a''+1	1022.425	1022.427	1.3
a ¹³	1219.518	1219.507	-9.3
b10	920.370	920.372	2.4
b11	1049.412	1049.415	2.9
b12	1146.465	1146.463	-1.7
b ¹³	1247.513	1247.517	3.1
b ¹⁴	1334.545	1334.550	4.1
b ¹⁵	1431.598	1431.603	3.7
b ¹⁶	1518.630	1518.637	4.8
b17	1681.693	1681.700	3.9
b ¹⁸	1810.736	1810.742	3.7
b ²⁰	2008.836	2008.846	4.8
b ²¹	2175.834	2175.847	5.8
b ²⁴	2522.983	2522.992	3.6
b ²⁵	2652.025	2652.035	3.6
b ₆	486.190	486.191	2.2
c ¹⁰	937.396	937.400	3.6
c ¹²	1163 491	1163 491	-0.1
c ¹³	1264 539	1264 543	3.4
C ¹⁶	1535 656	1535 662	4 1
c ²³	2376 945	2376 038	. 2 7
ر ح6	502 216	E02 210	-5.2
د ح ⁸	697 201	697 202	10
ر م	774 222	774 225	1.5
L 10	1196 442	1106 110	3.0
y .	1240 500	1240 512	3.9
y	1549.506	1549.512	4.5
y-3	1533.591	1533.599	5.0
y."	1533.591	1533.595	2.8
y''+1	1532.583	1532.588	3.1
У ¹⁴	1620.623	1620.631	4.5
Y10	1818.724	1818.729	3.0
y20	2294.914	2294.922	3.4
y ²¹	2381.946	2381.953	2.6
У ³	345.154	345.154	2.1
y ⁴	508.217	508.218	2.7
y ⁶	692.302	692.304	3.2
y7	859.300	859.301	1.6
y ⁹	1057.401	1057.404	2.8
z ³	329.135	329.138	9.9

329.138

dentified	Theoretical	Observed	Mass Difference
lons	mass	mass	(ppm)
a ¹¹ +1	1022.425	1022.419	-6.4
a14+1	1387.524	1387.516	-5.8
a ¹⁰	892.375	892.379	4.5
a ¹³	1219.518	1219.508	-8.0
a ²⁵	2624.030	2624.044	5.2
b ¹⁰	920.370	920.372	2.2
b11	1049.412	1049.416	3.6
b ¹³	1247.513	1247.516	2.8
b ¹⁶	1598.596	1598.604	4.7
b17	1761.659	1761.666	3.7
b ¹⁸	1890.702	1890.708	3.2
b ²⁰	2088.802	2088.803	0.2
b ²⁴	2522.983	2522.990	2.9
b ²⁵	2652.025	2652.035	3.8
b ⁹	757.307	757.309	3.6
C12	1163.491	1163.490	-1.6
c ¹⁶	1615.622	1615.629	4.2
C23	2376.945	2376.946	0.1
c ⁹	774.333	774.335	3.0
X ⁸	906.361	906.367	6.8
x ⁹	1003.413	1003.405	-8.2
V ¹⁰	1106.477	1106.483	5.3
v ¹⁰	1106.477	1106.480	2.8
y ¹²	1356.572	1356.567	-3.4
y ¹³	1453.625	1453.635	7.1
y ¹³ -1	1452.617	1452.613	-2.5
v ¹⁴	1620.623	1620.630	4.1
v ¹⁶	1818.724	1818.730	3.2
v ¹⁶ -1	1817.716	1817.718	1.4
v ²⁰	2294.914	2294.925	4.6
21	2381 946	2381.952	2.2
V ²¹	LUU1.010		
y ²¹ y ³	345.154	345.154	2.2
γ ²¹ γ ³ γ ⁵	345.154 595.249	345.154 595.251	2.2 3.8

779.334 779.337

977.434 977.437

3.8

2.7

4	N G P G S G I 20]P T C	W <mark>S</mark> PS]Y]E <u>)</u> PT	ls <mark>lpislyl</mark> eip 1	r <mark>lslpls]yl</mark> e 🕾
	Identified Ions	Theoretical mass	Observed mass	Mass Difference (ppm)
	a ¹⁰ +1	973 349	973 348	-1.6
	a ¹¹ +1	1102 392	1102 388	-3.7
	a ²²	2244.892	2244.913	9.1
	b ¹⁰	1000.336	1000.339	2.4
	b ¹¹	1129.379	1129.381	1.9
	b ¹⁴	1414.511	1414.513	1.6
	b^{16}	1598.596	1598.610	8.7
	b17	1761.659	1761.665	3.4
	b ¹⁸	1890.702	1890.706	2.0
	b ²⁰	2088.802	2088.803	0.1
	b ²³	2359.919	2359.937	7.5
	b ²⁴	2522.983	2522.988	2.3
	b ²⁵	2652.025	2652.033	3.2
	b ⁶	486.190	486.191	2.6
	b7	653.188	653.189	2.1
	C ¹³	1344.505	1344.513	5.5
	C ¹⁶	1615.622	1615.628	3.4
	C ²³	2376.945	2376.950	2.1
	C ⁹	854.299	854.301	1.8
	y ¹⁰	1106.477	1106.479	2.2
	y ¹¹	1269.540	1269.543	2.4
	y ¹²	1356.572	1356.574	1.6
	y ¹³	1453.625	1453.630	3.4
	y ¹⁴	1540.657	1540.663	3.7
	y ¹⁶	1738.757	1738.762	2.4
	y ²¹	2381.946	2381.950	1.4
	y ³	345.154	345.154	1.9
	y ⁵	595.249	595.250	1.9
	y ⁶	692.302	692.304	2.5
	y ⁷	779.334	779.336	3.3
	γ ⁹	977.434	977.437	2.5
	z ¹⁰	1090.458	1090.465	6.3
	z ³	329.135	329.138	9.9

Supplementary Figure 12: Lists of fragment ions for mono-phosphorylated peptides (m/z of 957.05) analyzed in Figure 6E by UVPD-MS. In each case, the 3+ charge state was selected, and UVPD was performed using 2 pulses (1.5 mJ per pulse). The identified site of phosphorylation is shaded in blue in the sequence map above each table.

γ⁷ γ⁹

NGPGSG ≈]PT⊂	ΜͺϛͺϷϡϛͺϒͺϛͺϷ;ϫ	lslp slylelp	TÌSÌP SÌY <mark>S</mark> ≃
Identified	Theoretical	Observed	Mass Difference
lons	mass	mass	(ppm)
a ¹¹ +1	980 415	980 411	-3.9
a ¹⁴ +1	1265 547	1265 539	-63
a ⁸ +1	643.287	643.288	1.1
a ¹⁰	892.375	892.366	-10.2
a ¹²	1076.460	1076.455	-4.4
a ¹³	1177.507	1177.497	-9.0
a ¹⁸	1740.730	1740.718	-6.8
a ²¹	2025.863	2025.872	4.8
b11	1007.402	1007.402	0.4
b ¹³	1205.502	1205.505	2.6
b ¹⁴	1292.534	1292.538	2.6
b ¹⁶	1476.619	1476.627	5.5
b17	1639.682	1639.687	2.6
b ¹⁸	1768.725	1768.726	0.8
b ²⁰	1966.825	1966.832	3.3
b ²⁴	2401.006	2401.011	2.1
b ²⁵	2568.004	2568.013	3.4
b ⁹	757.307	757.310	4.5
C ¹⁰	937.396	937.400	4.0
C ¹⁶	1493.645	1493.651	3.7
c ²³	2254.969	2254.976	3.2
c ⁸	687.301	687.302	2.3
C ⁹	774.333	774.335	2.2
y ¹⁰	1144.433	1144.435	2.6
y ¹¹	1307.496	1307.500	3.3
y ¹³	1491.581	1491.583	1.6
y ¹³	1491.581	1491.586	3.8
y ¹⁴	1578.613	1578.615	1.5
y ¹⁶	1776.713	1776.719	3.0
y ¹⁷	1863.745	1863.749	2.1
y ²⁰	2210.893	2210.901	3.7
y ²¹	2297.925	2297.933	3.5
γ ³	383.109	383.110	1.2
Y ⁶	730.258	730.259	1.8
¥ ⁷	817.290	817.292	2.6
y ⁹	1015.390	1015.392	1.9

817.290	817.292	2.6	
1015.390	1015.392	1.9	

3	Ν σρο s ο μίς[ρ]ς]γ<mark>ς</mark>[ρ]τ]ς[ρίς]γ]ε[ρίτ]ς]ρ s]γ] s 20]ρ τ ο				
	Identified Ions	Theoretical mass	Observed mass	Mass Difference (ppm)	
	a12+1	1157.434	1157.441	6.1	
	a ¹⁴ +1	1345.514	1345.508	-4.1	
	b ¹⁰	920.370	920.372	2.3	
	b ¹³	1285.469	1285.471	2.2	
	b ¹⁴	1372.501	1372.505	3.5	
	b ¹⁶	1556.585	1556.588	1.4	
	b ¹⁷	1719.649	1719.653	2.5	
	b ¹⁸	1848.691	1848.698	3.5	
	b ²⁰	2046.792	2046.797	2.7	
	b ²¹	2133.824	2133.830	2.7	
	b ²⁴	2480.972	2480.977	1.9	
	b ²⁵	2568.004	2568.013	3.7	
	b ⁹	757.307	757.309	2.9	
	C ¹³	1302.495	1302.501	4.7	
	C ¹⁶	1573.612	1573.616	2.9	
	C ²³	2334.935	2334.940	2.1	
	c ⁸	687.301	687.303	3.2	
	c ⁹	774.333	774.335	2.2	
	x ⁸ +1	865.358	865.359	0.7	
	y ¹⁰	1064.466	1064.469	2.7	
	y ¹¹	1227.530	1227.531	1.5	
	y ¹²	1314.562	1314.560	-1.6	
	y ¹³	1411.614	1411.619	3.2	
	y ¹⁴	1498.646	1498.653	4.2	
	y ¹⁶ -1	1695.739	1695.741	1.3	
	y ¹⁷	1863.745	1863.749	2.1	
	y ¹⁸	2026.808	2026.820	5.5	
	y ²⁰	2210.893	2210.900	3.2	
	y ²¹	2297.925	2297.933	3.4	
	γ ⁶	650.291	650.293	2.7	
	y ⁷	737.323	737.325	2.7	
	٧ ⁸	838.371	838.372	0.7	
	y ⁹	935.424	935.425	1.4	
	z ¹⁹	2097.822	2097.811	-5.0	

2097.822 2097.811

-5.0

T	N G P G S G M <mark>S</mark> P[S]Y[S]P T[S[P S[Y]E[P[T]S[P S[Y]S 24]P T C					
				Mass		
	Identified	Theoretical	Observed	Difference		
	lons	mass	mass	(ppm)		
	a''+1	973.349	973.357	8.1		
	p ₁₀	1000.336	1000.338	2.0		
	b''	1087.368	1087.370	1.4		
	b ¹³	1285.469	1285.472	2.6		
	b ¹⁴	1372.501	1372.505	2.9		
	b ¹⁶	1556.585	1556.595	6.2		
	b ¹⁷	1719.649	1719.653	2.5		
	b ¹⁸	1848.691	1848.698	3.7		
	b ²⁰	2046.792	2046.798	3.2		
	b ²¹	2133.824	2133.829	2.5		
	b ²⁴	2480.972	2480.978	2.6		
	b ²⁵	2568.004	2568.013	3.7		
	b6	486.190	486.191	1.7		
	C ¹⁶	1573.612	1573.617	3.0		
	C ²³	2334.935	2334.941	2.6		
	C ⁹	854.299	854.302	3.1		
	y ¹⁰	1064.466	1064.469	2.7		
	y ¹¹	1227.530	1227.531	1.3		
	v ¹³	1411.614	1411.617	1.9		
	y ¹⁴	1498.646	1498.651	3.3		
	V ¹⁷	1783.779	1783.784	2.9		
	y ²⁰	2130.927	2130.943	7.3		
	v ²¹	2297.925	2297.935	4.3		
	v ⁴	466.206	466.207	1.8		
	v ⁶	650.291	650.293	3.0		
	y ⁷	737.323	737.326	3.3		
	v ⁸	838.371	838.369	-2.4		
	v9	935.424	935.426	3.0		
	z ¹⁹	2017.855	2017.843	-6.0		

Supplementary Figure 13: Lists of fragment ions for mono-phosphorylated peptides (m/z of 929.05) analyzed in Figure 6F by UVPD-MS. In each case, the 3+ charge state was selected, and UVPD was performed using 2 pulses (1.5 mJ per pulse). The identified site of phosphorylation is shaded in blue in the sequence map above each table.