Peptide array: Histone H3(unmodified K27) (ab308248)
Area under curve (AUC) data for all peptides on the pad. Antigen-containing peptides are highlighted in green.

Cell Ref	AUC	Peptide
A1	6.37091875	ab10112 - H4 K91ac
A2	8.027561458	ab10121 - H2B K85ac
A3	5.374721875	ab10127 - H2A K95me2
A4	4.225045833	ab10138 - H1.4 T17p
A5	7.652720833	ab10139 - H1.4 T146p
A6	4.18680625	ab113997 - H2A.X Y142p
A7	5.649839583	ab11477 - H3 S10p
A8	6.354292708	ab12946 - H2B K5me3
A9	4.539488542	ab12949 - H2B K23me2
A10	4.7001125	ab12952 - H2A K127me1
A11	6.4725125	ab13211 - H2B K5me1
A12	6.12850625	ab1340 - H3 K4me1
A13	6.713865625	ab13570 - H2B K43me1
A14	6.426552083	ab13835 - HA
A15	5.273621875	ab13837 - Myc
A16	7.27958125	ab140408 - ScH2B T129p
A17	6.201865625	ab14103 - H3 T6p
A18	7.980865625	ab14724 - H4 S1p
A19	19.50888333	ab14799 - H3 T32p
A20	6.00874375	ab14943 - His
A21	4.721244792	ab14952 - H2A R77me2s
A22	5.738619792	ab14964 - H4 K20me2
A23	4.805257292	ab14997 - ScH2A K21ac
A24	5.349023958	ab154782 - H3 R26me2s
A25	6.957242708	ab15591 - H3 K14ac
B1	5.453926042	ab15644 - H3 K9me3 + S10p
B2	16.35892917	ab15645 - H2A.X S139p
B3	8.952439583	ab15824 - H4 K8ac
B4	6.043959375	ab15829 - V5
B5	6.653970833	ab16635 - H3 K9ac
B6	6.80074375	ab166685 - H2B S112GlcNAc
B7	7.23228125	ab166688 - H2B S36GlcNAc
B8	4.30350625	ab16935 - H3 R17me2a
B9	6.257771875	ab17043 - H4 K20me1
B10	64.72025	ab17162 - H3 P16hyd + P30hyd
B11	16.35427188	ab17567 - H4 K20me3
B12	6.249104167	ab17587 - H1.4 K25me3
B13	6.81343125	ab175905 - H2B K11ac
B14	5.974263542	ab176211 - H2B K23ac
B15	6.0953375	ab17632 - H4 K5bio
B16	6.178903125	ab1771 - H3 K9me1
B17	5.752758333	ab1773 - H3 K9me3
B18	6.278698958	ab17770 - H4 R3me1
B19	5.411166667	ab1781 - H3 K27me2
B20	5.567390625	ab1783 - H3 K36me1
B21	6.9009375	ab1784 - H3 K36me2
B22	5.486441667	ab1785 - H3 K36me3
B23	5.498907292	ab178667 - ScH2A unmod(1-30)
B24	4.794697917	ab178668 - ScH2A unmod(102-132)
B25	4.91664375	ab178670 - H2A.Z unmod(1-30)
C1	5.487971875	ab178833 - Flag
C2	5.340513542	ab178835 - H2B R99mutO
C3	6.433161458	ab178836 - H4 R23mutO
C4	6.263097917	ab178837 - H4 G48mutD
C5	5.46551875	ab178987 - H1.4 unmod(1-30)
C6	5.792870833	ab178988 - H1.4 S1p
C7	5.397802083	ab178989 - H1.4 T3p
C8	4.999470833	ab178990 - H1.4 S1p + T3p
C9	4.80424375	ab178991 - H1.4 K25me1
C10	6.672728125	ab178992 - H1.4 K25me3 + S26p
C11	8.834325	ab178993 - H1.4 unmod(130-160)
C12	7.4217	ab178994 - H2A unmod(1-30)
C13	6.586602083	ab178995 - H2A S1p
C14	7.529219792	ab178996 - H2A S1p + K5ac
C15	7.361159375	ab178997 - H2A S1p + R3me2a + K5ac
C16	6.138538542	ab178998 - H2A S1p + R3me2s + K5ac
C17	4.439557292	ab178999 - H2A S1p + R3cit + K5ac
C18	4.021201042	ab179000 - H2A S1ac + S1p + K5ac + K9ac + K13ac + K15ac
C19	3.638151042	ab179001 - H2A S1p + K5ac + K9ac + K13ac + K15ac
C20	5.619785417	ab179002 - H2A R3cit

C21	5.098833333	ab179003 - H2A R3cit + K5ac
C22	4.24045625	ab179004 - H2A R3me2a + K5ac
C23	5.57073125	ab179005 - H2A R3me2s + K5ac
C24	7.154603125	ab179006 - H2A S1ac + K5ac + K9ac + K13ac + K15ac
C25	4.285572917	ab179007 - H2A K5ac + K9ac + K13ac + K15ac
D1	4.354765625	ab179008 - H2A K9me2
D2	7.6872875	ab179009 - H2A K9me3
D3	3.890734375	ab179010 - H2A R11me1
D4	4.961632292	ab179011 - H2A R11me2s
D5	4.04129375	ab179012 - H2A R11me2a
D6	6.156863542	ab179013 - H2A K13ac
D7	3.85899375	ab179014 - H2A K15ac
D8	5.179804167	ab179015 - H2A R17me1
D9	7.4250875	ab179016 - H2A R17me2s
D10	7.277366667	ab179017 - H2A R17me2a
D11	7.175778125	ab179018 - H2A unmod(20-50)
D12	7.541623958	ab179019 - H2A P26hyd
D13	8.12291875	ab179020 - H2A R29me1
D14	6.976813542	ab179021 - H2A R29me2s
D15	9.104867708	ab179022 - H2A R29me2a
D16	7.013528125	ab179023 - H2A K36ac
D17	6.187252083	ab179024 - H2A K36cr
D18	3.848072917	ab179025 - H2A Y39hyd
D19	4.564840625	ab179026 - H2A R42me1
D20	4.307427083	ab179027 - H2A R42me2s
D21	5.678880208	ab179028 - H2A R42me2a
D22	3.981744792	ab179029 - H2A unmod(71-103)
D23	5.361397917	ab179030 - H2A R77me1
D24	5.3545375	ab179031 - H2A R77me2a
D25	5.559032292	ab179032 - H2A R88me1
E1	5.439289583	ab179033 - H2A R88me2s
E2	5.334611458	ab179034 - H2A R88me2a
E3	6.529988542	ab179035 - H2A K95me1
E4	4.80948125	ab179036 - H2A K95me3
E5	6.595301042	ab179037 - H2A unmod(92-123)
E6	3.533417708	ab179038 - H2A T101GlcNAc
E7	5.190291667	ab179039 - H2A Q104me1
E8	4.852547917	ab179040 - H2A unmod(112-129)
E9	4.922803125	ab179041 - H2A K118me1
E10	4.470890625	ab179042 - H2A K118me2
E11	6.27054375	ab179043 - H2A K118me3
E12	4.210292708	ab179044 - H2A K118fo
E13	5.686459375	ab179045 - H2A K118cr
E14	6.982986458	ab179046 - H2A K119cr
E15	7.077117708	ab179047 - H2A K118ubiq
E16	7.169570833	ab179048 - H2A K119ubiq
E17	6.884866667	ab179049 - H2A T120p
E18	7.135246875	ab179050 - H2A K125me1
E19	7.019227083	ab179051 - H2A K125me2
E20	4.928640625	ab179052 - H2A K125cr
E21	6.017842708	ab179053 - H2A.X unmod(1-30)
E22	4.649963542	ab179054 - H2A.X K5ac
E23	4.211308333	ab179055 - H2A.X unmod(114-142)
E24	4.1603625	ab179056 - H2A.X K118ubiq
E25	4.616648958	ab179057 - H2A.X K119ubiq
F1	5.062490625	ab179058 - H2A.X T120p
F2	5.127565625	ab179059 - H2A.X K127ubiq
F3	5.014023958	ab179060 - H2A.X S139p + Y142p
F4	6.359372917	ab179061 - H2A.Z K4ac
F5	4.470666667	ab179062 - H2A.Z K7ac
F6	4.705484375	ab179063 - H2A.Z K11ac
F7	5.447941667	ab179064 - H2A.Z K4ac + K7ac + K11ac
F8	4.61843125	ab179065 - H2B unmod(1-30)
F9	4.1065125	ab179066 - H2B K5ac
F10	5.359833333	ab179067 - H2B K5cr
F11	4.449427083	ab179068 - H2B K5fo
F12	5.404942708	ab179069 - H2B P10hyd
F13	5.294679167	ab179070 - H2B K11me1
F14	7.09523125	ab179071 - H2B K11me2
F15	5.019921875	ab179072 - H2B K11me3
F16	6.95453125	ab179073 - H2B K11cr
F17	5.52658125	ab179074 - H2B K12ac
F18	5.696139583	ab179075 - H2B K12cr
F19	5.960397917	ab179076 - H2B K12me1

F20	4.952	ab179077 - H2B K12me2
F21	6.299215625	ab179078 - H2B K12me3
F22	5.568441667	ab179079 - H2B K12ac + K15ac
F23	7.190377083	ab179080 - H2B S14p
F24	6.504107292	ab179081 - H2B K15ac
F25	7.723080208	ab179082 - H2B K15cr
G1	5.848909375	ab179083 - H2B K15me1
G2	6.788966667	ab179084 - H2B K15me2
G3	5.963782292	ab179085 - H2B K15me3
G4	4.987961458	ab179086 - H2B K16ac
G5	6.091458333	ab179087 - H2B K16cr
G6	4.004234375	ab179088 - H2B K20ac
G7	3.194492708	ab179089 - H2B K20cr
G8	3.969611458	ab179090 - H2B K20me1
G9	6.035257292	ab179091 - H2B K20me2
G10	5.6459375	ab179092 - H2B K20me3
G11	4.100215625	ab179093 - H2B K23cr
G12	6.265594792	ab179094 - H2B K23me1
G13	6.392420833	ab179095 - H2B K23me3
G14	2.741209375	ab179096 - H2B K24ac
G15	8.865471875	ab179097 - H2B unmod(26-56)
G16	6.590396875	ab179098 - H2B K34cr
G17	5.579323958	ab179099 - H2B K34fo
G18	4.829167708	ab179100 - H2B Y37hyd
G19	6.626454167	ab179101 - H2B Y37p
G20	6.029782292	ab179102 - H2B K43me2
G21	5.095585417	ab179103 - H2B K43me3
G22	6.893105208	ab179104 - H2B K46fo
G23	7.46715	ab179105 - H2B K46ac
G24	5.636216667	ab179106 - H2B K46me1
G25	7.554828125	ab179107 - H2B K46me2
H1	4.968944792	ab179108 - H2B K46me3
H2	7.626683333	ab179109 - H2B unmod(51-81)
H3	5.701783333	ab179110 - H2B K57me1
H4	6.50050625	ab179111 - H2B K57me2
H5	6.081325	ab179112 - H2B K57me3
H6	5.004541667	ab179113 - H2B unmod(73-103)
H7	5.990611458	ab179114 - H2B R79me1
H8	5.451347917	ab179115 - H2B R79me2s
H9	4.749253125	ab179116 - H2B R79me2a
H10	5.246041667	ab179117 - H2B Y83hyd
H11	4.395695833	ab179118 - H2B K85me1
H12	5.920238542	ab179119 - H2B K85me2
H13	5.409257292	ab179120 - H2B K85me3
H14	6.60823125	ab179121 - H2B unmod(94-125)
H15	4.46388125	ab179122 - H2B R99me1
H16	6.788878125	ab179123 - H2B R99me2s
H17	5.535601042	ab179124 - H2B R99me2a
H18	5.989244792	ab179125 - H2B K108fo
H19	4.99946875	ab179126 - H2B K116fo
H20	6.860383333	ab179127 - H2B K116me1
H21	5.871166667	ab179128 - H2B K116me2
H22	5.888070833	ab179129 - H2B K116me3
H23	5.94758125	ab179130 - H2B K116ubiq
H24	6.945686458	ab179131 - H2B K120ac
H25	5.757559375	ab179132 - H2B K120fo
I1	6.119611458	ab179133 - H2B K120ubiq
I2	8.20403125	ab179134 - H2B K125ubiq
I3	6.592308333	ab179135 - ScH2B unmod(102-132)
I4	79.17770104	ab179136 - H3 unmod(1-33)
I5	6.324439583	ab179137 - H3 R2cit
I6	9.906805208	ab179138 - H3 R2cit + K4me2
I7	6.287771875	ab179139 - H3 R2cit + K4me3
I8	5.879047917	ab179140 - H3 R2cit + K4me3 + K9ac + K14ac + K18ac
I9	7.438354167	ab179141 - H3 R2cit + K4ac + K9ac + K14ac + K18ac
I10	6.2377375	ab179142 - H3 R2me1
I11	6.298928125	ab179143 - H3 R2me1 + K4me2
I12	6.689077083	ab179144 - H3 R2me1 + K4me3
I13	7.760880208	ab179145 - H3 R2me1 + K4me3 + K9ac + K14ac + K18ac
I14	8.448942708	ab179146 - H3 R2me2s
I15	5.8314125	ab179147 - H3 R2me2s + K4me2
I16	6.452671875	ab179148 - H3 R2me2s + K4me3
I17	6.363872917	ab179149 - H3 R2me2s + K4me3 + K9ac + K14ac + K18ac
I18	7.05954375	ab179150 - H3 R2me2a

I19	7.468996875	ab179151 - H3 R2me2a + K4me2
I20	4.797867708	ab179152 - H3 R2me2a + K4me3
I21	5.807463542	ab179153 - H3 R2me2a + K4me3 + S10p
I22	6.044489583	ab179154 - H3 R2me2a + T3p + K4me3
I23	6.644154167	ab179155 - H3 R2me2a + K4ac + K9ac + K14ac + K18ac
I24	5.139619792	ab179156 - H3 R2me2a + K4me3 + K9ac + K14ac + K18ac
I25	7.403604167	ab179157 - H3 R2me2a + T3p + K4me3 + K9ac + K14ac + K18ac
J1	6.561258333	ab179158 - H3 R2me2a + K4me3 + K9ac + S10p + K14ac + K18ac
J2	4.826015625	ab179159 - H3 T3p
J3	5.327060417	ab179160 - H3 T3p + K4me1
J4	4.611461458	ab179161 - H3 T3p + K4me2
J5	5.43019375	ab179162 - H3 T3p + K4me3 + K9ac + K14ac + K18ac
J6	6.867880208	ab179163 - H3 K4ac
J7	6.357680208	ab179164 - H3 K4ac + K9ac
J8	7.18446875	ab179165 - H3 K4ac + K14ac
J9	6.709380208	ab179166 - H3 K4ac + K18ac
J10	6.626916667	ab179167 - H3 K4ac + K9me3
J11	5.79455	ab179168 - H3 K4ac + K9ac + K14ac
J12	6.722204167	ab179169 - H3 K4ac + K9ac + K18ac
J13	7.71451875	ab179170 - H3 K4ac + K14ac + K18ac
J14	6.318026042	ab179171 - H3 K4ac + K9ac + K14ac + K18ac
J15	6.2459125	ab179172 - H3 K4ac + K9me1 + K14ac + K18ac
J16	5.823189583	ab179173 - H3 K4ac + K9me2 + K14ac + K18ac
J17	7.57640625	ab179174 - H3 K4ac + K9me3 + K14ac + K18ac
J18	7.450455208	ab179175 - H3 K4ac + K9ac + S10p + K14ac + K18ac
J19	6.254957292	ab179176 - H3 K4cr
J20	6.508352083	ab179177 - H3 K4me1 + K9me2
J21	5.501882292	ab179178 - H3 K4me1 + K9ac + K18ac
J22	5.813754167	ab179179 - H3 K4me1 + K9ac + K14ac + K18ac
J23	3.432442708	ab179180 - H3 K4me2 + T6p
J24	4.836739583	ab179181 - H3 K4me2 + K9me2
J25	4.826823958	ab179182 - H3 K4me2 + K9ac + K18ac
K1	6.27530625	ab179183 - H3 K4me2 + K9ac + K14ac + K18ac
K2	7.374747917	ab179184 - H3 K4me3 + T6p
K3	6.392163542	ab179185 - H3 K4me3 + K9ac
K4	5.475201042	ab179186 - H3 K4me3 + K9me2
K5	6.424401042	ab179187 - H3 K4me3 + K9me3
K6	5.677516667	ab179188 - H3 K4me3 + S10p
K7	6.550228125	ab179189 - H3 K4me3 + K14ac
K8	8.367952083	ab179190 - H3 K4me3 + K18ac
K9	7.047819792	ab179191 - H3 K4me3 + R8me2s + K9me3
K10	7.318297917	ab179192 - H3 K4me3 + R8me2a + K9me3
K11	6.902330208	ab179193 - H3 K4me3 + K9ac + S10p
K12	5.650748958	ab179194 - H3 K4me3 + K9ac + K14ac
K13	6.181676042	ab179195 - H3 K4me3 + K9ac + K18ac
K14	7.450947917	ab179196 - H3 K4me3 + K14ac + K18ac
K15	6.917580208	ab179197 - H3 K4me3 + K9ac + K14ac + K18ac
K16	7.63766875	ab179198 - H3 K4me3 + K9ac + S10p + K14ac + K18ac
K17	7.046864583	ab179199 - H3 K4me3 + T6p + K9ac + K14ac + K18ac
K18	5.763329167	ab179200 - H3 K4me2 + T6p + K9ac + K14ac + K18ac
K19	4.5447625	ab179201 - H3 K4bio + K9bio + K18bio
K20	3.683697917	ab179202 - H3 T6p + K9me3
K21	5.061136458	ab179203 - H3 T6p + R8me2a + K9me3
K22	5.177207292	ab179204 - H3 T6p + K9ac + K14ac + K18ac
K23	5.538561458	ab179205 - H3 R8me1
K24	4.589010417	ab179206 - H3 R8me1 + K9me1
K25	5.037892708	ab179207 - H3 R8me1 + K9me2
L1	5.576513542	ab179208 - H3 R8me1 + K9me3
L2	6.543453125	ab179209 - H3 R8me2s
L3	5.991861458	ab179210 - H3 R8me2s + K9me1
L4	4.463165625	ab179211 - H3 R8me2s + K9me2
L5	4.034310417	ab179212 - H3 R8me2s + K9me3
L6	6.769317708	ab179213 - H3 R8me2a
L7	5.380691667	ab179214 - H3 R8me2a + K9me1
L8	6.958228125	ab179215 - H3 R8me2a + K9me2
L9	3.866955208	ab179216 - H3 R8me2a + K9me3
L10	6.736117708	ab179217 - H3 R8cit
L11	5.1125875	ab179218 - H3 K9ac + K14ac
L12	5.927277083	ab179219 - H3 K9ac + K18ac
L13	4.253529167	ab179220 - H3 K9ac + K14ac + K18ac
L14	5.0978875	ab179221 - H3 K9ac + K14ac + K18ac + K23ac + K27ac
L15	4.991416667	ab179222 - H3 K9ac + S10p
L16	5.826428125	ab179223 - H3 K9cr
L17	4.307983333	ab179224 - H3 K9me2

L18	4.535561458	ab179225 - H3 K9me2 + S10p
L19	8.763345833	ab179226 - H3 K9me2 + K27me2
L20	8.072958333	ab179227 - H3 S10p + S28p
L21	6.131003125	ab179228 - H3 S10p K14ac
L22	6.453576042	ab179229 - H3 K14me1
L23	6.073361458	ab179230 - H3 K14me2
L24	8.056698958	ab179231 - H3 K14me3
L25	7.393782292	ab179232 - H3 K14ac + K18ac
M1	7.263726042	ab179233 - H3 R17cit
M2	8.570345833	ab179234 - H3 R17me1
M3	6.630823958	ab179235 - H3 R17me2s
M4	6.521242708	ab179236 - H3 R17me2a + K18ac
M5	6.053248958	ab179237 - H3 K18cr
M6	4.194340625	ab179238 - H3 K18me1
M7	6.940266667	ab179239 - H3 K18me2
M8	142.1090135	ab179240 - H3 K18me3 + K36me3
M9	4.973614583	ab179241 - H3 K23cr
M10	4.727751042	ab179242 - H3 K23me1
M11	8.0232875	ab179243 - H3 K23me2
M12	5.212933333	ab179244 - H3 K23me3
M13	5.767058333	ab179245 - H3 T22p + K23me3
M14	6.63536875	ab179246 - H3 K23pr
M15	172.7927938	ab179247 - H3 unmod(20-51)
M16	27.61215938	ab179248 - H3 R26me1
M17	7.213528125	ab179249 - H3 R26me2a
M18	4.852990625	ab179250 - H3 R26me2a + K27me1
M19	6.092684375	ab179251 - H3 R26me2a + K27me3
M20	6.361564583	ab179252 - H3 K27ac + S28p
M21	8.731975	ab179253 - H3 K27cr
M22	7.564514583	ab179254 - H3 K27me1
M23	6.306921875	ab179255 - H3 K27me2 + S28p
M24	6.478380208	ab179256 - H3 K27me3
M25	7.069858333	ab179257 - H3 K27me3 + S28p
N1	4.841257292	ab179258 - H3 P30hpro
N2	7.072695833	ab179259 - H3 P30(5,5me2)
N3	5.784805208	ab179260 - H3 P30
N4	6.393846875	ab179261 - H3 P30mutA
N5	5.643305208	ab179262 - H3 T32GlcNAc
N6	7.67563125	ab179263 - H3 K36ac
N7	7.418461458	ab179264 - H3 P38hpro
N8	6.681640625	ab179265 - H3 P38(5,5me2)
N9	6.045392708	ab179266 - H3 P38
N10	5.656869792	ab179267 - H3 P38mutA
N11	5.718761458	ab179268 - H3 Y41p
N12	6.410720833	ab179269 - H3 unmod(50-80)
N13	4.9706875	ab179270 - H3 K56ac
N14	7.479838542	ab179271 - H3 K56cr
N15	4.446342708	ab179272 - H3 K56fo
N16	4.435104167	ab179273 - H3 K56me1
N17	5.572830208	ab179274 - H3 K56me2
N18	4.463723958	ab179275 - H3 K56me3
N19	6.463197917	ab179276 - H3 R63me1
N20	4.6873125	ab179277 - H3 R63me2s
N21	7.744944792	ab179278 - H3 R63me2a
N22	6.345140625	ab179279 - H3 K64ac
N23	7.148378125	ab179280 - H3 K64me2
N24	8.786992708	ab179281 - H3 K64me3
N25	6.389913542	ab179282 - H3 unmod(73-103)
O1	6.665946875	ab179283 - H3 K79ac
O2	7.019783333	ab179284 - H3 K79fo
O3	7.159869792	ab179285 - H3 T80p
O4	6.217703125	ab179286 - H3 K79me3 + T80p
O5	7.898558333	ab179287 - H3 unmod(99-135)
O6	6.39363125	ab179288 - H3 T118O-ac
O7	5.803059375	ab179289 - H3 K122fo
O8	6.216142708	ab179290 - H3 K122me1
O9	5.585872917	ab179291 - H3 R128me1
O10	5.895002083	ab179292 - H3 R128me2s
O11	108.7017781	ab179293 - H3.3 unmod(10-40)
O12	8.029119792	ab179294 - H3.3 S28p
O13	6.194172917	ab179295 - H3.3 S28p + S31p
O14	8.769815625	ab179296 - H3.3 S31p
O15	8.144813542	ab179297 - H4 unmod(1-30)
O16	4.603533333	ab179298 - H4 S1p + R3me1

O17	7.957282292	ab179299 - H4 S1p + R3me2s
O18	4.868709375	ab179300 - H4 S1p + R3me2a
O19	7.160642708	ab179301 - H4 S1p + K5ac + K8ac + K12ac + K16ac
O20	5.265709375	ab179302 - H4 R3cit
O21	6.109761458	ab179303 - H4 R3me1 + K5ac + K8ac + K12ac + K16ac + K20ac
O22	9.00426875	ab179304 - H4 R3me2s
O23	7.982828125	ab179305 - H4 R3me2s + K5ac + K8ac + K12ac + K16ac + K20ac
O24	4.787176042	ab179306 - H4 R3me2a
O25	5.419147917	ab179307 - H4 R3me2a + K5ac
P1	8.701459375	ab179308 - H4 R3me2a + K5ac + K8ac + K12ac + K16ac + K20ac
P2	5.606153125	ab179309 - H4 K5ac
P3	6.749770833	ab179310 - H4 K5ac + K8ac
P4	7.534867708	ab179311 - H4 K5ac + K12ac
P5	6.278988542	ab179312 - H4 K5ac + K16ac
P6	8.184617708	ab179313 - H4 K5ac + K8ac + K12ac
P7	6.10823125	ab179314 - H4 K5ac + K8me1 + K12ac
P8	7.045996875	ab179315 - H4 K5ac + K8ac + K12ac + K16ac
P9	4.751088542	ab179316 - H4 K5ac + K8me1 + K12ac + K16ac
P10	7.505876042	ab179317 - H4 K5ac + K8ac + K12me1 + K16ac
P11	6.954095833	ab179318 - H4 K5me1
P12	5.151313542	ab179319 - H4 K5me1 + K8ac + K12me1
P13	5.680161458	ab179320 - H4 K5me1 + K8me1 + K12me1
P14	5.402307292	ab179321 - H4 K5me1 + K8ac + K12ac + K16ac
P15	5.288078125	ab179322 - H4 K5me2
P16	6.233013542	ab179323 - H4 K5me3
P17	4.842848958	ab179324 - H4 K5cr
P18	7.610090625	ab179325 - H4 K8ac + K12ac
P19	8.394057292	ab179326 - H4 K8ac + K16ac
P20	6.052345833	ab179327 - H4 K8cr
P21	8.001653125	ab179328 - H4 K8me1
P22	7.427107292	ab179329 - H4 K12ac
P23	8.233615625	ab179330 - H4 K12ac + K16ac
P24	5.395188542	ab179331 - H4 K12ac + K16ac + K20me2
P25	4.671870833	ab179332 - H4 K12ac + K16ac + K20me3
Q1	6.015446875	ab179333 - H4 K5mutQ + K8mutQ + K12ac + K16mutQ + K20mutQ
Q2	5.104094792	ab179334 - H4 K12cr
Q3	7.018621875	ab179335 - H4 K12fo
Q4	5.088536458	ab179336 - H4 K12me1
Q5	7.4982375	ab179337 - H4 K12me2
Q6	5.363094792	ab179338 - H4 K12me3
Q7	6.318180208	ab179339 - H4 K16ac
Q8	6.450344792	ab179340 - H4 K16ac + K20ac
Q9	6.982760417	ab179341 - H4 K16cr
Q10	7.705920833	ab179342 - H4 K16me1
Q11	7.744875	ab179343 - H4 K16me3
Q12	6.561736458	ab179344 - H4 R17me1
Q13	8.634763542	ab179345 - H4 R17me2s
Q14	5.490009375	ab179346 - H4 R17me2a
Q15	8.70679375	ab179347 - H4 R19me1
Q16	5.70000625	ab179348 - H4 R19me2s
Q17	7.9248	ab179349 - H4 R19me2a
Q18	6.468236458	ab179350 - H4 K20ac
Q19	6.782782292	ab179351 - H4 R23me1
Q20	8.3164375	ab179352 - H4 R23me2s
Q21	6.076128125	ab179353 - H4 R23me2a
Q22	7.456182292	ab179354 - H4 unmod(24-54)
Q23	6.727376042	ab179355 - H4 K31fo
Q24	7.235405208	ab179356 - H4 K31me1
Q25	8.067222917	ab179357 - H4 K31me2
R1	7.082314583	ab179358 - H4 K31me3
R2	7.091388542	ab179359 - H4 P32hyd
R3	6.94885	ab179360 - H4 R35me1
R4	6.18805625	ab179361 - H4 R35me2s
R5	5.479282292	ab179362 - H4 R35me2a
R6	6.186011458	ab179363 - H4 S47O-ac
R7	5.767178125	ab179364 - H4 S47GlcNAc
R8	5.729041667	ab179365 - H4 S47p
R9	6.113544792	ab179366 - H4 unmod(45-75)
R10	9.085514583	ab179367 - H4 Y51hyd
R11	5.503080208	ab179368 - H4 R55me1
R12	7.301170833	ab179369 - H4 R55me2s
R13	6.188180208	ab179370 - H4 R55me2a
R14	7.650630208	ab179371 - H4 K59fo
R15	5.888697917	ab179372 - H4 K59me1

R16	6.384001042	ab179373 - H4 K59me2
R17	6.812114583	ab179374 - H4 R67me1
R18	8.6180125	ab179375 - H4 R67me2s
R19	6.721911458	ab179376 - H4 R67me2a
R20	6.712255208	ab179377 - H4 unmod(71-102)
R21	5.460786458	ab179378 - H4 K77me1
R22	5.991408333	ab179379 - H4 K77me2
R23	7.777286458	ab179380 - H4 K77me3
R24	7.206084375	ab179381 - H4 K79fo
R25	7.075777083	ab179382 - H4 K79me1
S1	6.092451042	ab179383 - H4 K79me3
S2	7.759014583	ab179384 - H4 T82O-ac
S3	6.685867708	ab179385 - H4 Y88hyd
S4	5.663094792	ab179386 - H4 Y88nitro
S5	7.497983333	ab179387 - H4 K91fo
S6	4.801444792	ab179403 - H2A K5ac
S7	5.067292708	ab179404 - H2A K9me1
S8	4.528302083	ab18504 - H2B S32p
S9	5.423803125	ab19828 - ScH2A S129p
S10	5.005919792	ab20631 - H3 R26cit
S11	6.487227083	ab20635 - H3 T3p + K4me3
S12	6.2996	ab21043 - H3 K37me1
S13	4.484457292	ab21998 - H1.4 K25me2
S14	6.030564583	ab22079 - H4 K59me3
S15	6.934458333	ab22080 - H3 K122me2
S16	6.125616667	ab22082 - H3 K122me3
S17	7.860295833	ab22214 - H3 K37me2
S18	5.869875	ab22398 - H2A R3me2a
S19	6.37085	ab22399 - H2A R3me2s
S20	7.687692708	ab23539 - H3 T22p
S21	6.05678125	ab24003 - H3 K18ac
S22	6.736698958	ab24404 - H3 K27ac
S23	6.600791667	ab24417 - H3 K37me3
S24	5.923175	ab24444 - H3 T11p
S25	5.143015625	ab24659 - H2A K127me3
T1	9.075263542	ab24660 - H2A K125me3
T2	5.709833333	ab26307 - H3 T45p
T3	4.295972917	ab26342 - H2B K5me2
T4	6.820382292	ab26739 - H4 T30p
T5	6.343670833	ab27896 - H4 T96p
T6	7.305104167	ab27898 - H4 T80p
T7	5.503704167	ab28756 - H2A R3me1
T8	5.2098125	ab32876 - H3 R2cit + R8cit + R17cit
T9	5.022327083	ab33047 - H2A K127me2
T10	5.739946875	ab33505 - H3 T118p
T11	5.978157292	ab33594 - H3 R128me2a
T12	5.8335625	ab34466 - H3 K122ac
T13	6.493363542	ab36927 - H2A S122p
T14	6.941144792	ab4555 - H3 K79me1
T15	6.77888125	ab4556 - H3 K79me2
T16	4.614766667	ab4557 - H3 K79me3
T17	16.28026875	ab4560 - H4 K79me2
T18	6.059827083	ab46854 - H3 K18me3
T19	9.634907292	ab48359 - H3 K23ac
T20	6.442441667	ab53244 - H3 K64me1
T21	5.886821875	ab54016 - H2A K9ac
T22	6.115694792	ab5499 - H3 S28p
T23	5.414192708	ab7768 - H3 K4me2
T24	6.646723958	ab89018 - H1.4 S26p
T25	5.208876042	ab92374 - H3 K4me3 (HEXI)
U1	5.68890625	ab95444 - H4 K16me2