

Poly-acetylated chromatin signatures are preferred epitopes for site-specific histone H4 acetyl antibodies

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SUPPLEMENTARY INFORMATION

Supplemental Figures 1-2 – Expanded array results

Supplemental Figure 3 – Expanded mass spectrometry results

Supplemental Table 1 – Peptide List

Supplemental Table 2 – Antibodies used in this study

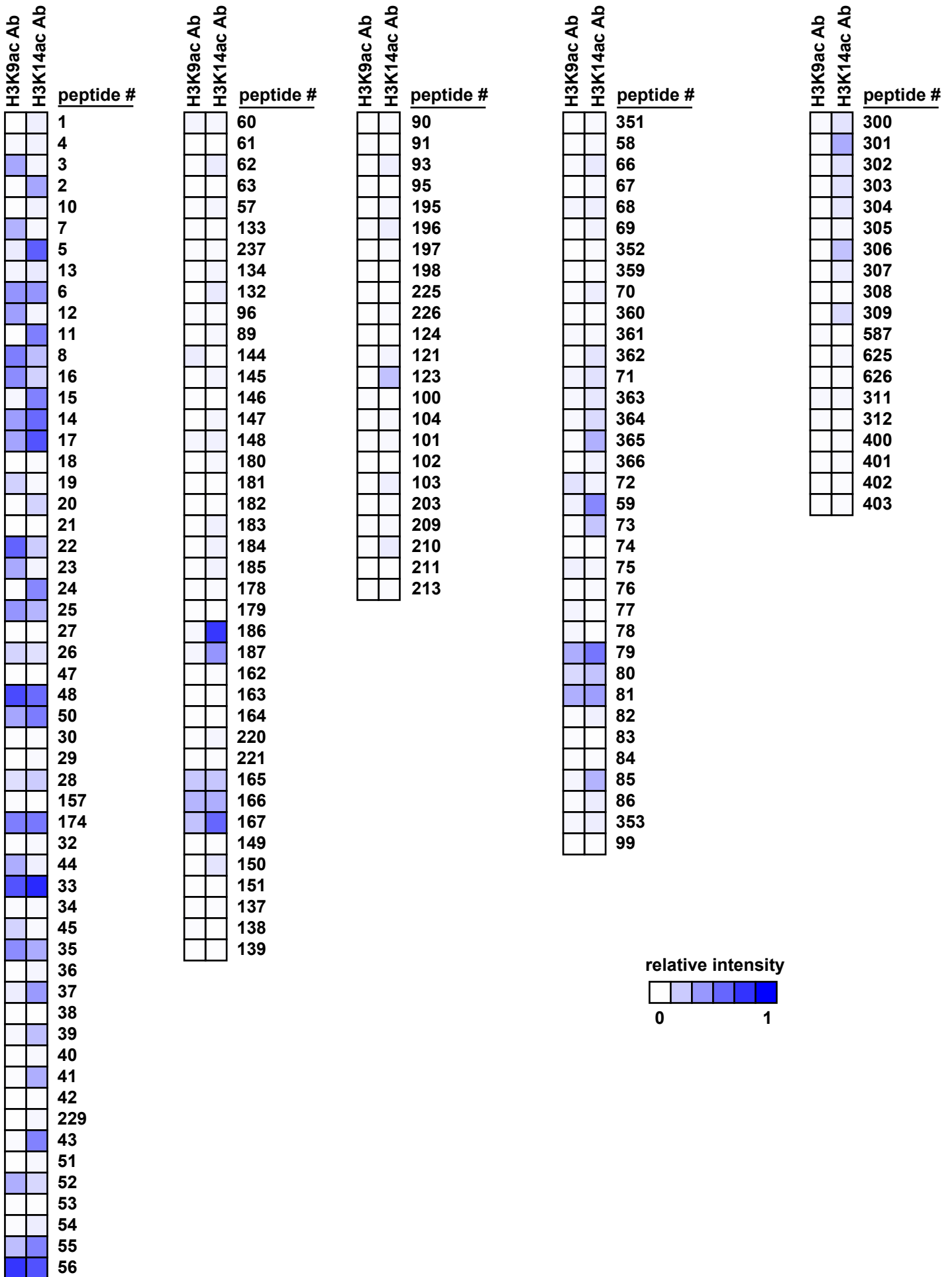
Supplemental Figure Legends

Supplemental Figure 1. Rothbart *et al*

H3 peptides

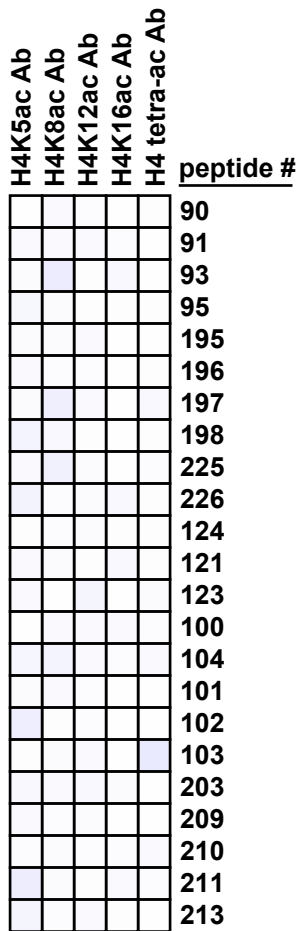
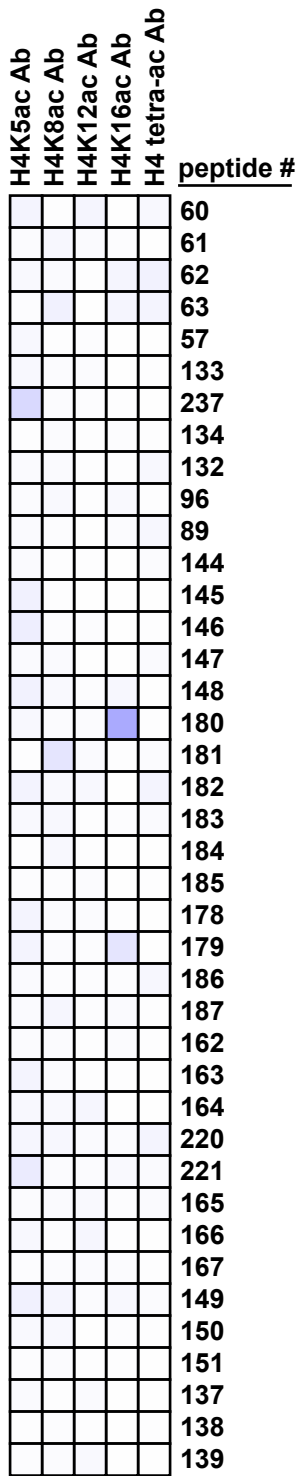
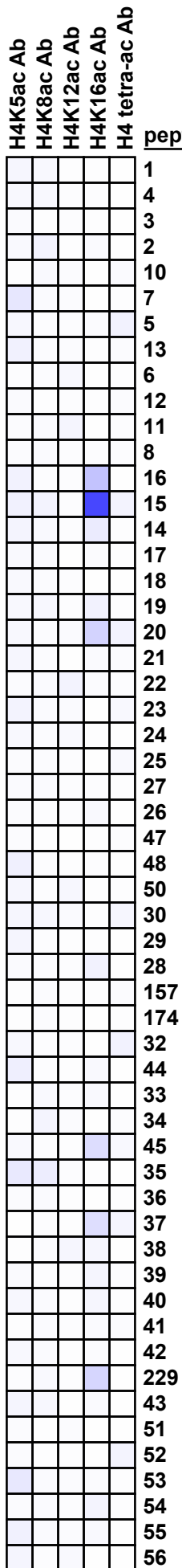
H4 peptides

H2A/H2B peptides

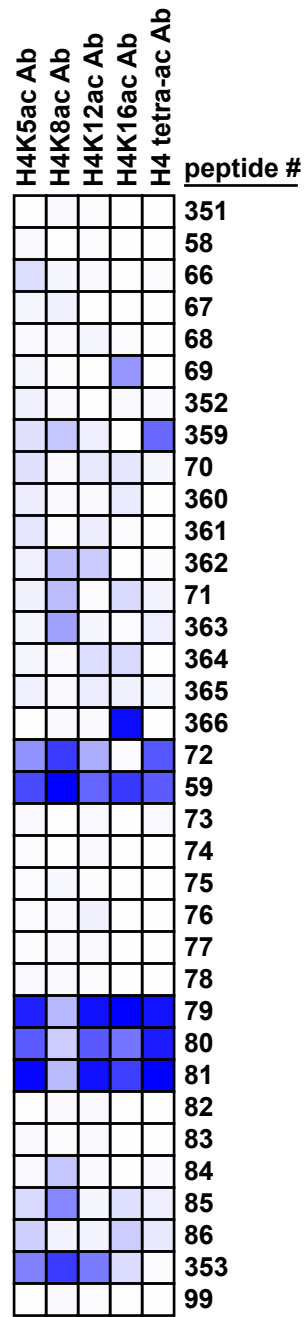


Supplemental Figure 2. Rothbart et al

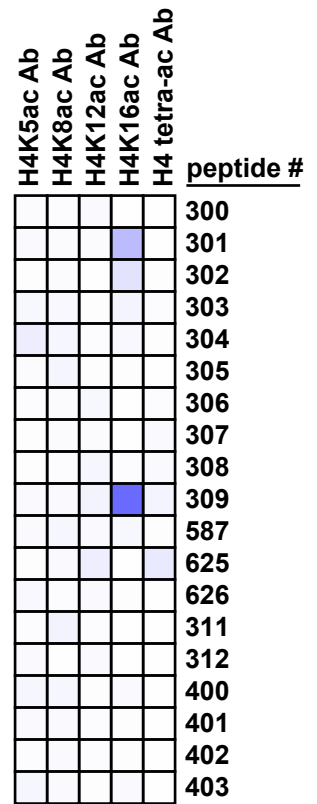
H3 peptides



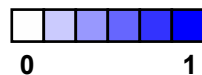
H4 peptides



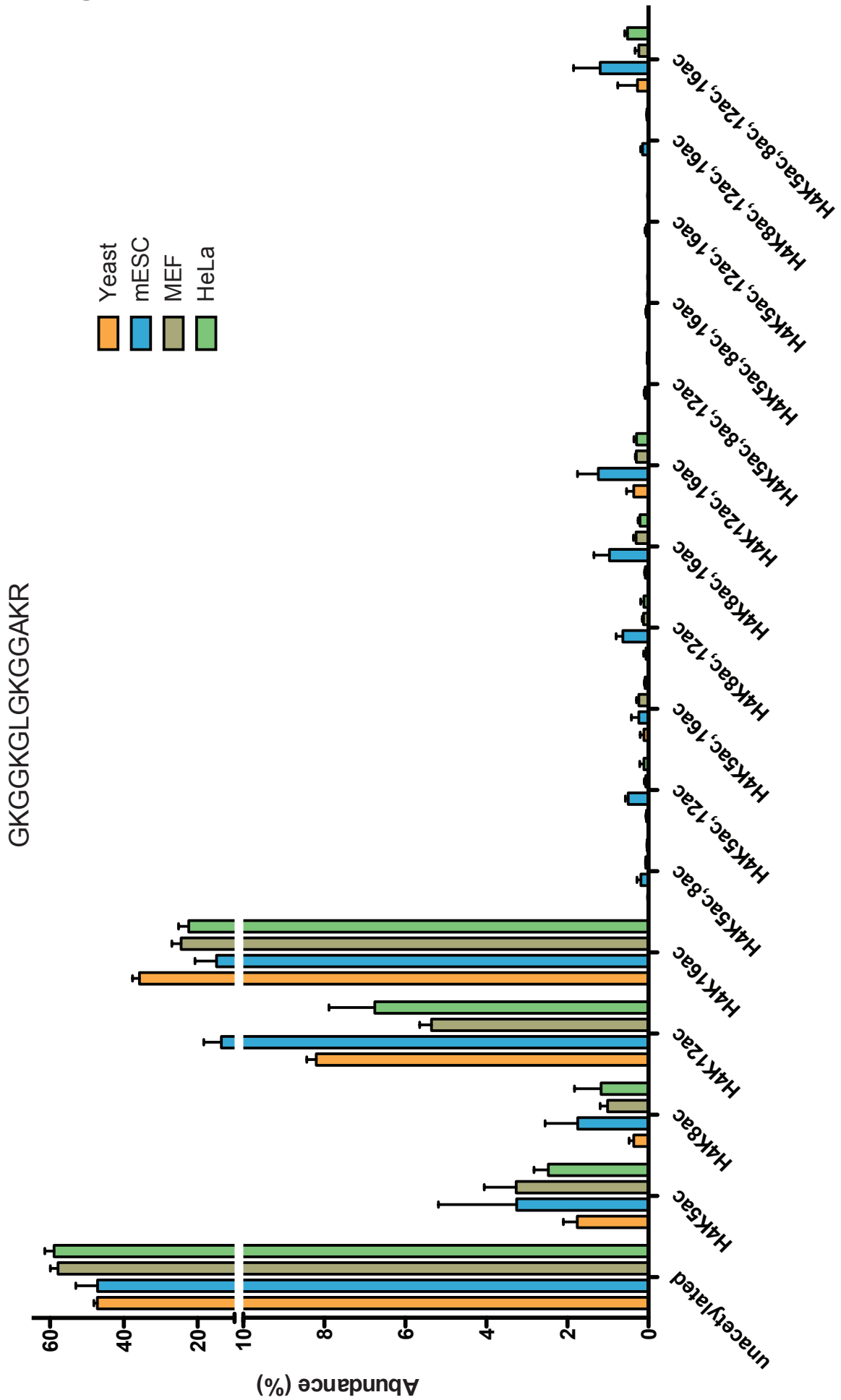
H2A/H2B peptides



relative intensity



Supplemental Figure 3. Rothbart *et al*



Supplemental Table 1. List of Peptides

| Peptide # | Sequence | annotation |
|-----------|---|--|
| 1 | H3 1-20 AR ² TK ⁴ QTAR ⁸ K ⁹ S ¹⁰ TGGK ¹⁴ APRK ¹⁸ QL-K(Biot)-NH ₂ | H3 (1-20) |
| 2 | H3 1-20 ARTKQTARKSTGGK(Ac)APRKQL-K(Biot)-NH ₂ | H3K14ac |
| 3 | H3 1-20 ARTKQTARK(Ac)STGGKAPRKQL-K(Biot)-NH ₂ | H3K9ac |
| 4 | H3 1-20 ARTK(Ac)QTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3K4ac |
| 5 | H3 1-20 ARTK(Ac)QTARKSTGGK(Ac)APRKQL-K(Biot)-NH ₂ | H3K4ac + K14ac |
| 6 | H3 1-20 ARTKQTARK(Ac)STGGK(Ac)APRKQL-K(Biot)-NH ₂ | H3K9ac + K14ac |
| 7 | H3 1-20 ARTK(Ac)QTARK(Ac)STGGKAPRKQL-K(Biot)-NH ₂ | H3K4ac + K9ac |
| 8 | H3 1-20 ARTK(Ac)QTARK(Ac)STGGK(Ac)APRKQL-K(Biot)-NH ₂ | H3K4ac + K9ac + K14ac |
| 10 | H3 1-20 ARTKQTARKSTGGKAPRK(Ac)QL-K(Biot)-NH ₂ | H3K18ac |
| 11 | H3 1-20 ARTKQTARKSTGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K14ac + K18ac |
| 12 | H3 1-20 ARTKQTARK(Ac)STGGKAPRK(Ac)QL-K(Biot)-NH ₂ | H3K9ac + K18ac |
| 13 | H3 1-20 ARTK(Ac)QTARKSTGGKAPRK(Ac)QL-K(Biot)-NH ₂ | H3K4ac + K18ac |
| 14 | H3 1-20 ARTKQTARK(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K9ac + K14ac + K18ac |
| 15 | H3 1-20 ARTK(Ac)QTARKSTGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K4ac + K14ac + K18ac |
| 16 | H3 1-20 ARTK(Ac)QTARK(Ac)STGGKAPRK(Ac)QL-K(Biot)-NH ₂ | H3K4ac + K9ac + K18ac |
| 17 | H3 1-20 ARTK(Ac)QTARK(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K4ac + K9ac + K14ac + K18ac |
| 18 | H3 1-20 ARTK(Me ₃)QTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3K4me3 |
| 19 | H3 1-20 ARTK(Me ₃)QTARK(Ac)STGGKAPRKQL-K(Biot)-NH ₂ | H3K4me3 + K9ac |
| 20 | H3 1-20 ARTK(Me ₃)QTARKSTGGK(Ac)APRKQL-K(Biot)-NH ₂ | H3K4me3 + K14ac |
| 21 | H3 1-20 ARTK(Me ₃)QTARKSTGGKAPRK(Ac)QL-K(Biot)-NH ₂ | H3K4me3 + K18ac |
| 22 | H3 1-20 ARTK(Me ₃)QTARK(Ac)STGGK(Ac)APRKQL-K(Biot)-NH ₂ | H3K4me3 + K9ac + K14ac |
| 23 | H3 1-20 ARTK(Me ₃)QTARK(Ac)STGGKAPRK(Ac)QL-K(Biot)-NH ₂ | H3K4me3 + K9ac + K18ac |
| 24 | H3 1-20 ARTK(Me ₃)QTARKSTGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K4me3 + K14ac + K18ac |
| 25 | H3 1-20 ARTK(Me ₃)QTARK(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K4me3 + K9ac + K14ac + K18ac |
| 26 | H3 1-20 ARpTK(Me ₃)QTARK(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3T3p + K4me3 + K9ac + K14ac + K18ac |
| 27 | H3 1-20 ARpTK(Me ₃)QTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3T3p + K4me3 |
| 28 | H3 1-20 AR(Me ₂ a)pTK(Me ₃)QTARK(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3R2me2a + T3p + K4me3 + K9ac + K14ac + K18ac |
| 29 | H3 1-20 AR(Me ₂ a)pTK(Me ₃)QTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3R2me2a + T3p + K4me3 |
| 30 | H3 1-20 AR(Me ₂ a)TK(Me ₃)QTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3R2me2a + K4me3 |
| 32 | H3 1-20 ARTK(Me ₂)QTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3K4me2 |
| 33 | H3 1-20 ARTK(Me ₂)QTARK(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K4me2 + K9ac + K14ac + K18ac |
| 34 | H3 1-20 ARTK(Me)QTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3K4me1 |
| 35 | H3 1-20 ARTK(Me)QTARK(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K4me1 + K9ac + K14ac + K18ac |
| 36 | H3 1-20 ARTKQTARKpSTGGKAPRKQL-K(Biot)-NH ₂ | H3S10p |
| 37 | H3 1-20 ARTK(Ac)QTARK(Ac)pSTGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K4ac + K9ac + S10p + K14ac + K18ac |
| 38 | H3 1-20 ARTK(Me ₃)QTARKpSTGGKAPRKQL-K(Biot)-NH ₂ | H3K4me3 + S10p |
| 39 | H3 1-20 ARTK(Me ₃)QTARK(Ac)pSTGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K4me3 + K9ac + S10p + K14ac + K18ac |
| 40 | H3 1-20 AR(Me ₂ a)TK(Me ₃)QTARKpSTGGKAPRKQL-K(Biot)-NH ₂ | H3R2me2a + K4me3 + S10p |
| 41 | H3 1-20 AR(Me ₂ a)TK(Me ₃)QTARK(Ac)pSTGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3R2me2a + K4me3 + K9ac + S10p + K14ac + K18ac |
| 42 | H3 1-20 ARTKQTARK(Me ₃)STGGKAPRKQL-K(Biot)-NH ₂ | H3K9me3 |
| 43 | H3 1-20 ARTK(Ac)QTARK(Me ₃)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K4ac + K9me3 + K14ac + K18ac |
| 44 | H3 1-20 ARTK(Me ₂)QTARK(Ac)STGGKAPRK(Ac)QL-K(Biot)-NH ₂ | H3K4me2 + K9ac + K18ac |

| | | | |
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| 45 | H3 1-20 | ARTK(Me)QTARK(Ac)STGGKAPRK(Ac)QL-K(Biot)-NH ₂ | H3K4me1 + K9ac + K18ac |
| 47 | H3 1-20 | AR(Me ₂ a)TKQTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3R2me2a |
| 48 | H3 1-20 | AR(Me ₂ a)TK(Ac)QTARK(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3R2me2a + K4ac + K9ac + K14ac + K18ac |
| 50 | H3 1-20 | AR(Me ₂ a)TK(Me ₃)QTARK(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3R2me2a + K4me3 + K9ac + K14ac + K18ac |
| 51 | H3 1-20 | AR(Me)TK(Me ₃)QTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3R2me1 + K4me3 |
| 52 | H3 1-20 | AR(Me)TK(Me ₃)QTARK(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3R2me1 + K4me3 + K9ac + K14ac + K18ac |
| 53 | H3 1-20 | ACitTKQTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3Cit2 |
| 54 | H3 1-20 | ACitTK(Me ₃)QTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3Cit2 + K4me3 |
| 55 | H3 1-20 | ACitTK(Me ₃)QTARK(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3Cit2 + K4me3 + K9ac + K14ac + K18ac |
| 56 | H3 1-20 | ACitTK(Ac)QTARK(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3Cit2 + K4ac + K9ac + K14ac + K18ac |
| 57 | H3 1-20 | ARpTKQTARKSTGGKAPRKQL-Peg-K(Biot)-NH ₂ | H3T3p |
| 58 | H4 1-23 | Ac-SGRGK ⁵ GGKGLGKGGAKRHRKVLR-Peg-Biot | H4 (1-23) |
| 59 | H4 1-23 | Ac-SGRGK(Ac)GGK(Ac)GLGK(Ac)GGAK(Ac)RHRKVLR-Peg-Biot | H4K5ac + K8ac + K12ac + K16ac |
| 60 | H3 1-20 | AR(Me ₂ a)TK(Me ₂)QTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3R2me2a + K4me2 |
| 61 | H3 1-20 | AR(Me ₂ s)TK(Me ₂)QTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3R2me2s + K4me2 |
| 62 | H3 1-20 | AR(Me)TK(Me ₂)QTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3R2me1 + K4me2 |
| 63 | H3 1-20 | ACitTK(Me ₂)QTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3Cit2 + K4me2 |
| 66 | H4 1-23 | Ac-SGRGK(Ac)GGKGLGKGGAKRHRKVLR-Peg-Biot | H4K5ac |
| 67 | H4 1-23 | Ac-SGRGKGGK(Ac)GLGKGGAKRHRKVLR-Peg-Biot | H4K8ac |
| 68 | H4 1-23 | Ac-SGRGKGGKGLGK(Ac)GGAKRHRKVLR-Peg-Biot | H4K12ac |
| 69 | H4 1-23 | Ac-SGRGKGGKGLGKGGAK(Ac)RHRKVLR-Peg-Biot | H4K16ac |
| 70 | H4 1-23 | Ac-SGRGK(Ac)GGKGLGK(Ac)GGAKRHRKVLR-Peg-Biot | H4K5ac + K12ac |
| 71 | H4 1-23 | Ac-SGRGKGGK(Ac)GLGKGGAK(Ac)RHRKVLR-Peg-Biot | H4K8ac + K16ac |
| 72 | H4 1-23 | Ac-SGRGK(Ac)GGK(Ac)GLGK(Ac)GGAKRHRKVLR-Peg-Biot | H4K5ac + K8ac + K12ac |
| 73 | H4 1-23 | Ac-SGR(Me ₂ a)GKGGKGLGKGGAKRHRKVLR-K(Biot)-NH ₂ | H4R3me2a |
| 74 | H4 1-23 | Ac-SGR(Me ₂ s)GKGGKGLGKGGAKRHRKVLR-K(Biot)-NH ₂ | H4R3me2s |
| 75 | H4 1-23 | Ac-SGR(Me)GKGGKGLGKGGAKRHRKVLR-K(Biot)-NH ₂ | H4R3me1 |
| 76 | H4 1-23 | Ac-pSGR(Me ₂ a)GKGGKGLGKGGAKRHRKVLR-K(Biot)-NH ₂ | H4S1p + R3me2a |
| 77 | H4 1-23 | Ac-pSGR(Me ₂ s)GKGGKGLGKGGAKRHRKVLR-K(Biot)-NH ₂ | H4S1p + R3me2s |
| 78 | H4 1-23 | Ac-pSGR(Me)GKGGKGLGKGGAKRHRKVLR-K(Biot)-NH ₂ | H4S1p + R3me1 |
| 79 | H4 1-23 | Ac-SGR(Me ₂ a)GK(Ac)GGK(Ac)GLGK(Ac)GGAK(Ac)RHRK(Ac)VLR-K(Biot)-NH ₂ | H4R3me2a + K5ac + K8ac + K12ac + K16ac + K20ac |
| 80 | H4 1-23 | Ac-SGR(Me ₂ s)GK(Ac)GGK(Ac)GLGK(Ac)GGAK(Ac)RHRK(Ac)VLR-K(Biot)-NH ₂ | H4R3me2s + K5ac + K8ac + K12ac + K16ac + K20ac |
| 81 | H4 1-23 | Ac-SGR(Me)GK(Ac)GGK(Ac)GLGK(Ac)GGAK(Ac)RHRK(Ac)VLR-K(Biot)-NH ₂ | H4R3me1 + K5ac + K8ac + K12ac + K16ac + K20ac |
| 82 | H4 11-27 | Ac-GKGGAKRHRK(Me ₃)VLRDNIQ-Peg-Biot | H4K20me3 |
| 83 | H4 11-27 | Ac-GKGGAKRHRK(Me ₂)VLRDNIQ-Peg-Biot | H4K20me2 |
| 84 | H4 11-27 | Ac-GKGGAKRHRK(Me)VLRDNIQ-Peg-Biot | H4K20me1 |
| 85 | H4 11-27 | Ac-GK(Ac)GGAK(Ac)RHRK(Me ₃)VLRDNIQ-Peg-Biot | H4K12ac + K16ac + K20me3 |
| 86 | H4 11-27 | Ac-GK(Ac)GGAK(Ac)RHRK(Me ₂)VLRDNIQ-Peg-Biot | H4K12ac + K16ac + K20me2 |
| 89 | H3 1-20 | ARTK(Me ₃)QTAR(Me ₂ s)K(Me ₃)STGGKAPRKQL-K(Biot)-NH ₂ | H3K4me3 + R8me2s + K9me3 |
| 90 | H3 15-43 | Ac-APRK ¹⁸ QLATK ²³ AARK ²⁷ SAPSTGGVK ³⁶ K ³⁷ PHRYGGK(Biot)-NH ₂ | H3 (15-41) |
| 91 | H3 15-43 | Ac-APRK(Me ₃)QLATKAARKSAPSTGGVKKPHRY-GG-K(Biot)-NH ₂ | H3K18me3 |
| 93 | H3 15-43 | Ac-APRKQLATKAARKSAPSTGGVKK(Me ₃)KPHRY-GG-K(Biot)-NH ₂ | H3K36me3 |
| 95 | H3 15-43 | Ac-APRK(Me ₃)QLATKAARKSAPSTGGVKK(Me ₃)KPHRY-GG-K(Biot)-NH ₂ | H3K18me3 + K36me3 |
| 96 | H3 1-20 | ARTK(Me ₃)QTAR(Me ₂ a)K(Me ₃)STGGKAPRKQL-K(Biot)-NH ₂ | H3K4me3 + R8me2a + K9me3 |
| 99 | H4 11-27 | Ac-GKGGAKRHRKVLRDNIQ-Peg-Biot | H4 (11-27) |
| 100 | H3 74-84 | Ac-IAQDFKTDLRF-Peg-K(Biot)-NH ₂ | H3 (74-84) N-ac |
| 101 | H3 74-84 | Ac-IAQDFK(Me ₃)TDLRF-Peg-K(Biot)-NH ₂ | H3K79me3 |
| 102 | H3 74-84 | Ac-IAQDFK(Me ₂)TDLRF-Peg-K(Biot)-NH ₂ | H3K79me2 |

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| 103 | H3 74-84 | Ac-IAQDFK(Me)TDLRF-Peg-K(Biot)-NH ₂ | H3K79me1 |
| 104 | H3 74-84 | IAQDFKTDLRF-Peg-K(Biot)-NH ₂ | H3 (74-84) |
| 121 | H3 27-45 | KSAPSTGGVK(Me ₂)KPHRYKPGT-GG-K(Biot)-NH ₂ | H3K36me2 |
| 123 | H3 27-45 | KSAPSTGGVK(Ac)KPHRYKPGT-GG-K(Biot)-NH ₂ | H3K36ac |
| 124 | H3 27-45 | KSAPSTGGVKKPHRYKPGT-GG-K(Biot)-NH ₂ | H3 (27-45) |
| 132 | H3 1-20 | ARTK(Me ₃)QTARK(Me ₃)STGGKAPRKQL-K(Biot)-NH ₂ | H3K4me3 + K9me3 |
| 133 | H3 1-20 | ARTKQTARK(Me ₂)STGGKAPRKQL-K(Biot)-NH ₂ | H3K9me2 |
| 134 | H3 1-20 | ARTKQTARK(Me)STGGKAPRKQL-K(Biot)-NH ₂ | H3K9me1 |
| 137 | H3 1-20 | ARTKQTARKSTGGKAPRK(Me ₃)QL-K(Biot)-NH ₂ | H3K18me3 |
| 138 | H3 1-20 | ARTKQTARKSTGGKAPRK(Me ₂)QL-K(Biot)-NH ₂ | H3K18me2 |
| 139 | H3 1-20 | ARTKQTARKSTGGKAPRK(Me)QL-K(Biot)-NH ₂ | H3K18me1 |
| 144 | H3 1-20 | ARTKQTARK(Ac)pSTGGKAPRKQL-K(Biot)-NH ₂ | H3K9ac + S10p |
| 145 | H3 1-20 | ARTKQTARK(Me ₃)pSTGGKAPRKQL-K(Biot)-NH ₂ | H3K9me3 + S10p |
| 146 | H3 1-20 | ARTKQTARK(Me ₂)pSTGGKAPRKQL-K(Biot)-NH ₂ | H3K9me2 + S10p |
| 147 | H3 1-20 | ARTKQTARK(Me)pSTGGKAPRKQL-K(Biot)-NH ₂ | H3K9me1 + S10p |
| 148 | H3 1-20 | ARTK(Me ₃)QTARK(Ac)pSTGGKAPRKQL-K(Biot)-NH ₂ | H3K4me3 + K9ac + S10p |
| 149 | H3 1-22 | ARTKQTARKSTGGKAPR(Me ₂ a)KQLAT-K(Biot)-NH ₂ | H3R17me2a |
| 150 | H3 1-22 | ARTKQTARKSTGGKAPR(Me ₂ s)KQLAT-K(Biot)-NH ₂ | H3R17me2s |
| 151 | H3 1-22 | ARTKQTARKSTGGKAPR(Me)KQLAT-K(Biot)-NH ₂ | H3R17me1 |
| 157 | H3 1-20 | AR(Me ₂ s)TK(Me ₃)QTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3R2me2s + K4me3 |
| 162 | H3 1-20 | ARTKQpTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3T6p |
| 163 | H3 1-20 | ARTK(Me ₃)QpTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3K4me3 + T6p |
| 164 | H3 1-20 | ARTK(Me ₂)QpTARKSTGGKAPRKQL-K(Biot)-NH ₂ | H3K4me2 + T6p |
| 165 | H3 1-20 | ARTKQpTARK(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3T6p + K9ac + K14ac + K18ac |
| 166 | H3 1-20 | ARTK(Me ₃)QpTAR(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K4me3 + T6p + K9ac + K14ac + K18ac |
| 167 | H3 1-20 | ARTK(Me ₂)QpTAR(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K4me2 + T6p + K9ac + K14ac + K18ac |
| 174 | H3 1-20 | AR(Me ₂ s)TK(Me ₃)QTARK(Ac)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3R2me2s + K4me3 + K9ac + K14ac + K18ac |
| 178 | H3 1-20 | ARTKQTAR(Me)K(Me ₃)STGGKAPRKQL-K(Biot)-NH ₂ | H3R8me1 + K9me3 |
| 179 | H3 1-20 | ARTKQTAR(Me)K(Me ₂)STGGKAPRKQL-K(Biot)-NH ₂ | H3R8me1 + K9me2 |
| 180 | H3 1-20 | ARTKQTAR(Me ₂ a)K(Me ₃)STGGKAPRKQL-K(Biot)-NH ₂ | H3R8me2a + K9me3 |
| 181 | H3 1-20 | ARTKQTAR(Me ₂ a)K(Me ₂)STGGKAPRKQL-K(Biot)-NH ₂ | H3R8me2a + K9me2 |
| 182 | H3 1-20 | ARTKQTAR(Me ₂ a)K(Me)STGGKAPRKQL-K(Biot)-NH ₂ | H3R8me2a + K9me1 |
| 183 | H3 1-20 | ARTKQTAR(Me ₂ s)K(Me ₃)STGGKAPRKQL-K(Biot)-NH ₂ | H3R8me2s + K9me3 |
| 184 | H3 1-20 | ARTKQTAR(Me ₂ s)K(Me ₂)STGGKAPRKQL-K(Biot)-NH ₂ | H3R8me2s + K9me2 |
| 185 | H3 1-20 | ARTKQTAR(Me ₂ s)K(Me)STGGKAPRKQL-K(Biot)-NH ₂ | H3R8me2s + K9me1 |
| 186 | H3 1-20 | ARTK(Ac)QTARK(Me ₂)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K4ac + K9me2 + K14ac + K18ac |
| 187 | H3 1-20 | ARTK(Ac)QTARK(Me)STGGK(Ac)APRK(Ac)QL-K(Biot)-NH ₂ | H3K4ac + K9me1 + K14ac + K18ac |
| 195 | H3 15-34 | Ac-APRK ¹⁸ QLATK ²³ AARK(Me ₃) ²⁷ SAPSTGG-Peg-Biot | H3K27me3 |
| 196 | H3 15-34 | Ac-APRK ¹⁸ QLATK ²³ AARK(Me ₂) ²⁷ SAPSTGG-Peg-Biot | H3K27me2 |
| 197 | H3 15-34 | Ac-APRK ¹⁸ QLATK ²³ AARK(Me) ²⁷ SAPSTGG--Peg-Biot | H3K27me1 |
| 198 | H3 15-34 | Ac-APRK ¹⁸ QLATK ²³ AAR(Me ₂ a)K(Me ₃) ²⁷ SAPSTGG--Peg-Biot | H3R26me2a + K27me3 |
| 203 | H3 30-49 | Ac-PATGGVKKPHRYRPGTVALR-Peg-K(Biot)-NH ₂ | H3 (30-49) |
| 209 | H3.3 15-34 | Ac-APRKQLATKAARKSAPSTGG-Peg-K(Biot)-NH ₂ | H3.3 (15-34) |
| 210 | H3 30-49 | Ac-PSTGGVKKPHRYRPGTVALR-Peg-K(Biot)-NH ₂ | H3.3 (30-49) |
| 211 | H3.3 75-94 | Ac-AQDFKTDLRFQSAAGALQE-Peg-K(Biot)-NH ₂ | H3.3 (75-94) |
| 213 | H3 120-135 | (Biot)Peg-MPKDIQLARRIRGERA-OH | H3 (120-135) |
| 220 | H3 1-20 | ARTKQpTARK(Me ₃)STGGKAPRKQL-K(Biot)-NH ₂ | H3T6p + K9me3 |
| 221 | H3 1-20 | ARTKQpTAR(Me ₂ a)K(Me ₃)STGGKAPRKQL-K(Biot)-NH ₂ | H3T6p + R8me2a + K9me3 |

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| 225 | H3 15-34 | Ac-APRK ¹⁸ QLATK ²³ AARK(Me ₃) ²⁷ pSAPSTGG-Peg-Biot | H3K27me3 + S28p |
| 226 | H3 15-34 | Ac-APRK ¹⁸ QLATK ²³ AARK(Me ₂) ²⁷ pSAPSTGG-Peg-Biot | H3S27me2 + S28p |
| 229 | H3 1-20 | ARTK(Ac)QTARK(Me ₃)STGGKAPRKQL-K(Biot)-NH ₂ | H3K4ac + K9me3 |
| 237 | H3 1-32 | ARTKQTARK(Me ₂)STGGKAPRKQLATKAARKSAPAT-Peg-Biot | H3K9me2 (1-32) |
| 300 | H2A 1-17 | Ac-SGRGKQGGKARAKAKTR-Peg-Biot | H2A (1-17) |
| 301 | H2A 1-17 | Ac-SGRGK(Ac)QGGK(Ac)ARAK(Ac)AK(Ac)TR-Peg-Biot | H2AK5ac + K9ac + K13ac + K15ac |
| 302 | H2A 1-17 | Ac-SGRGK(Ac)QGGKARAKAKTR-Peg-Biot | H2AK5ac |
| 303 | H2A 1-17 | Ac-pSGRGK(Ac)QGGKARAKAKTR-Peg-Biot | H2AS1p + K5ac |
| 304 | H2A 1-17 | Ac-SGR(Me ₂ a)GK(Ac)QGGKARAKAKTR-Peg-Biot | H2AR3me2a + K5ac |
| 305 | H2A 1-17 | Ac-pSGR(Me ₂ a)GK(Ac)QGGKARAKAKTR-Peg-Biot | H2AS1p + R3me2a + K5ac |
| 306 | H2A 1-17 | Ac-SGCitGK(Ac)QGGKARAKAKTR-Peg-Biot | H2ACit3 + K5ac |
| 307 | H2A 1-17 | Ac-pSGCitGK(Ac)QGGKARAKAKTR-Peg-Biot | H2AS1p + Cit3 + K5ac |
| 308 | H2A 1-17 | Ac-pSGRGK(Ac)QGGK(Ac)ARAK(Ac)AK(Ac)TR-Peg-Biot | H2AS1p + K5ac + K9ac + K13ac + K15ac |
| 309 | H2A 1-17 | SGRGK(Ac)QGGK(Ac)ARAK(Ac)AK(Ac)TR-Peg-Biot | H2AK5ac + K9ac + K13ac + K15ac (no N-ac) |
| 311 | H2A.X | Biot-Peg-G ¹³² KKATQAS ¹³⁹ QEY ¹⁴² -OH | H2AX (132-142) |
| 312 | H2A.X | Biot-Peg-G ¹³² KKATQApS ¹³⁹ QEY ¹⁴² -OH | H2AX (S139p) |
| 351 | H4 1-23 | SGRGKGGKGLGKGGAKRHRKVLR-Peg-Biot | H4 (1-23) (no N-ac) |
| 352 | H4 1-23 | Ac-SGRGKGGKGLGKGGAKRHRK(Ac)VLRD-Peg-Biot | H4K20ac |
| 353 | H4 1-23 | Ac-pSGRGK(Ac)GGK(Ac)GLGK(Ac)GGAK(Ac)RHRKVLR-Peg-Biot | H4S1p + K5ac + K8ac + K12ac + K16ac |
| 359 | H4 1-23 | Ac-SGRGK(Ac)GGK(Ac)GLGKGGAKRHRKVLR-Peg-Biot | H4K5ac + K8ac |
| 360 | H4 1-23 | Ac-SGRGK(Ac)GGKGLGKGGAK(Ac)RHRKVLR-Peg-Biot | H4K5ac + K16ac |
| 361 | H4 1-23 | Ac-SGRGK(Ac)GGKGLGKGGAKRHRK(Ac)VLR-Peg-Biot | H4K5ac + K20ac |
| 362 | H4 1-23 | Ac-SGRGKGGK(Ac)GLGK(Ac)GGAKRHRKVLR-Peg-Biot | H4K8ac + K12ac |
| 363 | H4 1-23 | Ac-SGRGKGGK(Ac)GLGKGGAKRHRK(Ac)VLR-Peg-Biot | H4K8ac + K20ac |
| 364 | H4 1-23 | Ac-SGRGKGGKGLGK(Ac)GGAK(Ac)RHRKVLR-Peg-Biot | H4K12ac + K16ac |
| 365 | H4 1-23 | Ac-SGRGKGGKGLGK(Ac)GGAKRHRK(Ac)VLR-Peg-Biot | H4K12ac + K20ac |
| 366 | H4 1-23 | Ac-SGRGKGGKGLGKGGAK(Ac)RHRK(Ac)VLR-Peg-Biot | H4K16ac + K20ac |
| 400 | H2B 1-24 | PEPAK ¹⁸ SAPAPKKGSKKAVTKAQKK-Peg-Biot | H2B (1-24) |
| 401 | H2B 1-24 | PEPAK(Me ₃)SAPAPKKGSKKAVTKAQKK-Peg-Biot | H2BK5me3 |
| 402 | H2B 1-24 | PEPAK(Me ₂)SAPAPKKGSKKAVTKAQKK-Peg-Biot | H2BK5me2 |
| 403 | H2B 1-24 | PEPAK(Me)SAPAPKKGSKKAVTKAQKK-Peg-Biot | H2BK5me1 |
| 587 | H2A10-25 | Ac-ARAKAKTRSSRAGLQF-Peg-Biotin | H2A (10-25) |
| 625 | H2A.X 1-17 | Ac-SGRGKTGGKARAKAKSR-Peg-Biotin | H2A.X (1-17) |
| 626 | H2A.X 1-17 | Ac-SGRGK(Ac)TGGKARAKAKSR-Peg-Biotin | H2A.X K5ac |

Supplemental Table 2. Rothbart *et al*

Supplemental Table 2. Details of antibodies tested

| Antibody | Company | Catalog # | Host and antibody type | Concentration | Results Location | tetra-ac/ mono-ac (\pm s.e.m.) |
|-------------|--------------|-----------|------------------------|---------------|--------------------------------------|---|
| H3K9ac | Millipore | 07-352 | Rabbit polyclonal | 1:1000 | Fig. 1a Supp. Fig. 1 | 1.8 \pm 0.3 |
| H3K14ac | Millipore | 07-353 | Rabbit polyclonal | 1:1000 | Fig. 1a Supp. Fig. 1 | 1.1 \pm 0.2 |
| H4K5ac | Active Motif | 39169 | Rabbit polyclonal | 1:1000 | Fig. 1b-c Fig. 2c Supp. Fig. 2 | 4.0 \pm 0.3 |
| H4K5ac | Millipore | 07-327 | Rabbit polyclonal | 1:1000 | not shown | 15.9 \pm 4.1 |
| H4K8ac | Active Motif | 39172 | Rabbit polyclonal | 1:1000 | Fig. 1b-c Supp. Fig. 2 | 21.9 \pm 5.0 |
| H4K12ac | Active Motif | 39165 | Rabbit polyclonal | 1:1000 | Fig. 1b-d Fig. 2b Supp. Fig. 2 | 18.6 \pm 2.1 |
| H4K12ac | Millipore | 04-119 | Rabbit monoclonal | 1:1000 | not shown | 11.2 \pm 2.5 |
| H4K12ac | Millipore | 07-595 | Rabbit polyclonal | 1:1000 | not shown | 2.8 \pm 0.2 |
| H4K16ac | Active Motif | 39168 | Rabbit polyclonal | 1:2500 | Fig. 1b-c Supp. Fig. 2 | 1.7 \pm 0.3 |
| H4 tetra-ac | Active Motif | 39179 | Rabbit polyclonal | 1:2000 | Fig. 1b-c Supp. Fig. 2 | n/a |

SUPPLEMENTAL FIGURE LEGENDS

Supplemental Figures 1-2. Expanded heat map of peptide array results for H3 acetyl antibodies (Supplemental Fig. 1) and H4 acetyl antibodies (Supplemental Fig. 2). For each array, the most intense series of peptide spots (12 individual spots per peptide) is assigned a value of 1 (blue), and all values are normalized to this peptide. Each interaction is presented as an averaged normalized intensity from at least two independent arrays ($r^2 > 0.9$).

Supplemental Figure 3. Quantitative mass spectrometry of the histone H4 tail (amino acids 3-17) across species. Error is represented as standard deviations from three biological replicates.