

Correction: Novel insights into the recognition of acetylated histone H4 tail by the TRIM24 PHD-Bromo module

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Due to an error in the production process, [Table 1](#) was omitted from the published article. The complete [Table 1](#) is presented here. Portland Press apologises for the error.

Table 1. Binding affinities of the TRIM24 PHD-Bromo and Bromo alone to acetylated histone H4 peptides as measured by ITC

Histone peptide	Peptide sequence	TRIM24-PHD-Bromo ITC K_D (μM)	TRIM24-Bromo ITC K_D (μM)
H4(1-20)unmodified	SGRGKGGKGLGKGGAKRHRK	No binding	No binding
H4(1-15)K5ac	SGRGKacGGKGLGKGGGA	7.5 ± 1.0	6.5 ± 1.2
H4(1-15)K8ac	SGRGKGGKacGLGKGGGA	4.2 ± 1.3	5.5 ± 1.0
H4(1-15)K12ac	SGRGKGGKGLGKacGGA	> 500	> 500
H4(11-21)K16ac	GKGGAKacRHRKVY	6.0 ± 1.0	6.5 ± 1.0
H4(1-15)K5acK8ac	SGRGKacGGKacGLGKGGGA	4.2 ± 1.4	5.2 ± 1.0
H4(1-15)K5acK12ac	SGRGKacGGKGLGKacGGA	9.4 ± 1.0	8.6 ± 1.0
H4(1-15)K8acK12ac	SGRGKGGKacGLGKacGGA	8.6 ± 1.0	9.4 ± 1.2
H4(1-20)Kac4 (K5/K8/K12/K16)	SGRGKacGGKacGLGKacGGAKacRHRK	12.8 ± 1.0	14.5 ± 1.0

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