

Identification, Characterization, and Heritability of Murine Metastable Epialleles: Implications for Non-genetic Inheritance

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In the Introduction to this article, we inadvertently neglected to cite Faulk et al., 2013. This study carried out a phylogenetic analysis of the IAPLTR1_Mm subclass of IAP elements, which includes the *A^{vy}* and *Cabp^{IAP}* loci. The authors show that this family is divided into three clades and provide data suggesting that the younger clades may be enriched for inter-individual methylation variation. This was corrected online on November 7, 2018.

REFERENCE

Faulk, C., Barks, A., and Dolinoy, D.C. (2013). Phylogenetic and DNA methylation analysis reveal novel regions of variable methylation in the mouse IAP class of transposons. *BMC Genomics* 14, 48.

