



Figure S1. Sequence divergence of IAPEY_LTR genomic copies.

Sequence divergence (%) from the respective consensus sequences for genomic copies of IAPEY_LTR (A), IAPLTR1_Mm (B), IAPLTR2a2_Mm (c), and IAPLTR3 (D) are shown. For IAPEY_LTR sequences, statistics were analyzed separately for Y-chromosomal (E), X-chromosomal (F), and autosomal (G) copies. Although IAPEY copies have been accumulated in the Y chromosome, these copies are more divergent from the consensus sequence than those in autosomes.