Supplemental Figure S2: $P_{CT}$ scores of X-Y pairs across 8 mammals. (A) $P_{CT}$ score distributions of all gene-miRNA interactions involving X-Y pairs across eight sequenced mammalian Y Chromosomes (n = 647 interactions from 32 genes) and other ancestral X genes (n = 8,831 interactions from 457 genes). ** p < 0.01, two-sided Kolmogorov-Smirnov test. (B) Gene-level mean $P_{CT}$ scores. * p < 0.05, two-sided Wilcoxon rank-sum test.