



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.