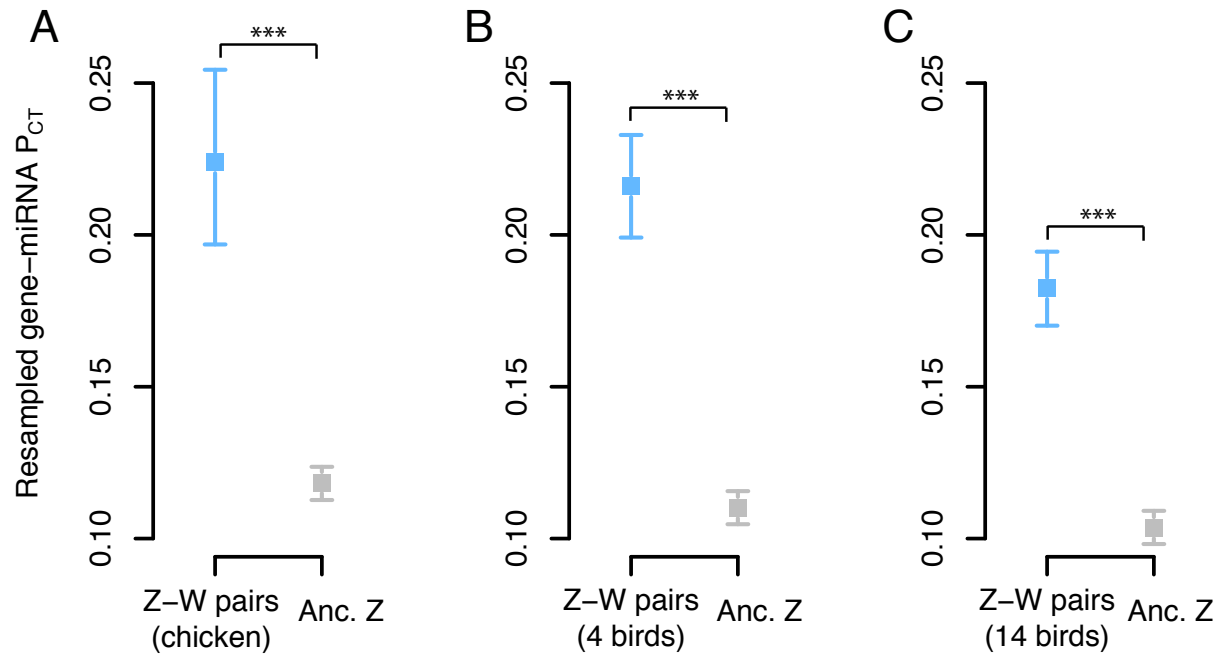




Supplemental Figure S10: Resampled mean P_{CT} scores of Z-linked genes. Gene sets: (A) chicken Z-W pairs ($n = 28$ genes) and other ancestral Z genes ($n = 657$ genes), (B) Z-W pairs across four birds ($n = 78$ genes) compared to the remainder of ancestral Z genes ($n = 607$ genes), and (C) Z-W pairs across 14 birds ($n = 157$ genes) compared to the remainder of ancestral Z genes ($n = 528$ genes). Points and error bars represent the median and 95% confidence intervals from 1,000 gene samplings with replacement. *** $p < 0.001$, empirical p-value computed as the fraction of random non-overlapping gene sets with a median difference in P_{CT} score at least as large as the true difference.