

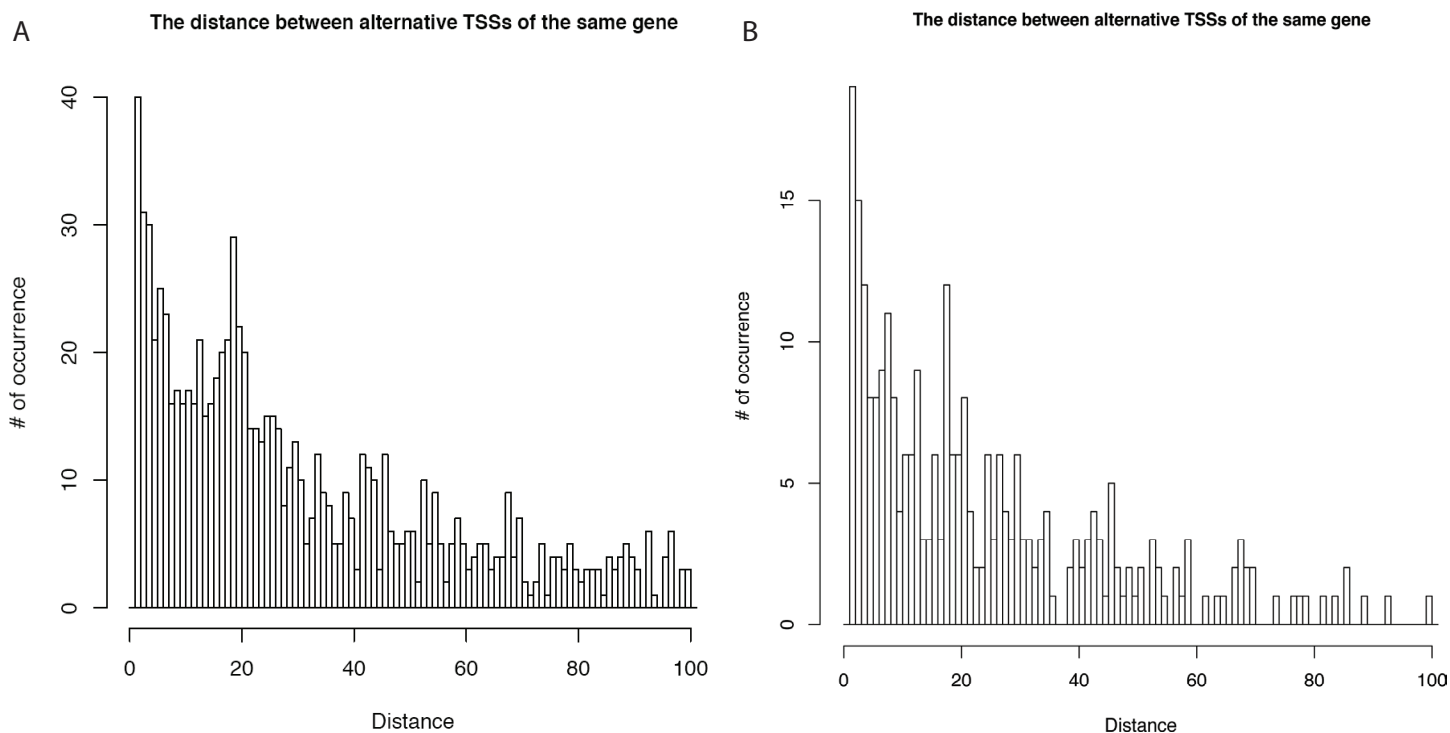


Figure S5: Histogram of distances between alternative TSSs for the same gene in the fly genome when (A) all promoters are considered and (B) when only the “cap-supported” promoters are considered. Alternative TSSs whose distances are shorter than a cutoff determined by the position of the second peak in the distribution were considered to derive from the same promoter.