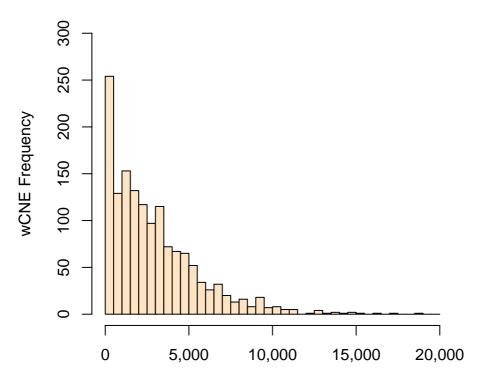
Figure S3. The distribution of distances between intergenic wCNEs and their nearest genes reveals many distal predicted *cis*-regulatory elements in the *C. elegans* genome. For each of the 1,460 intergenic wCNEs, we calculated the distance from the nearest transcription start site of a protein-coding gene.



Distance from transcription start site of nearest gene (bp)