



Supplemental Figure S35. Mapping resolution. (A) Distance between genome location of *TK1* and its retention peak, all samples shown. (B) Distance between assigned centromere genome locations and retention peaks, all samples shown. (C) Distance between *TK1* and its retention peak averaged across the six RH lineages. (D) Distance between assigned centromere locations and retention peaks averaged across the six RH lineages. (E) Distance between assigned centromere locations and retention peaks averaged across chromosomes.