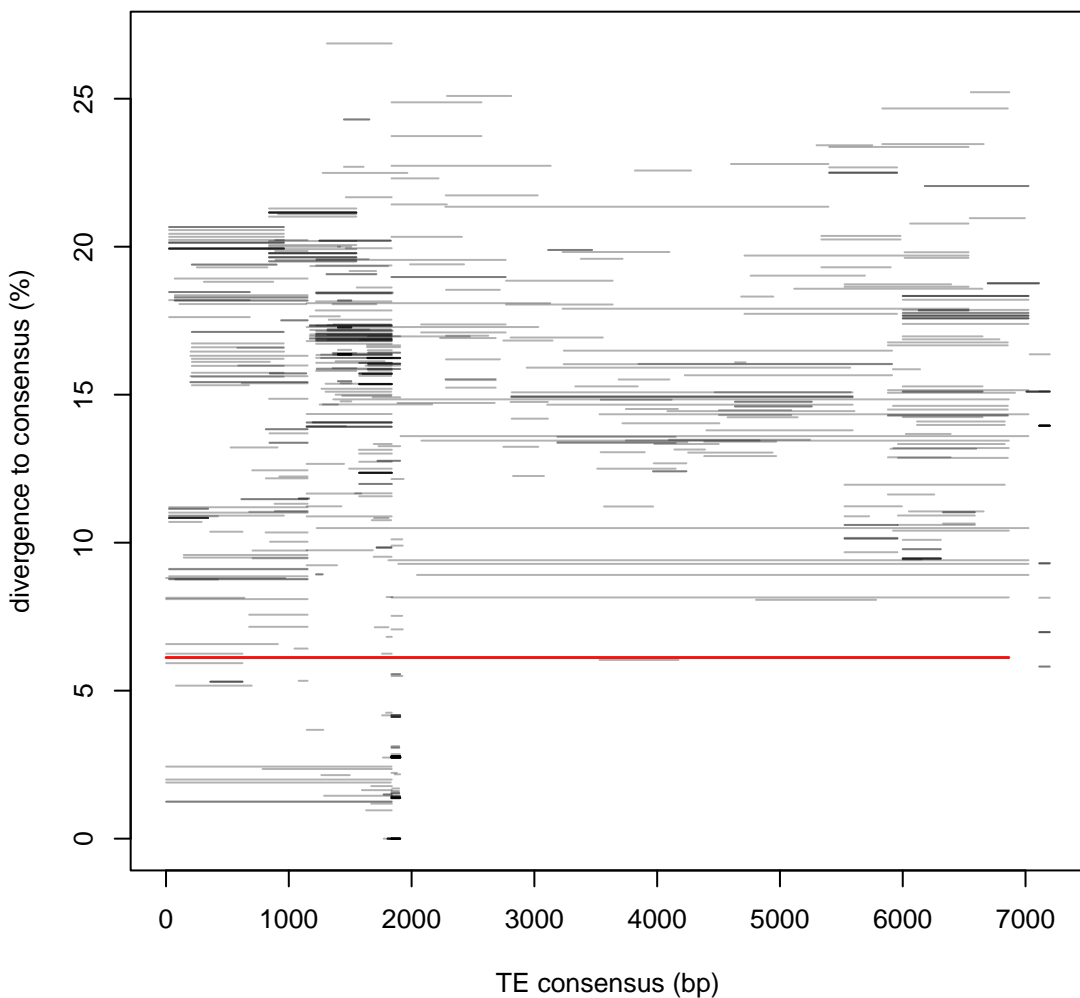
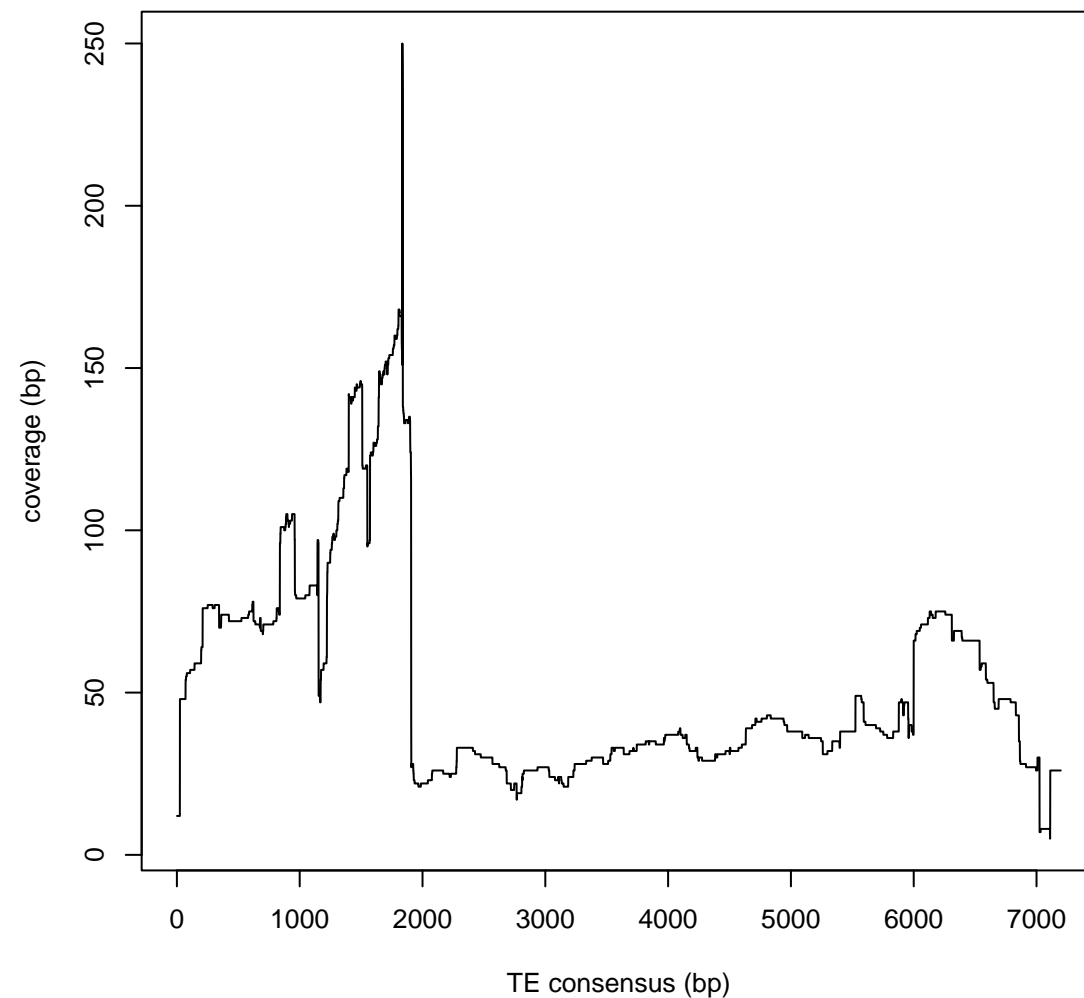


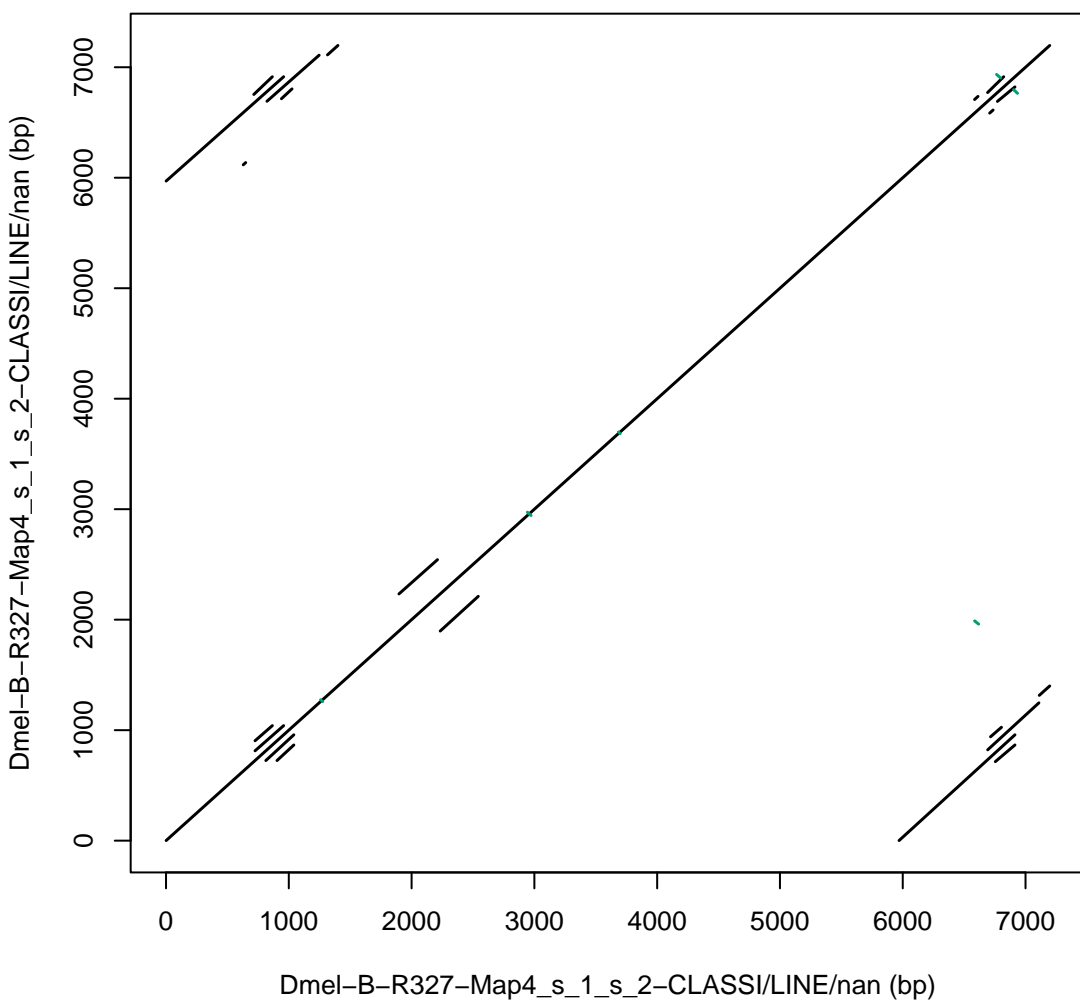
TE: Dmel-B-R327-Map4_s_1_s_2-CLASSI/LINE/nan
consensus size: 7197bp; fragments: 651; full length: 1 (>=6477.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

