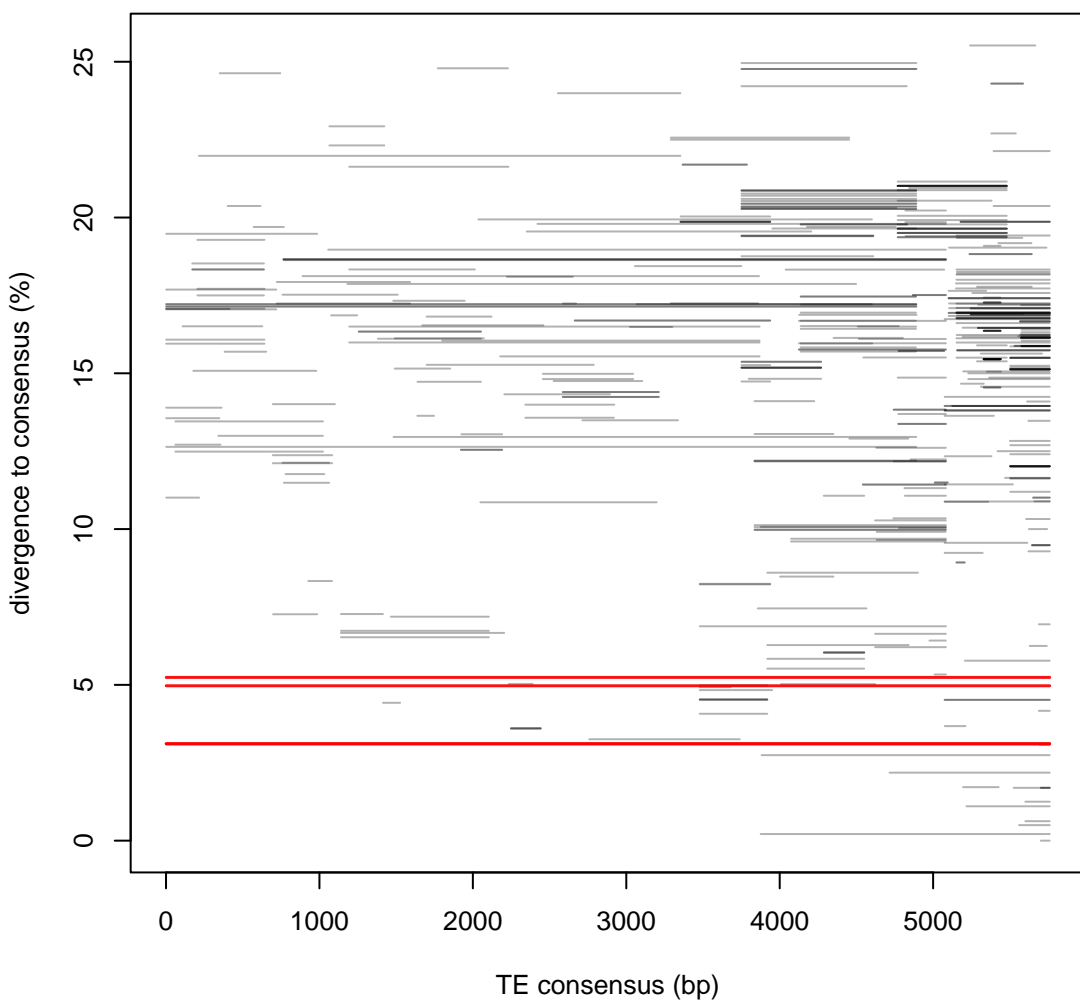
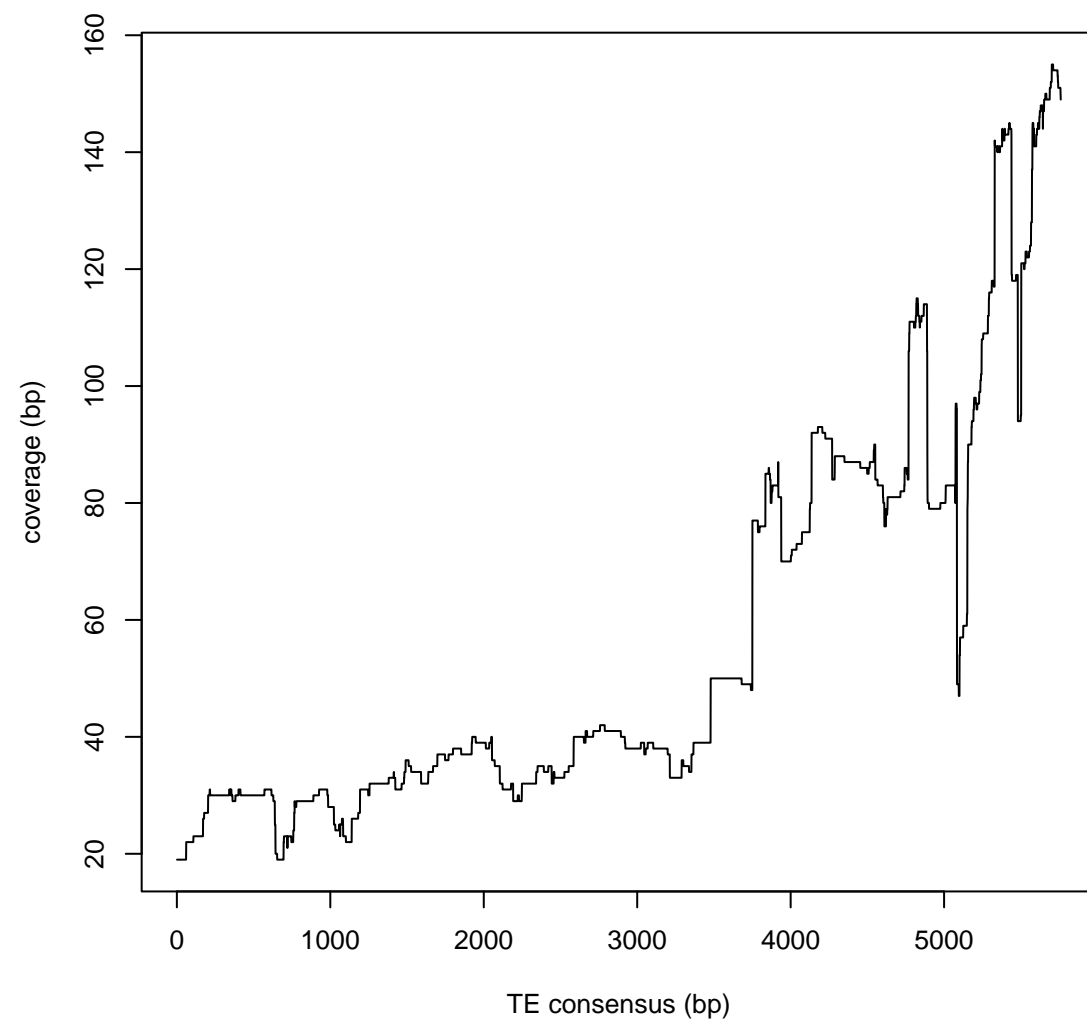


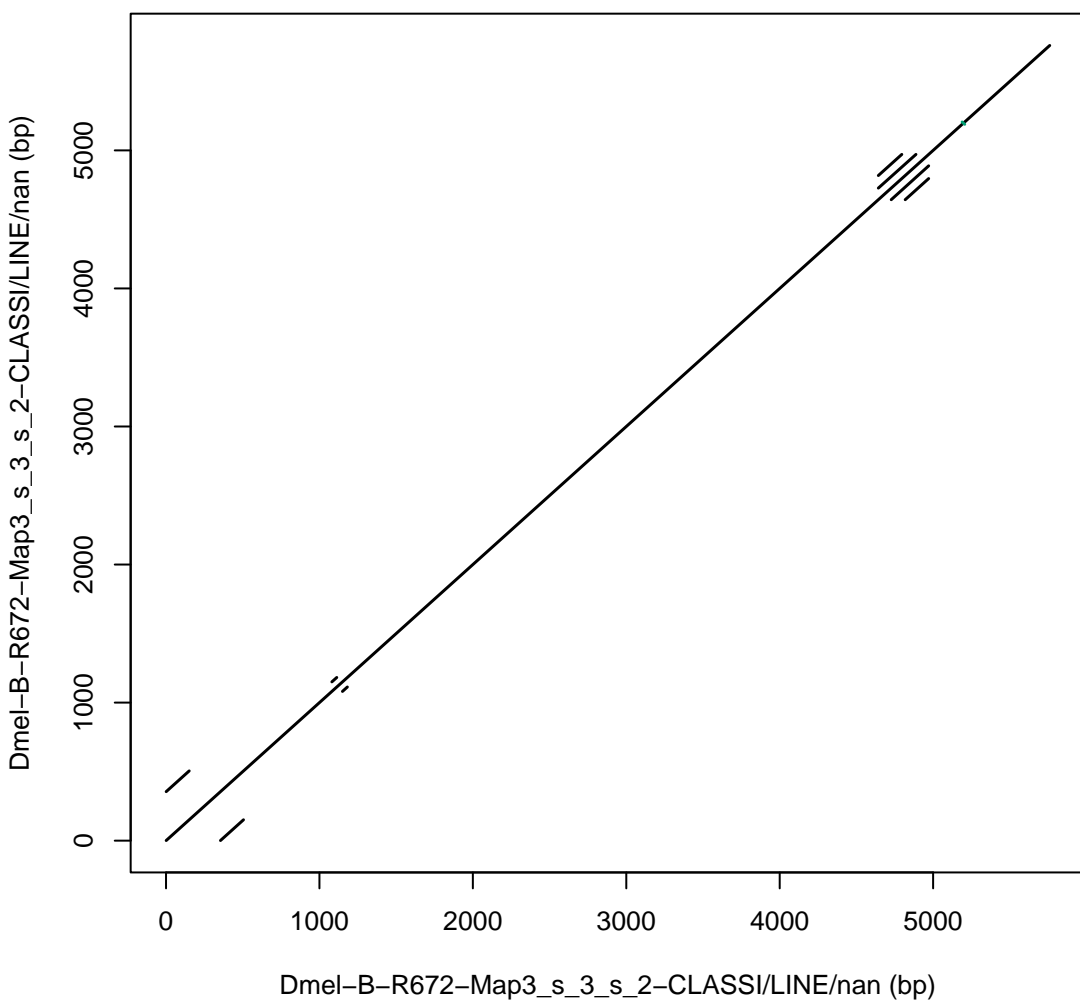
TE: Dmel-B-R672-Map3_s_3_s_2-CLASSI/LINE/nan
consensus size: 5760bp; fragments: 468; full length: 4 (>=5184bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

