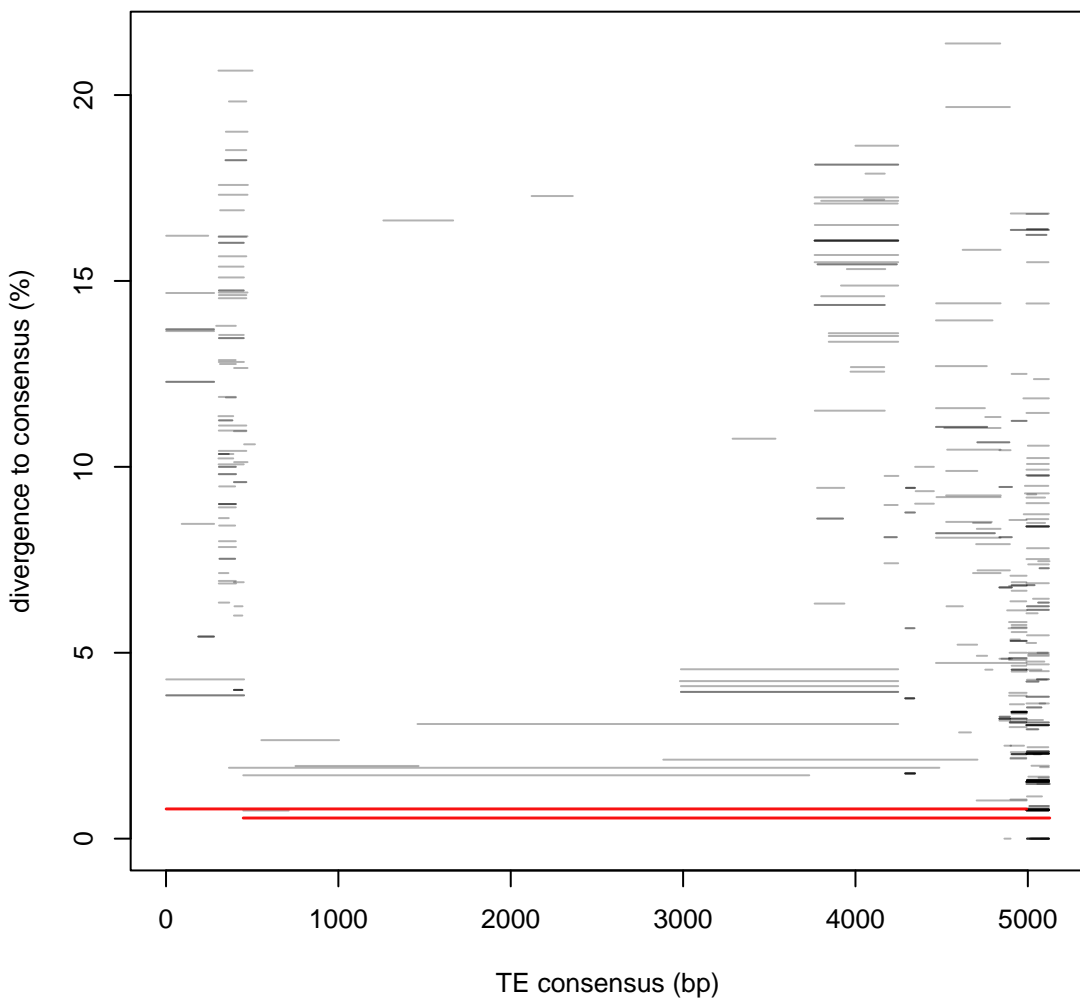
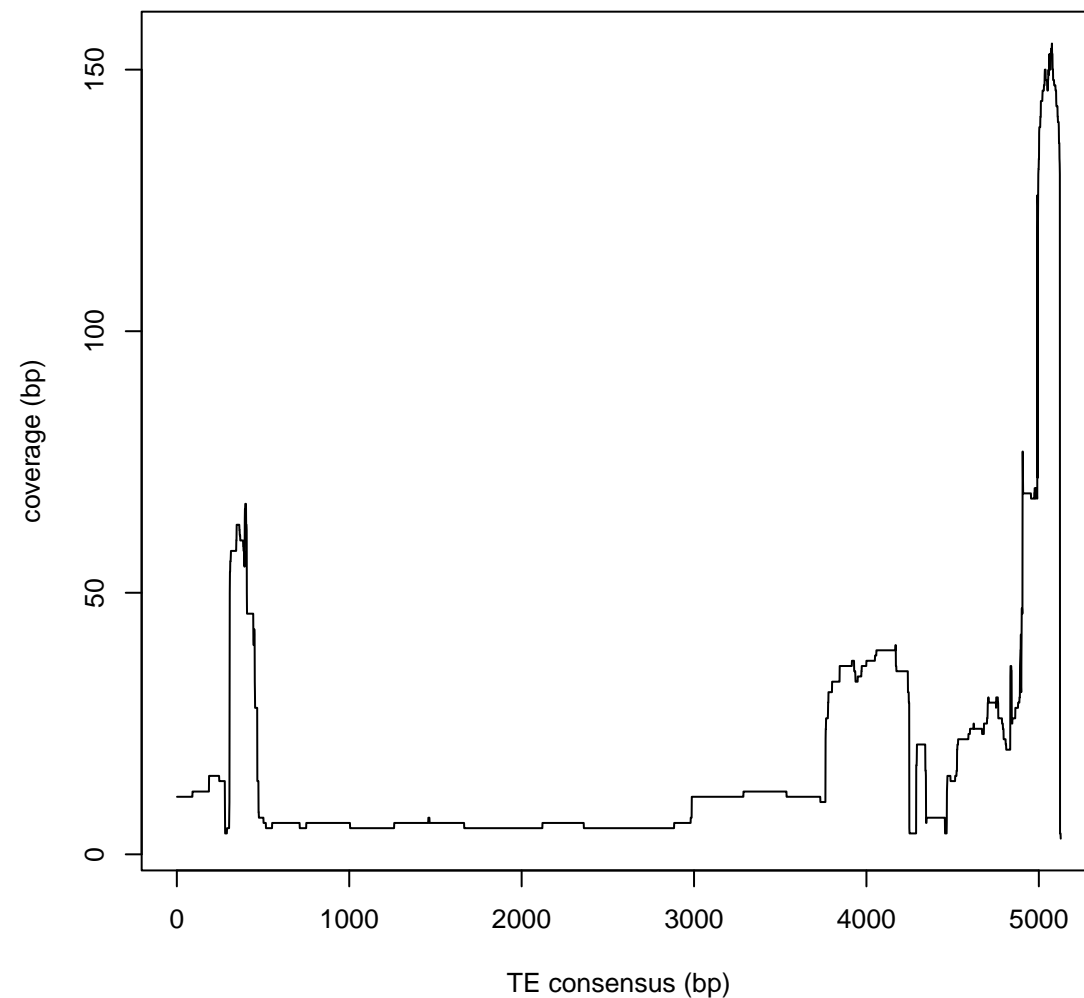


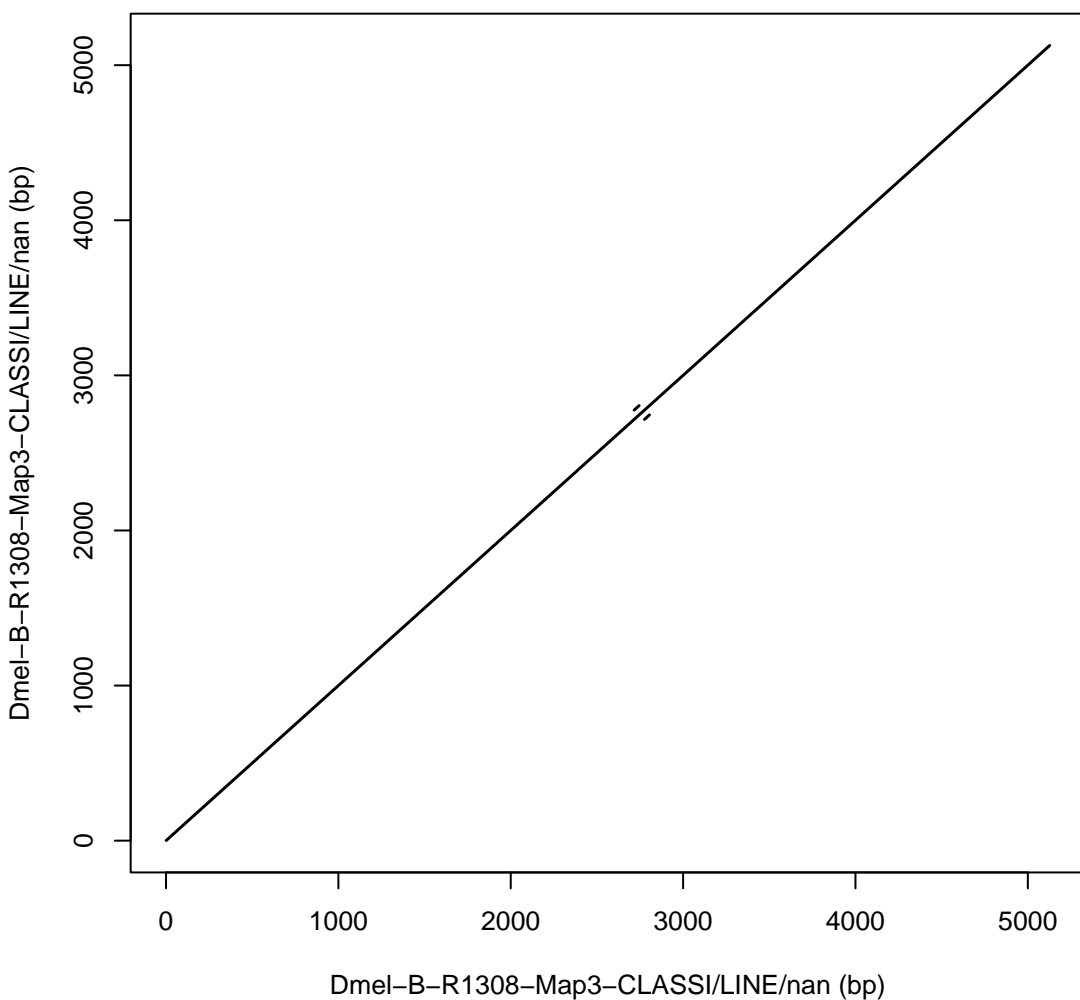
TE: Dmel-B-R1308-Map3-CLASSI/LINE/nan
consensus size: 5127bp; fragments: 439; full length: 2 (>=4614.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

