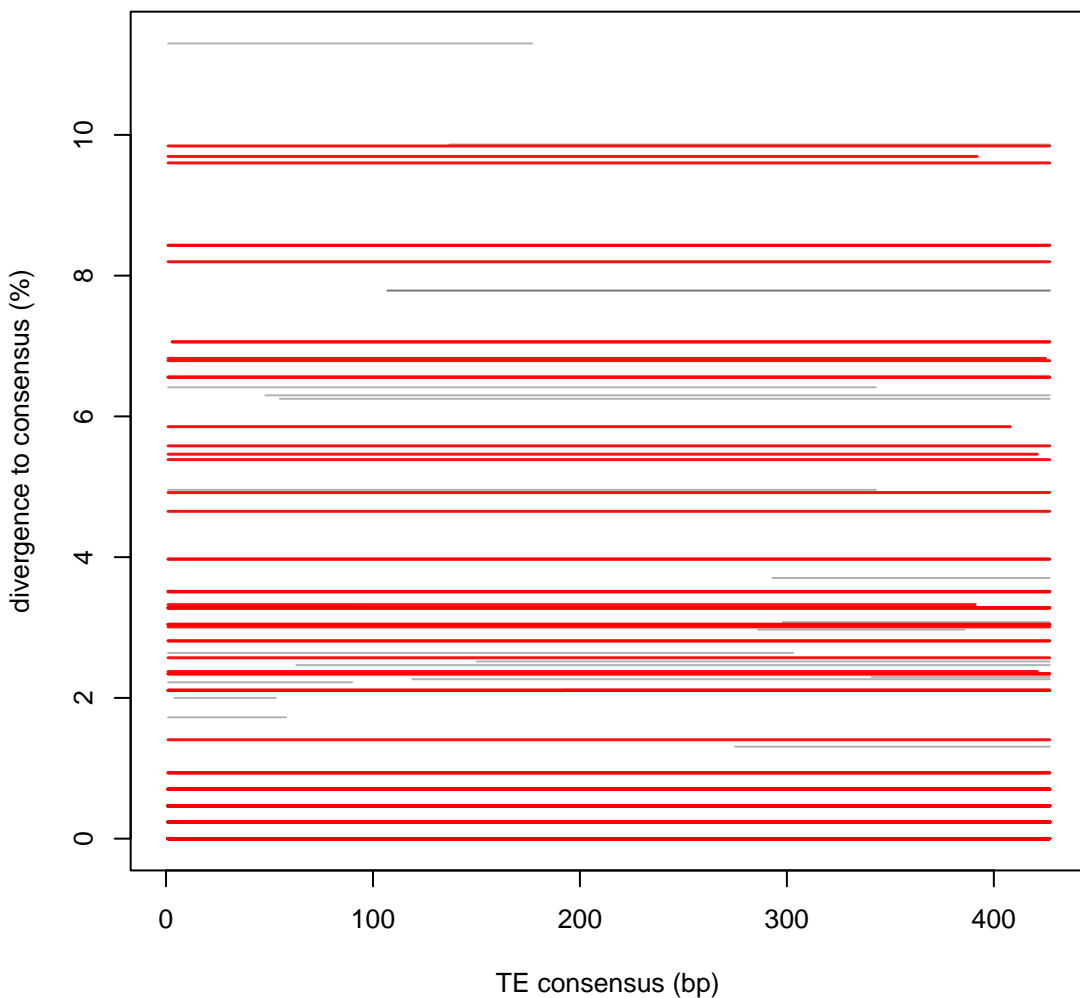
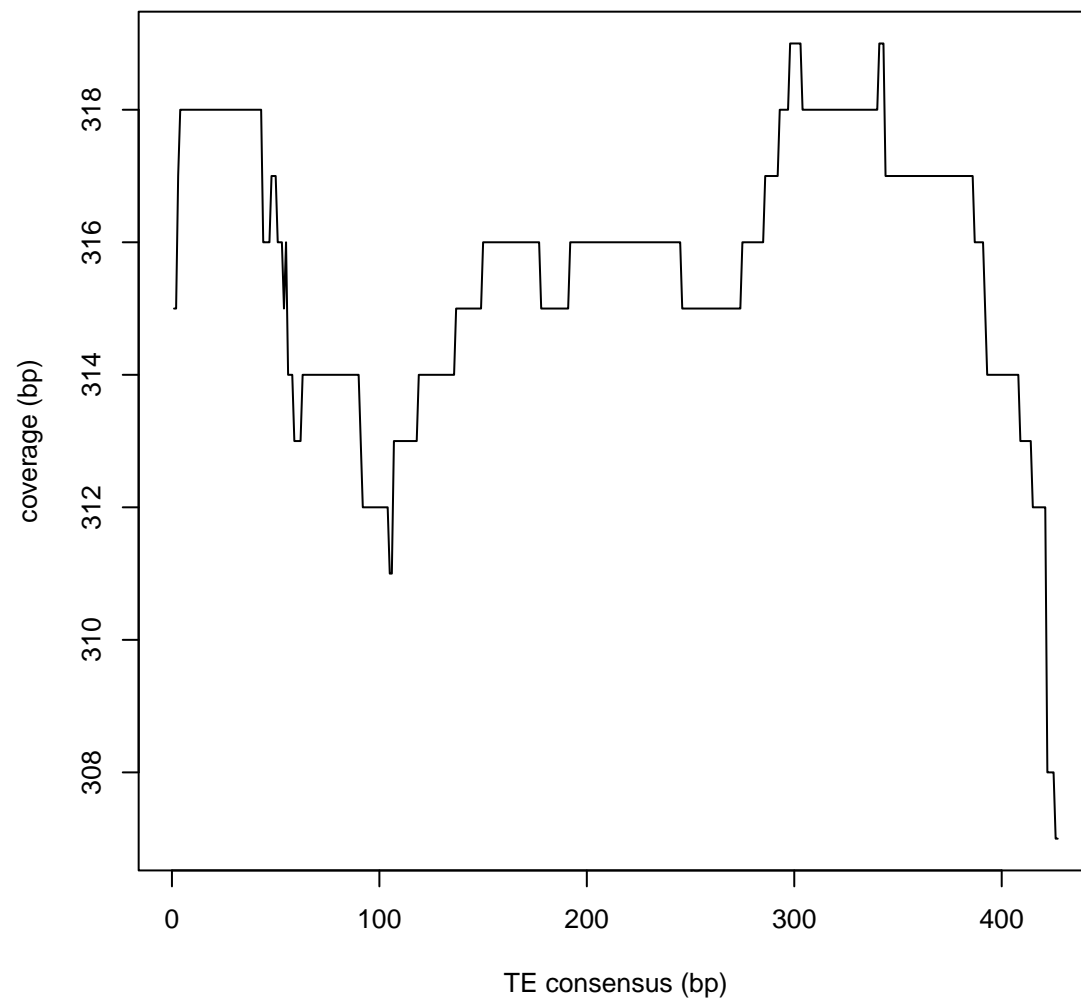


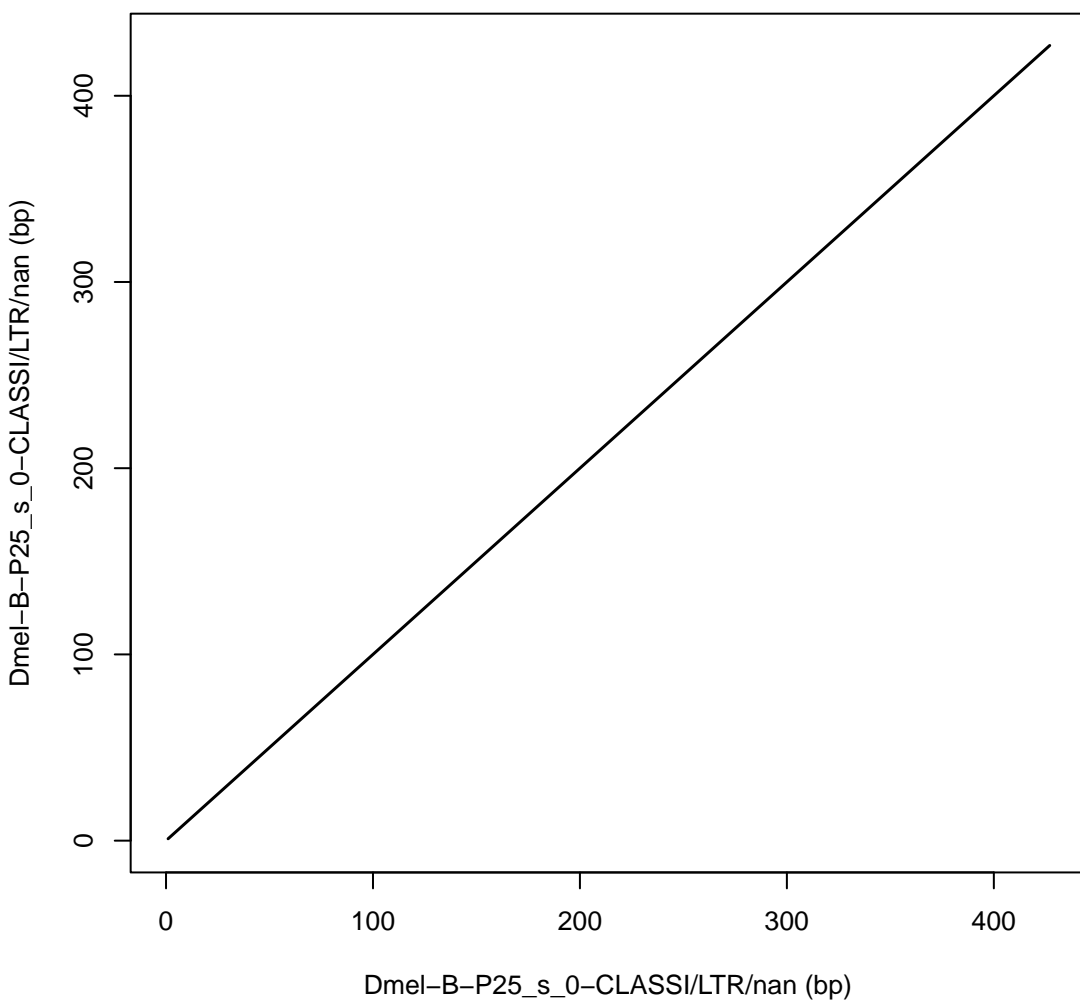
TE: Dmel-B-P25_s_0-CLASSI/LTR/nan
consensus size: 427bp; fragments: 332; full length: 303 (>=384.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

