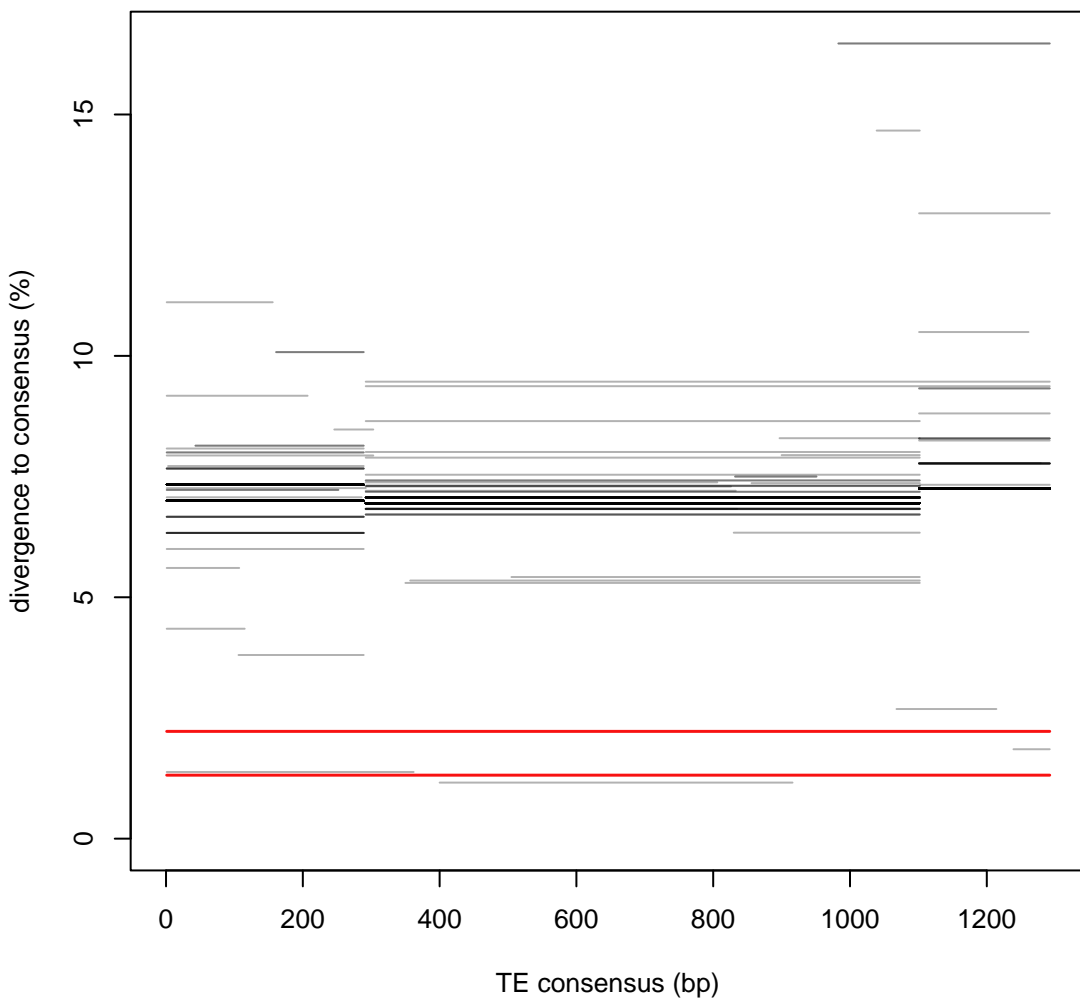
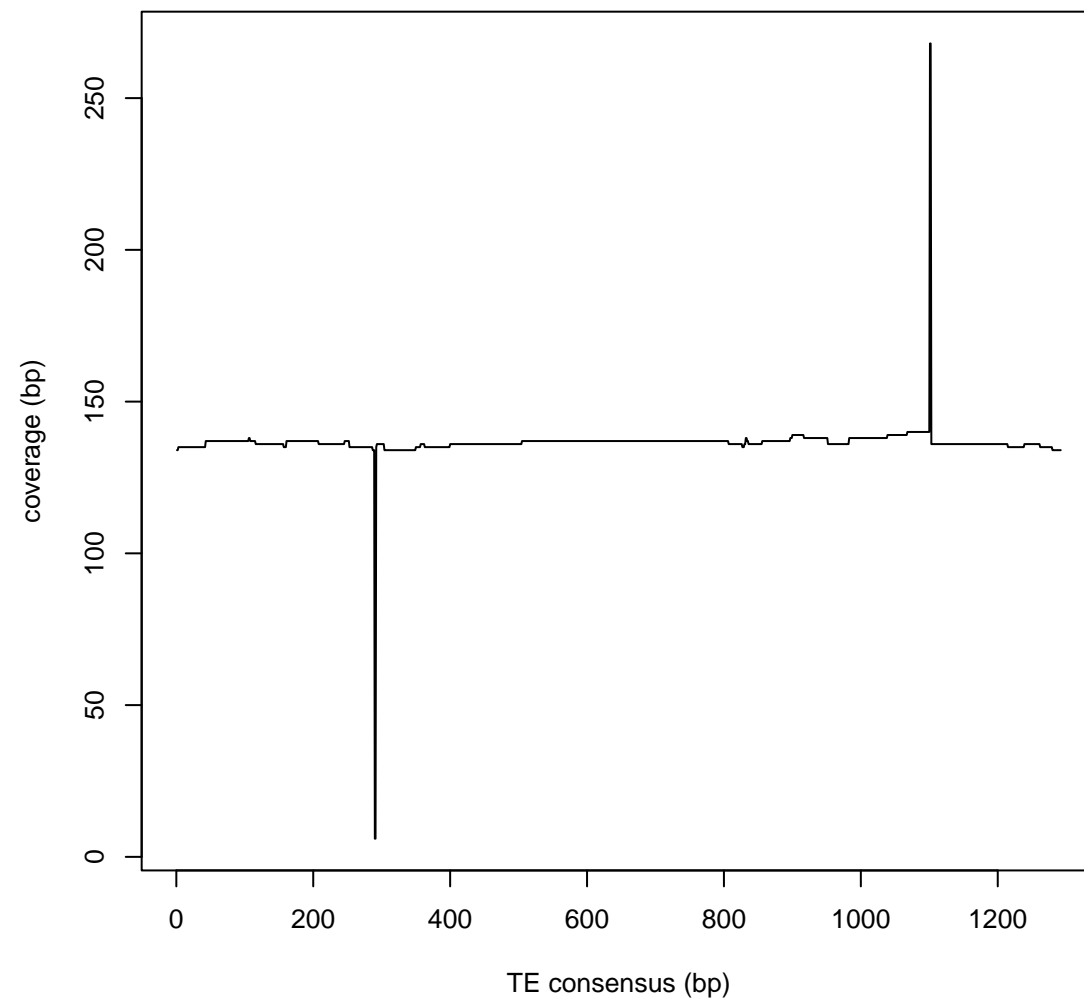


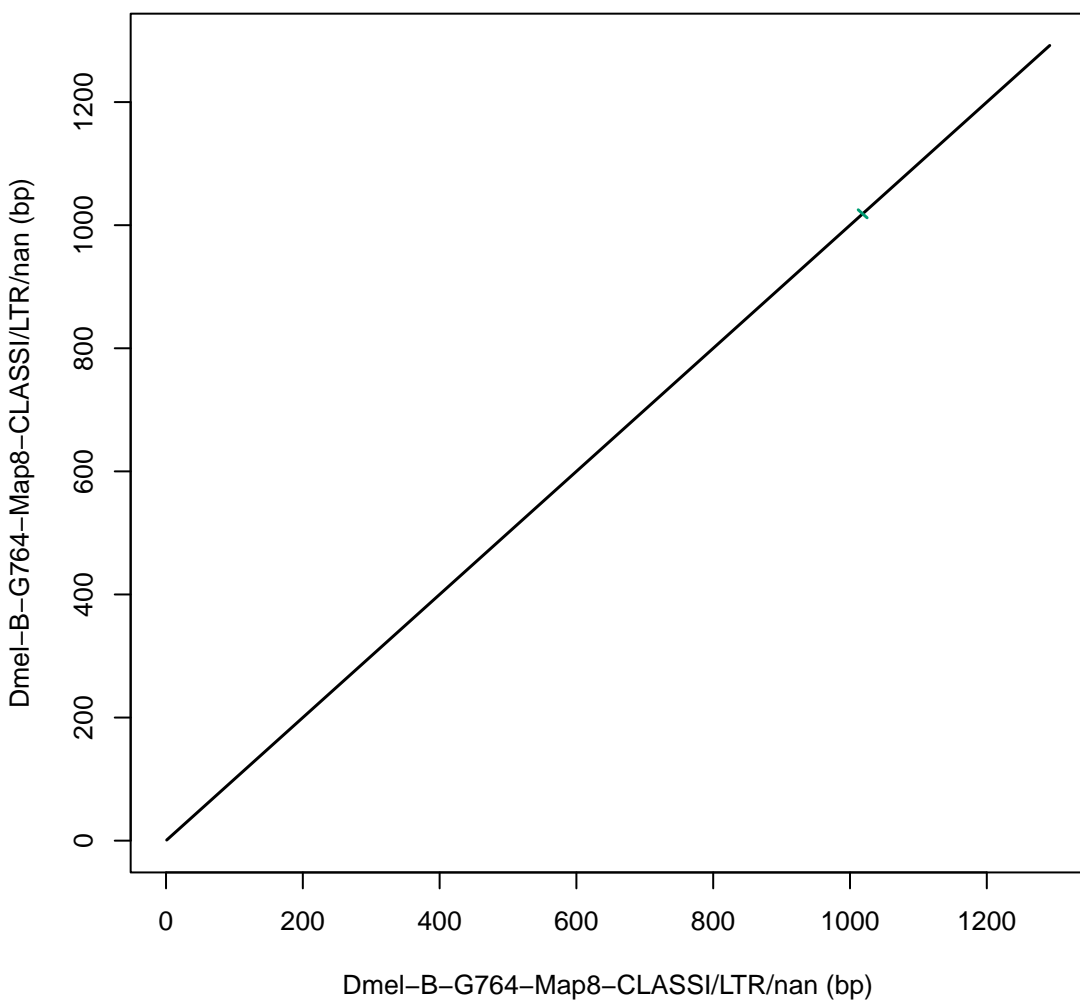
TE: Dmel-B-G764-Map8-CLASSI/LTR/nan
consensus size: 1292bp; fragments: 416; full length: 2 (>=1162.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

