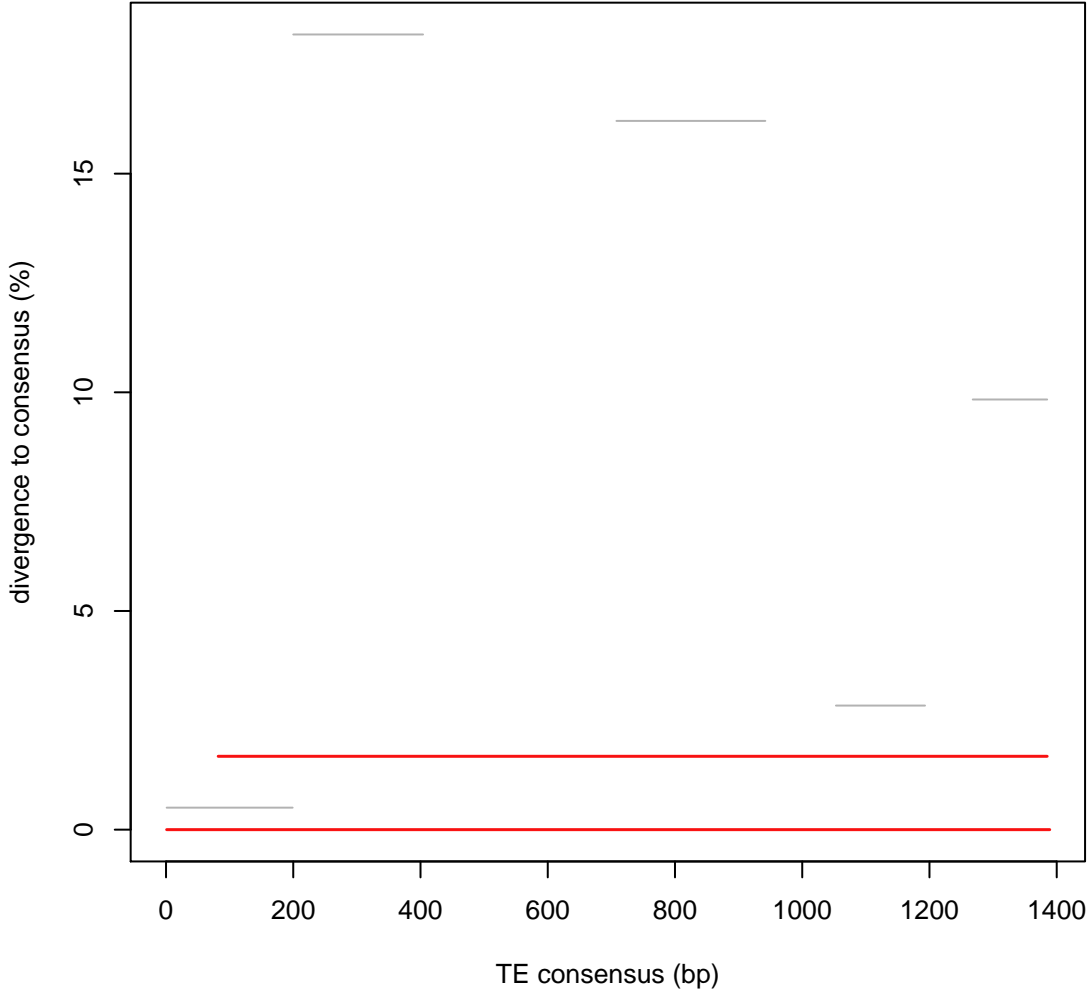
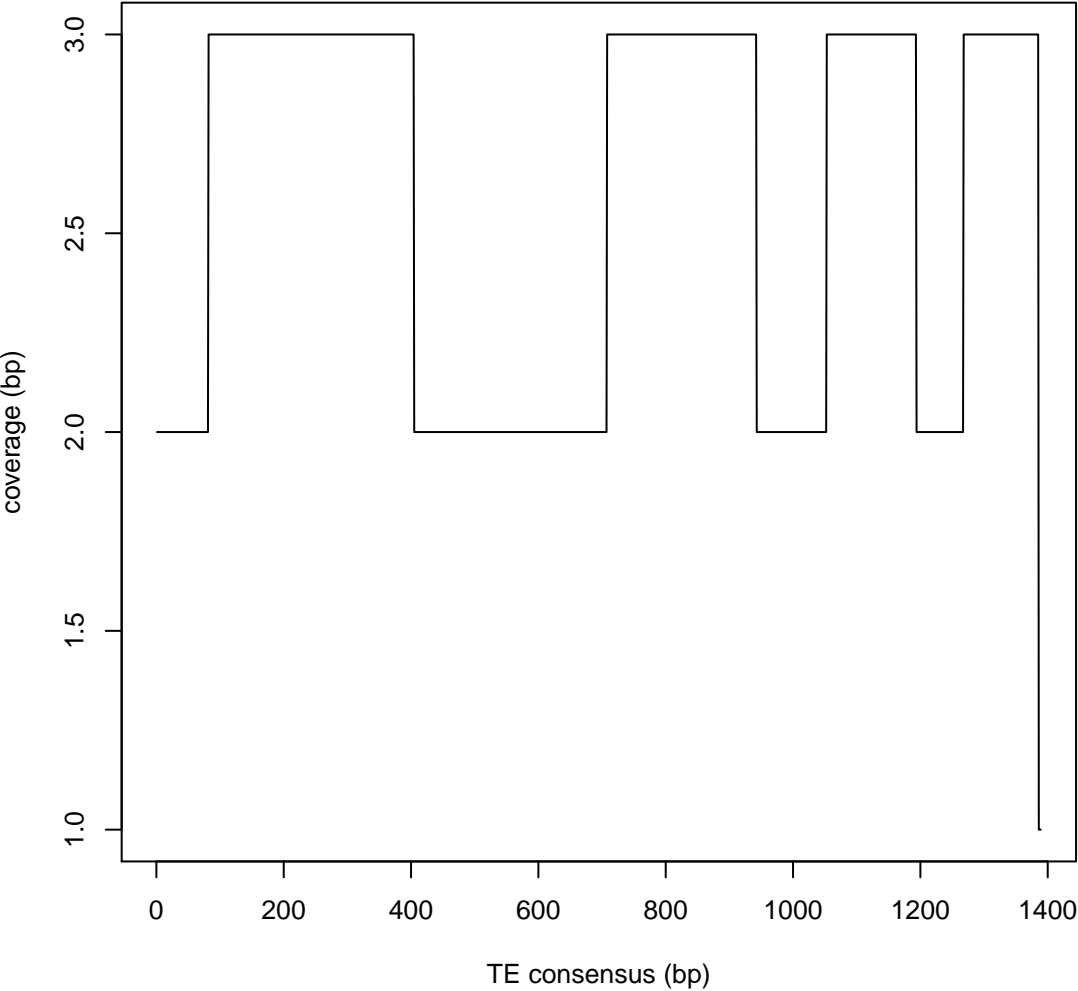


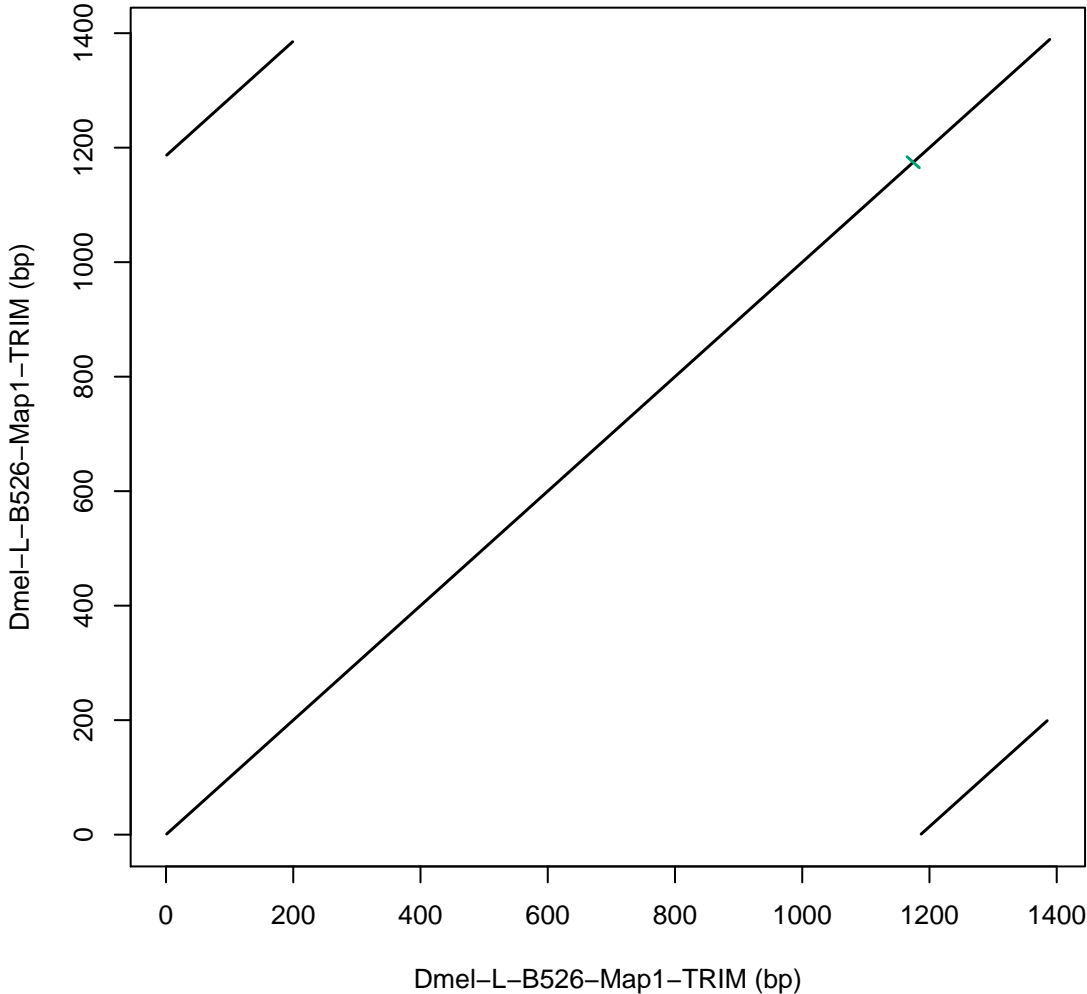
TE: Dmel-L-B526-Map1-TRIM
consensus size: 1389bp; fragments: 7; full length: 2 (≥ 1250.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

