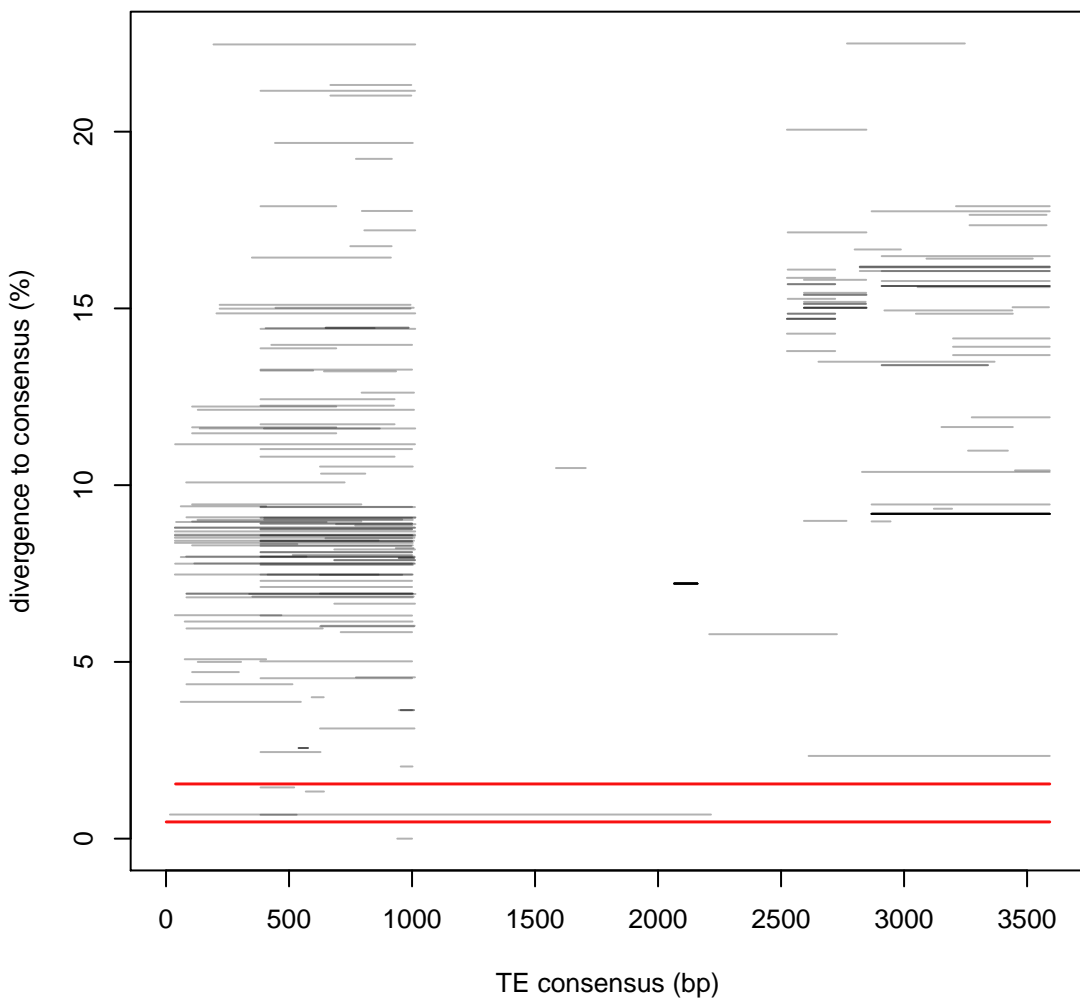
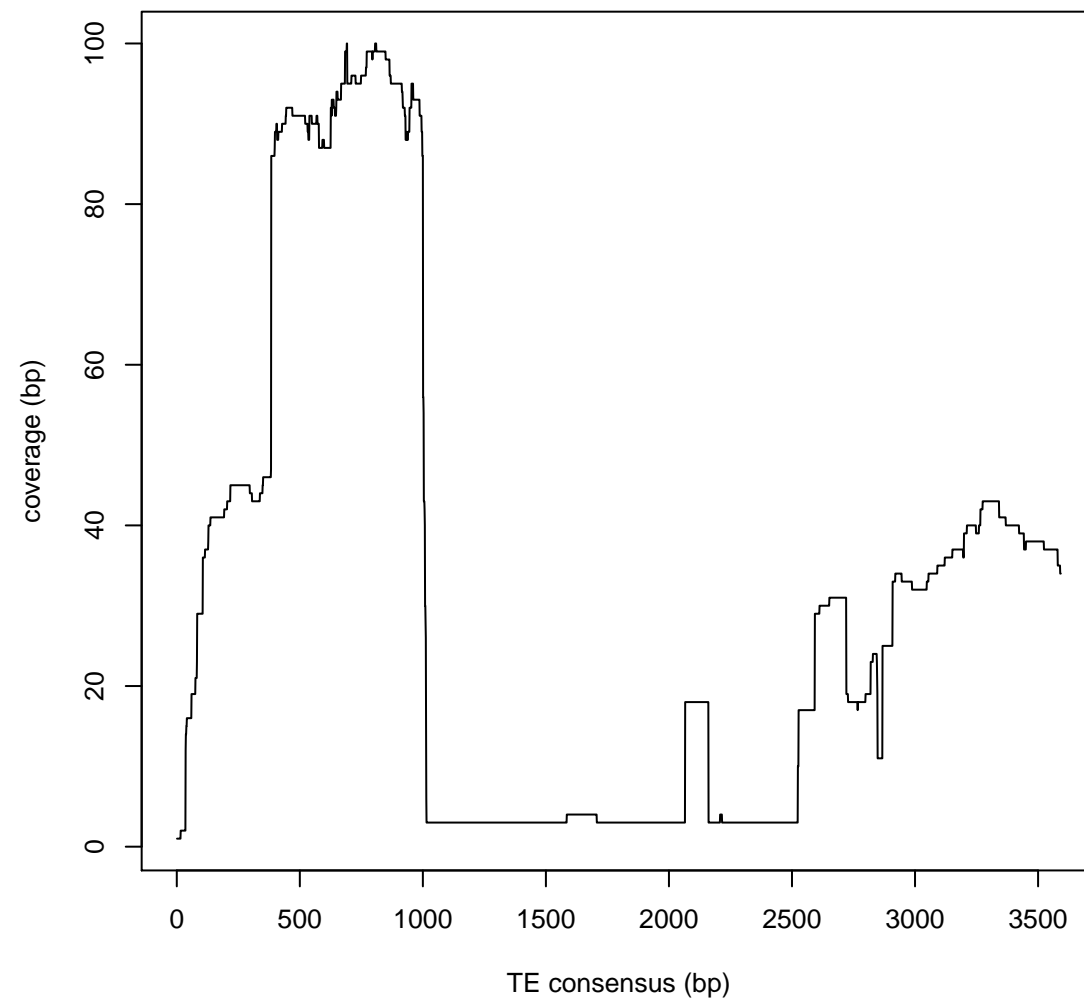


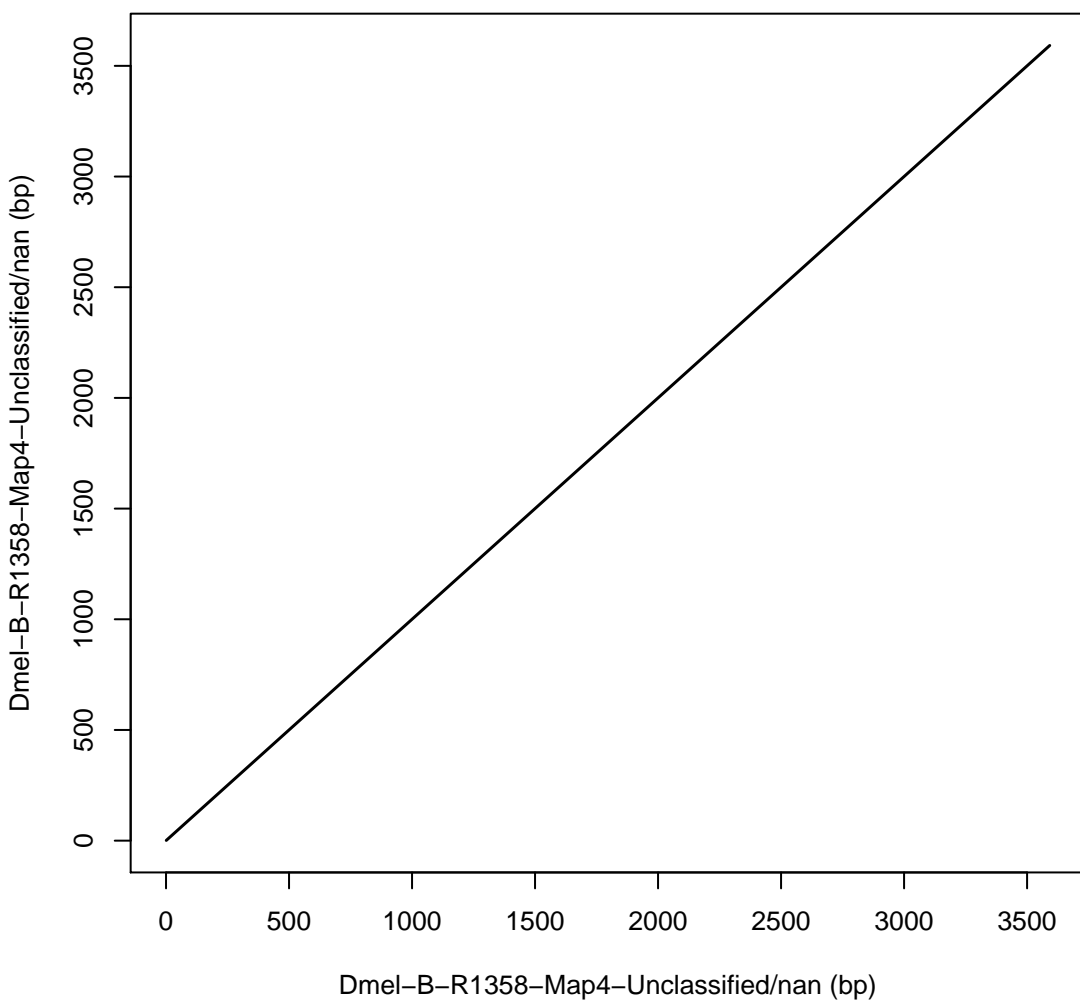
TE: Dmel-B-R1358-Map4-Unclassified/nan  
consensus size: 3592bp; fragments: 226; full length: 2 (>=3232.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

