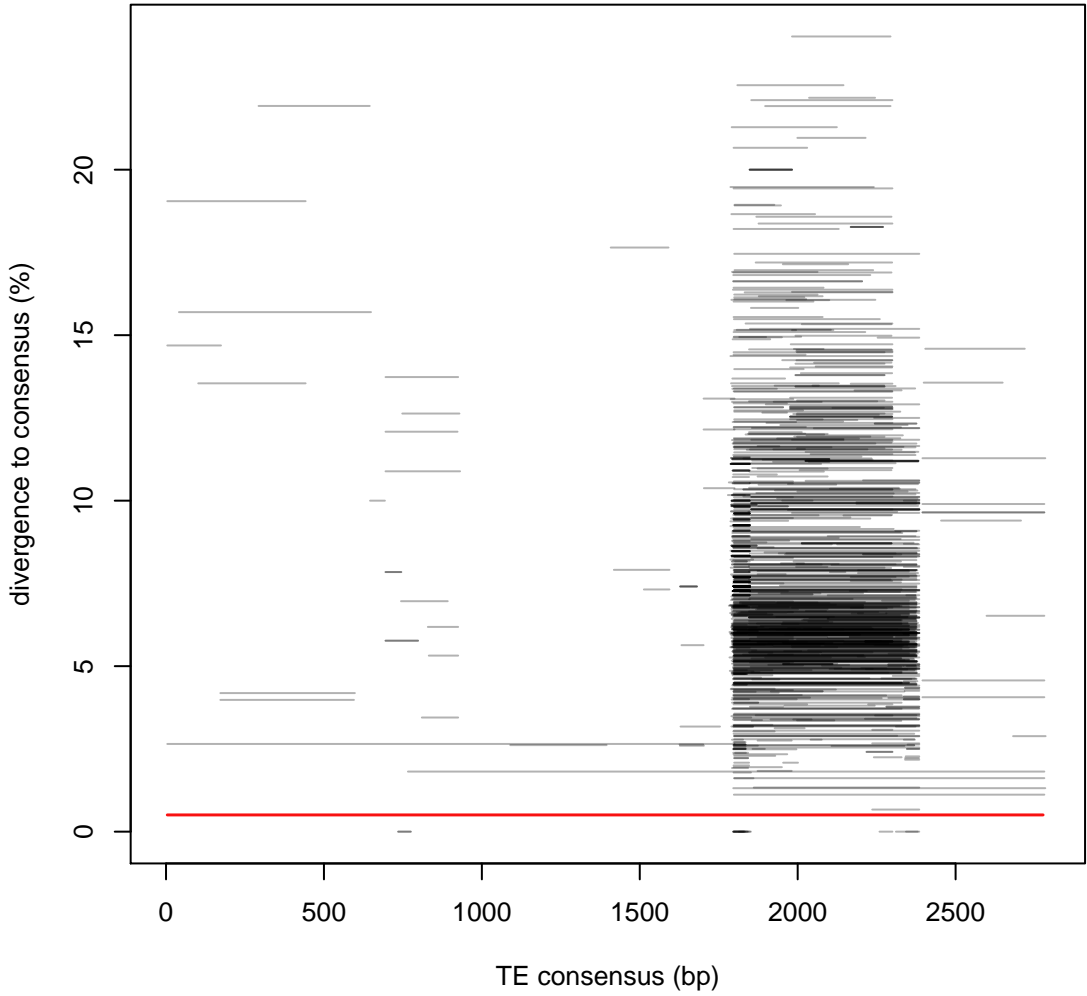
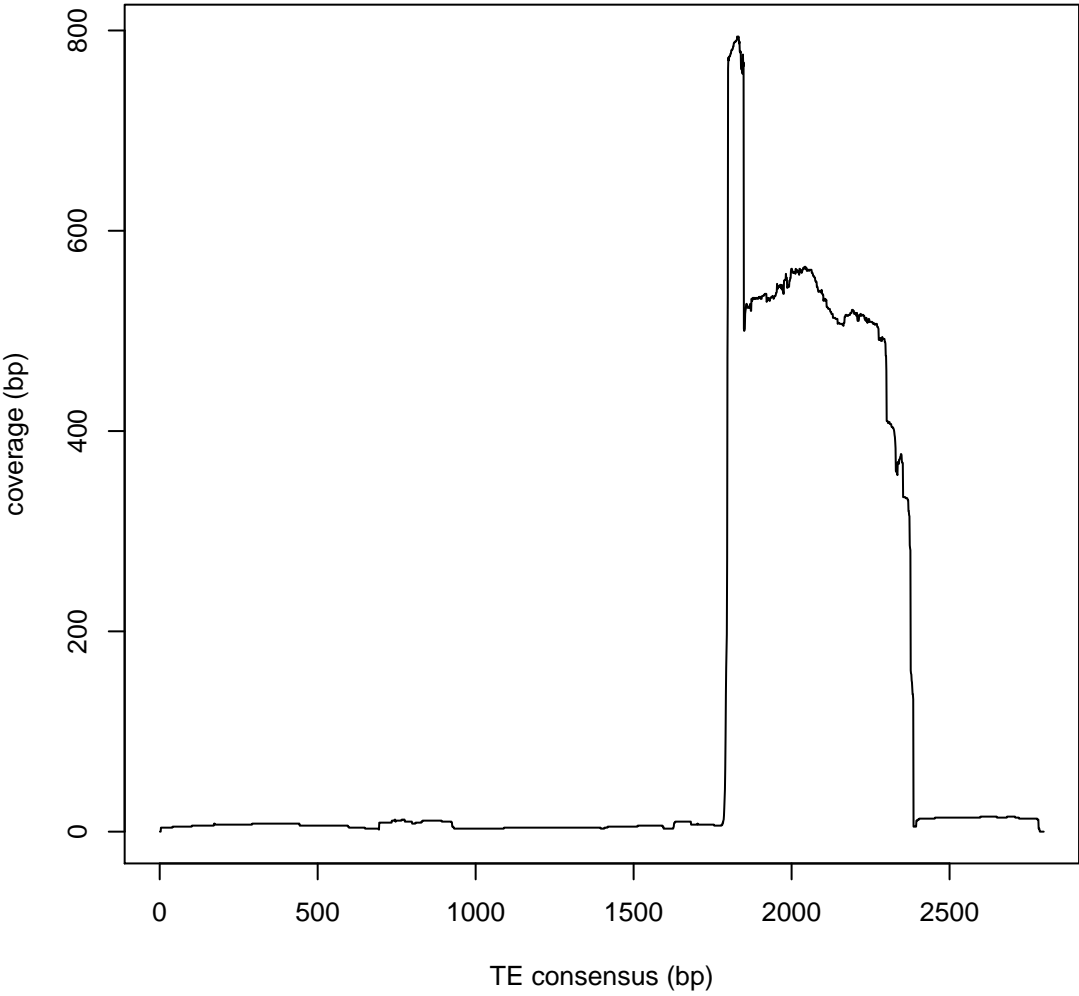


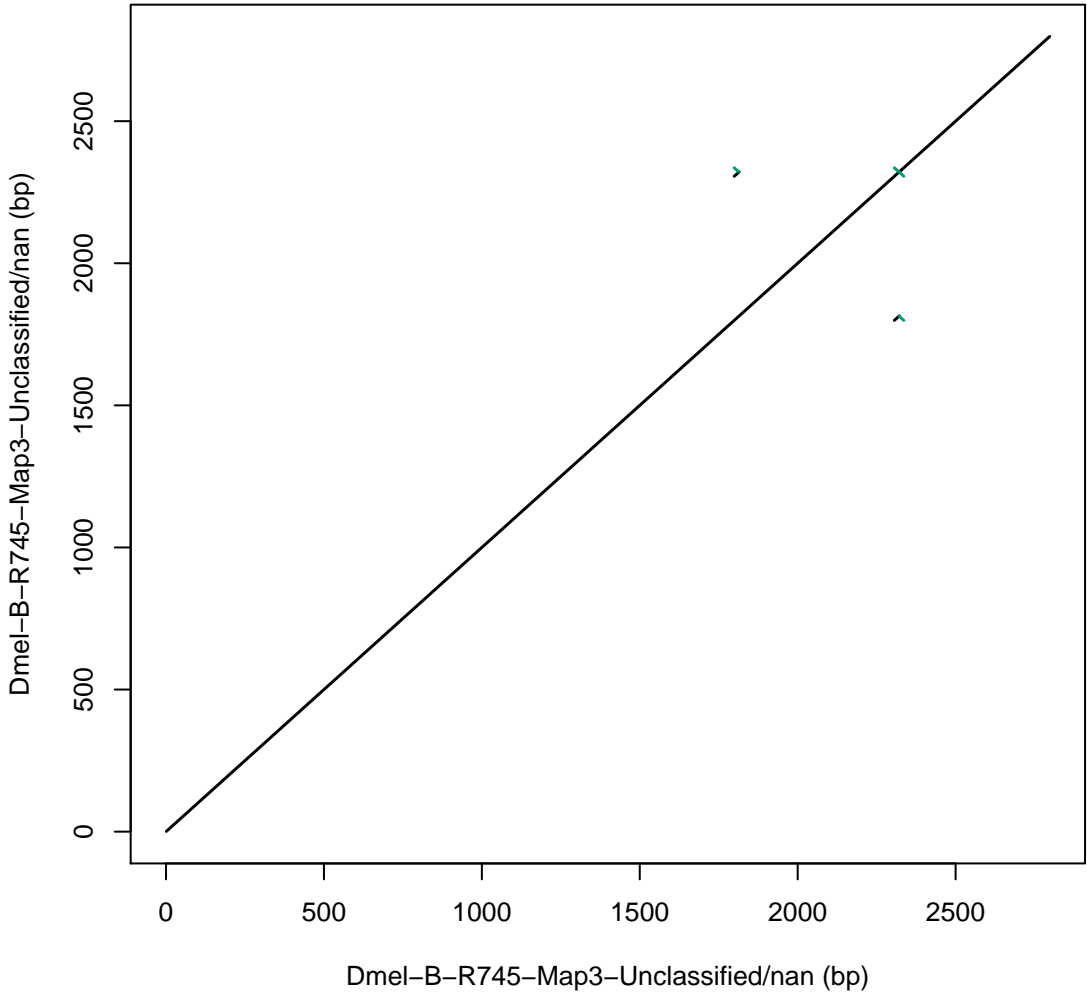
TE: Dmel-B-R745-Map3-Unclassified/nan
consensus size: 2798bp; fragments: 1185; full length: 1 (>=2518.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

