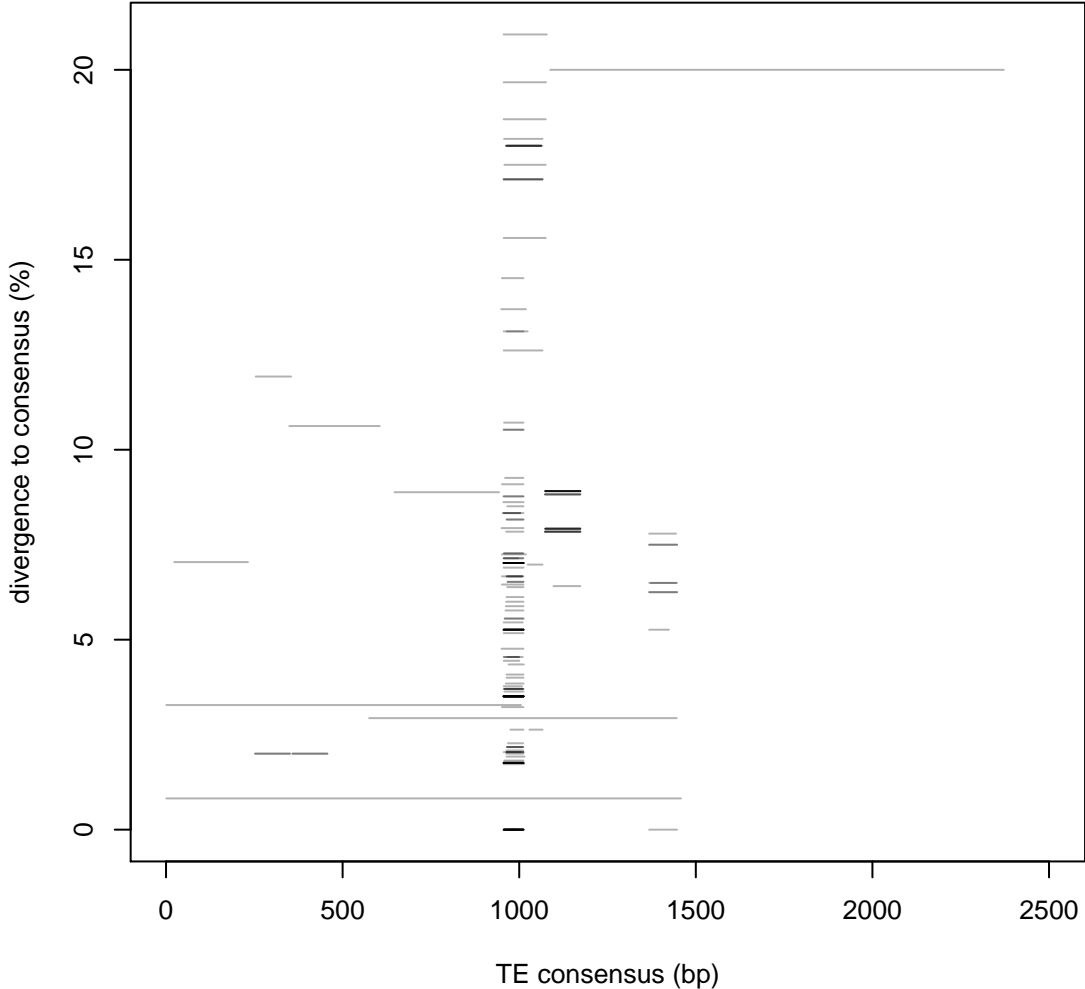
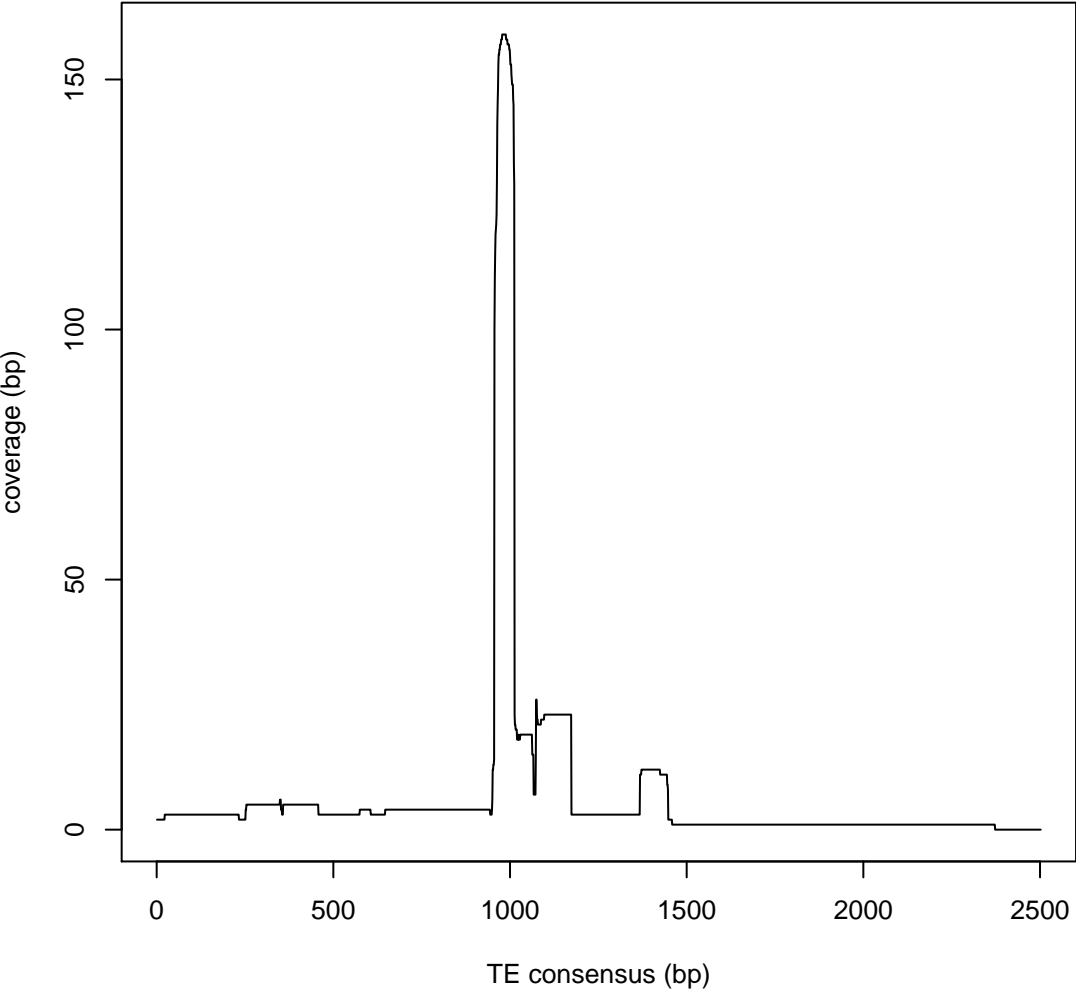


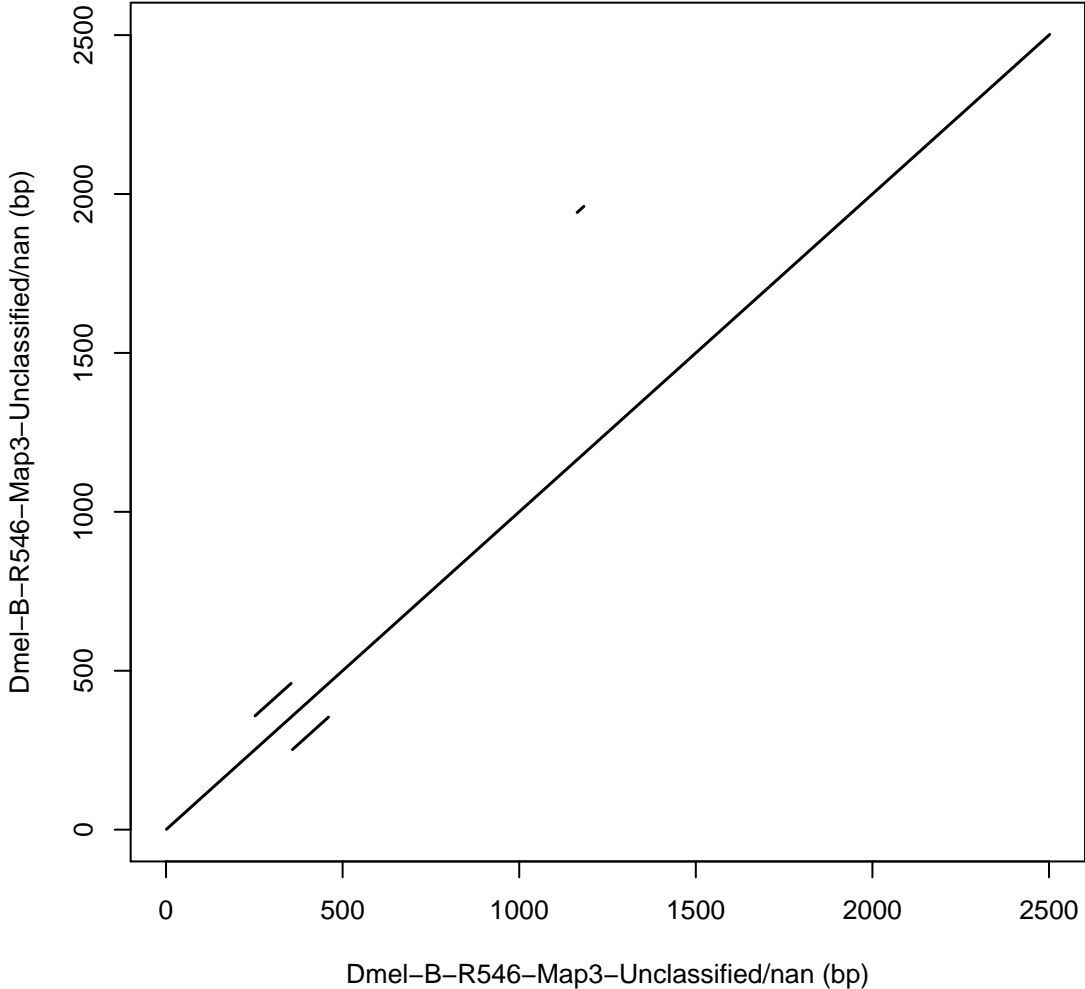
TE: Dmel-B-R546-Map3-Unclassified/nan
consensus size: 2502bp; fragments: 199; full length: 0 (>=2251.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

