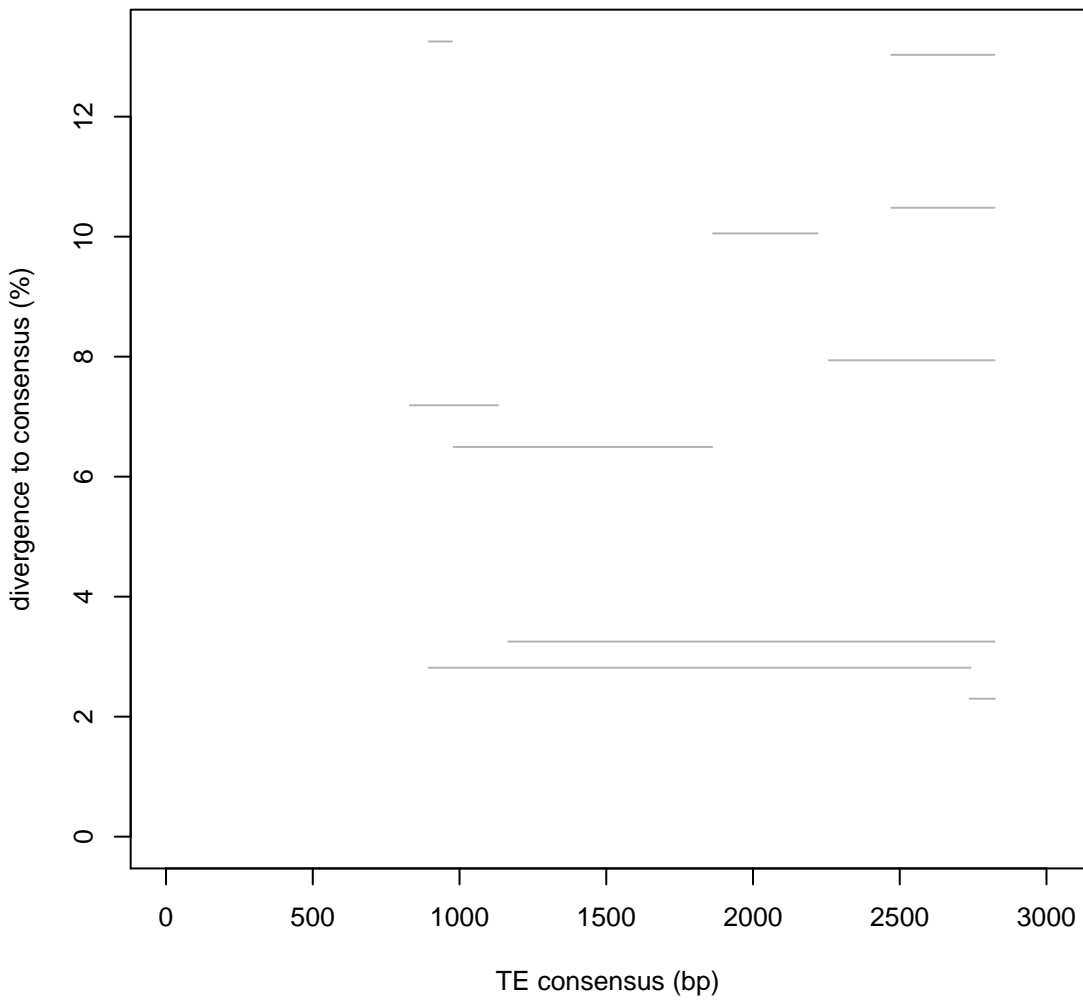
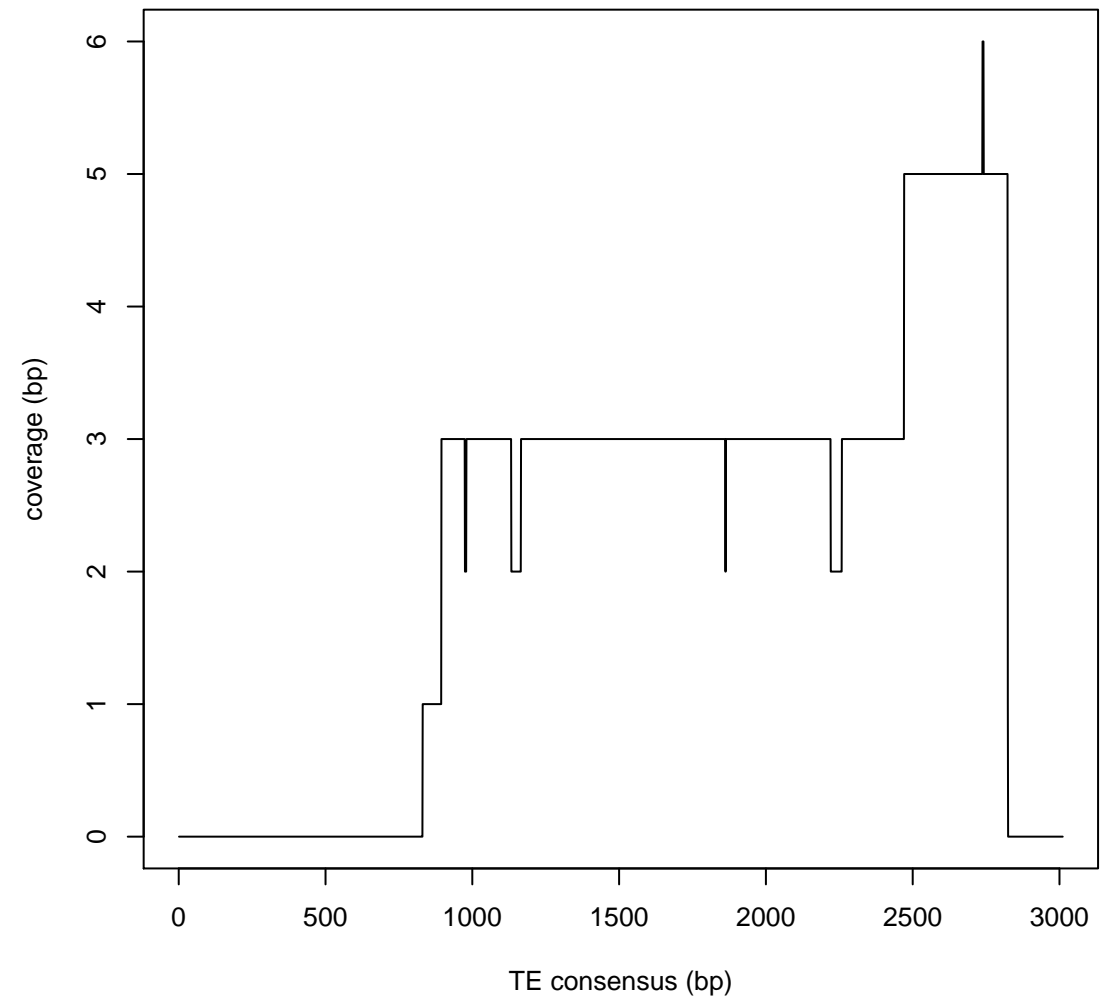


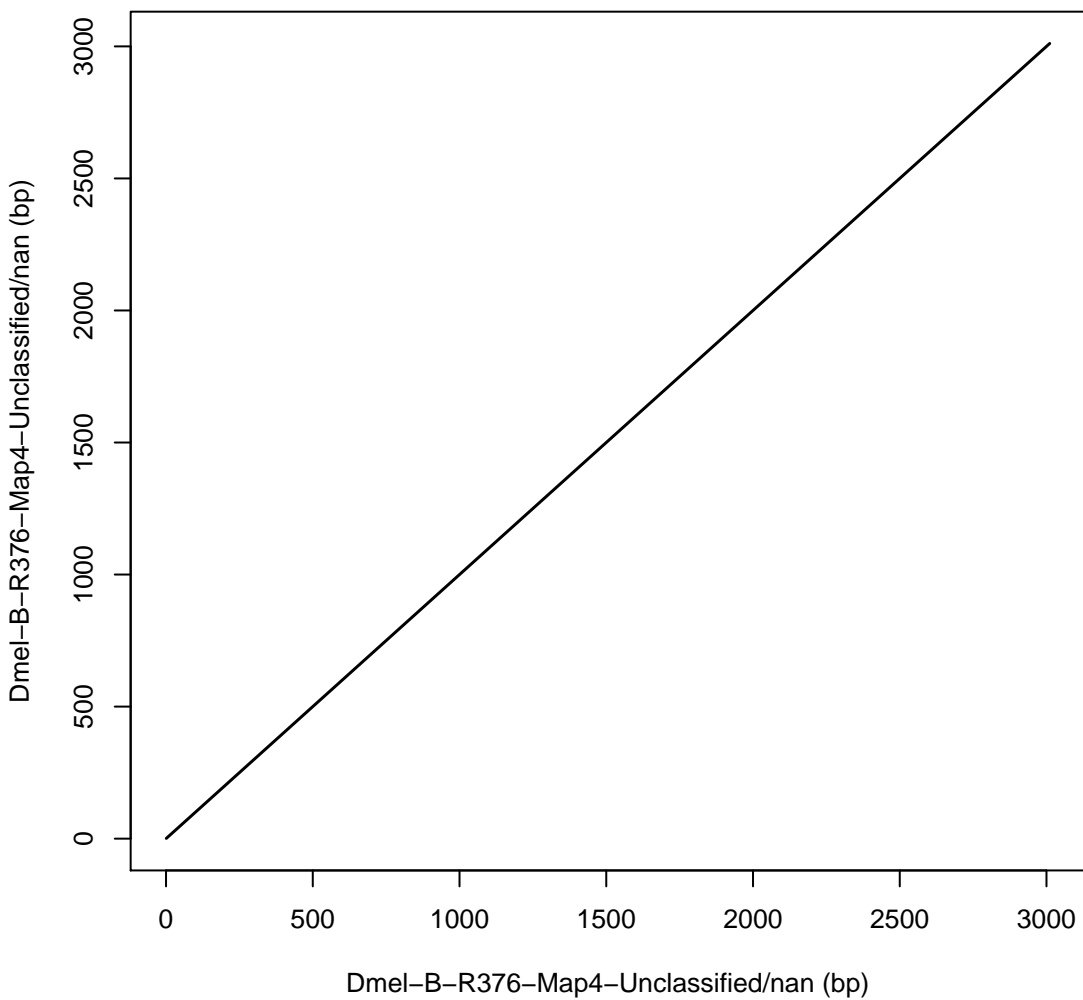
TE: Dmel-B-R376-Map4-Unclassified/nan
consensus size: 3011bp; fragments: 10; full length: 0 (>=2709.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

