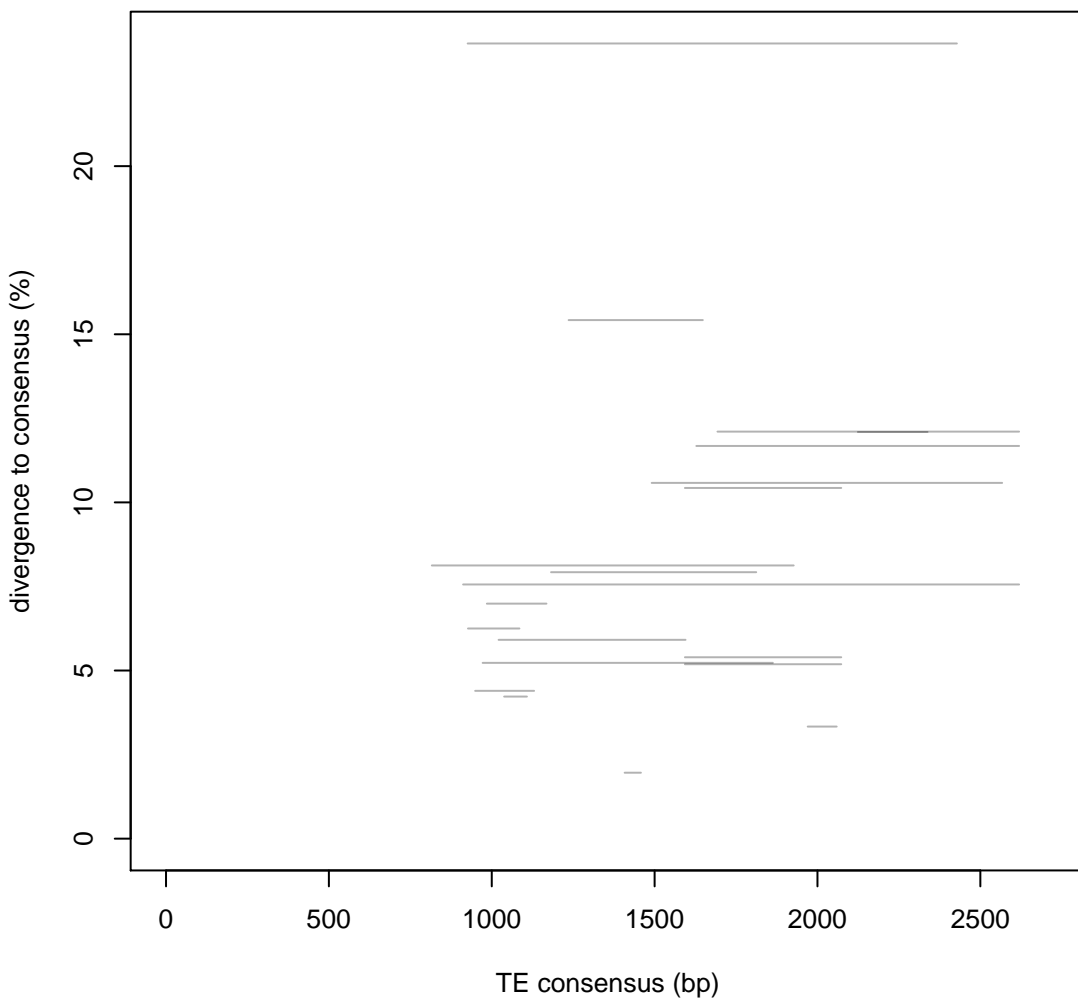
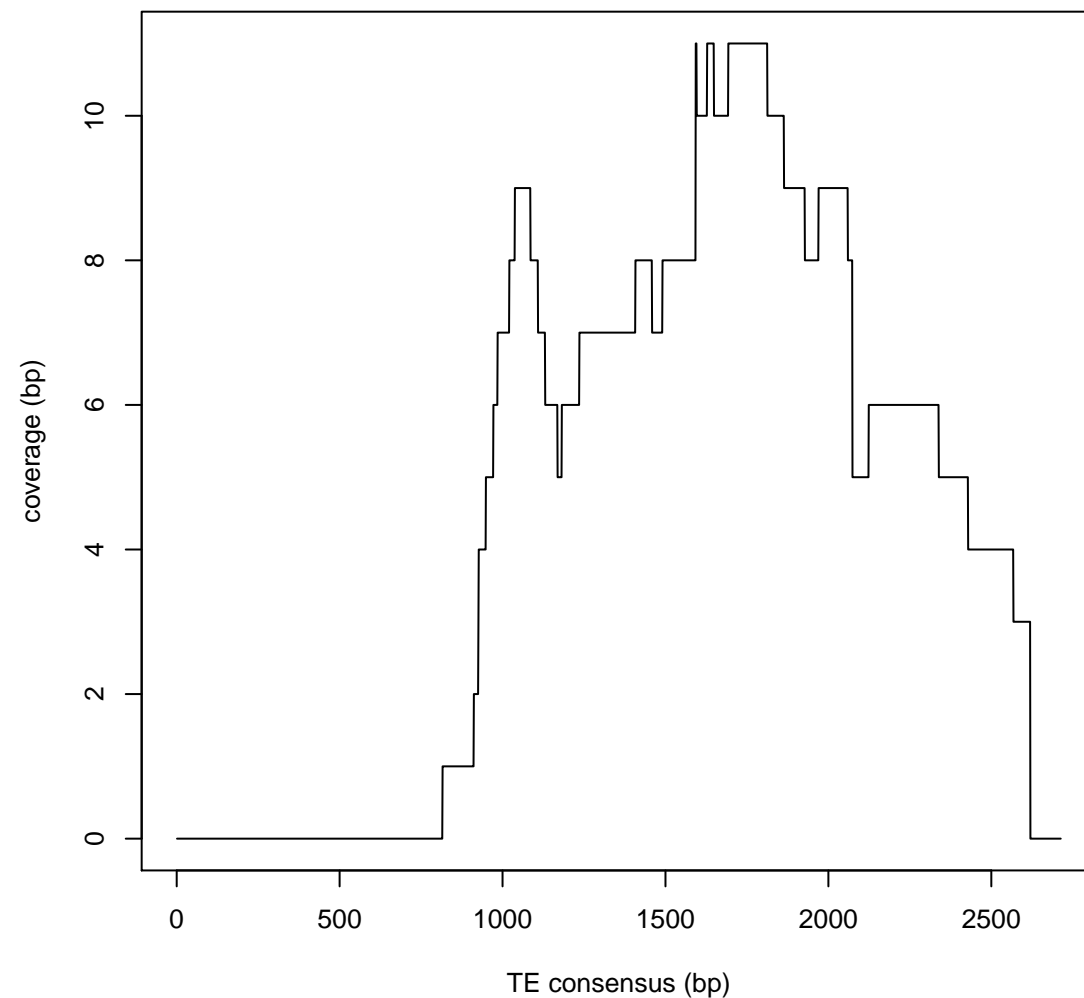


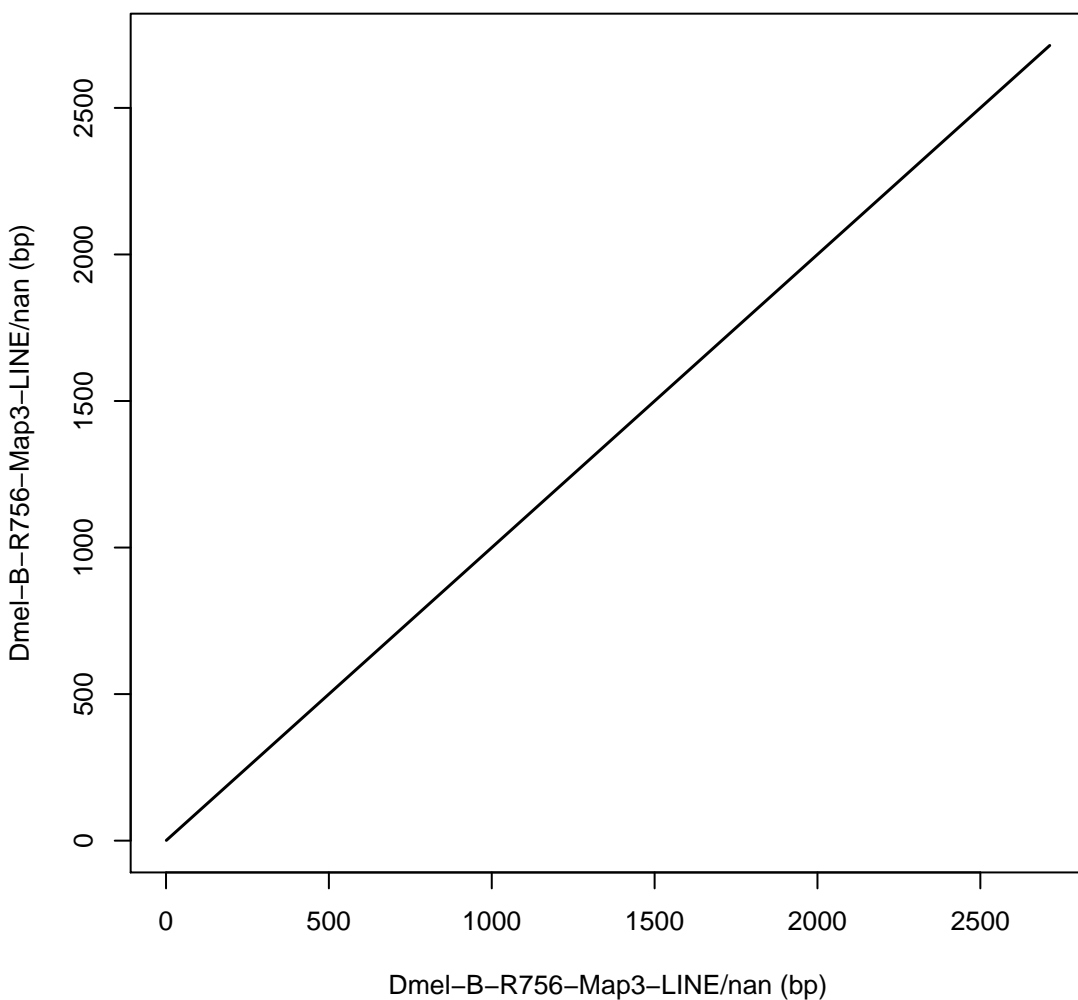
TE: Dmel-B-R756-Map3-LINE/nan
consensus size: 2713bp; fragments: 20; full length: 0 (≥ 2441.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

