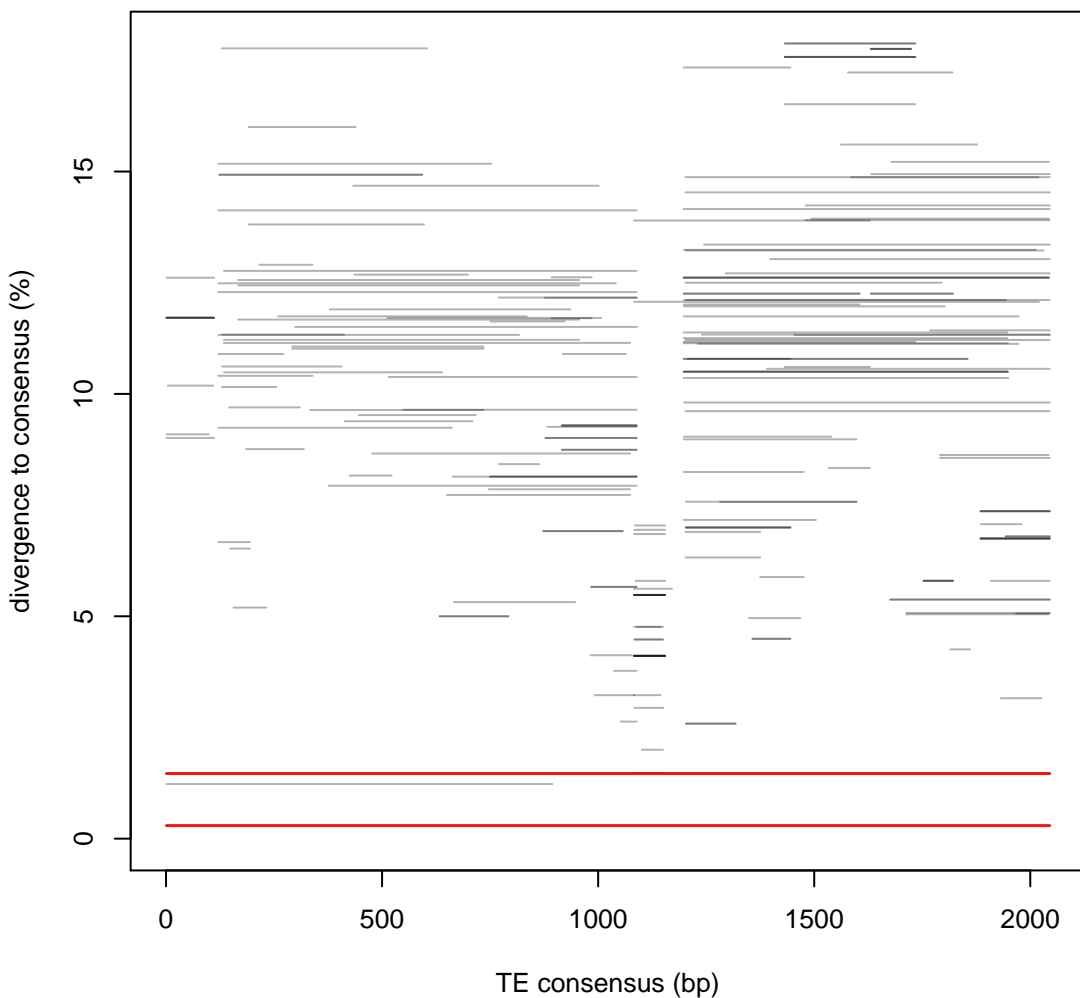
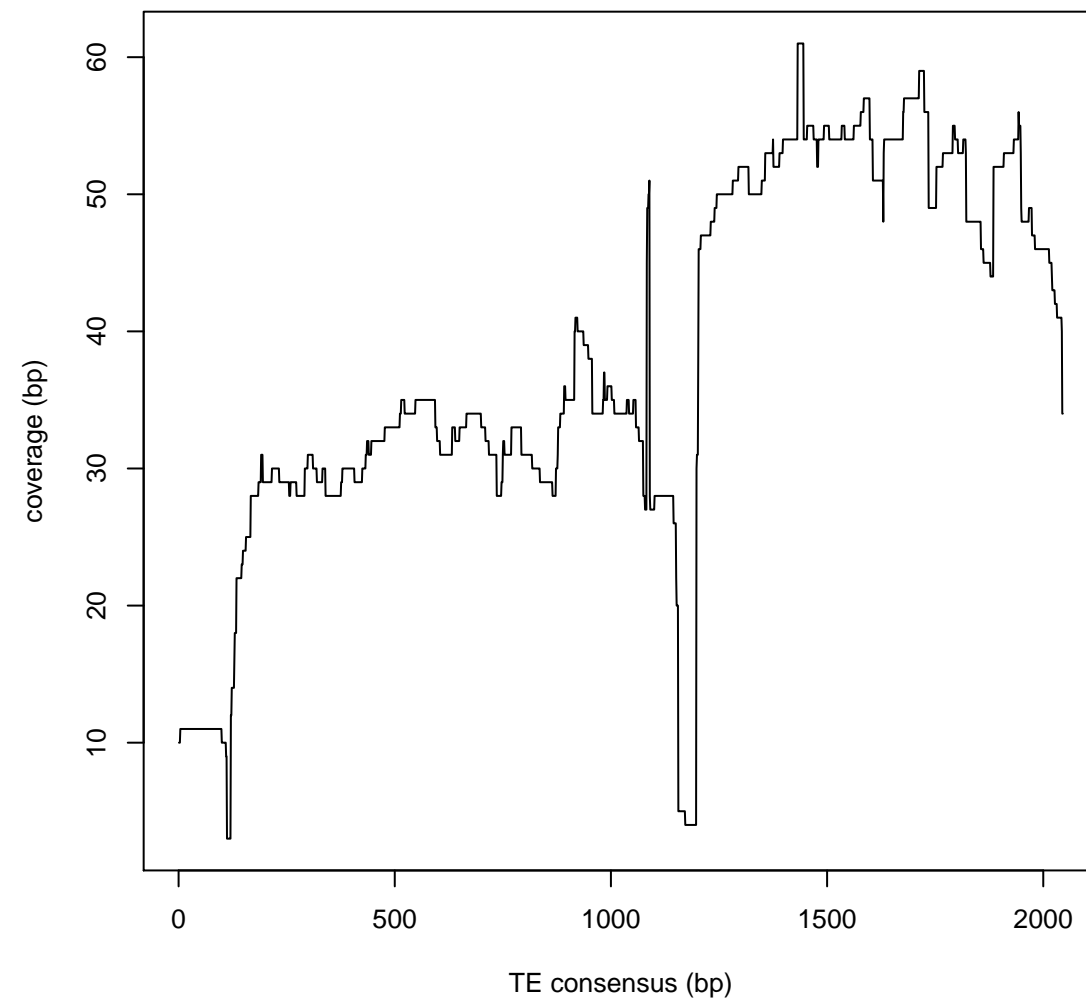


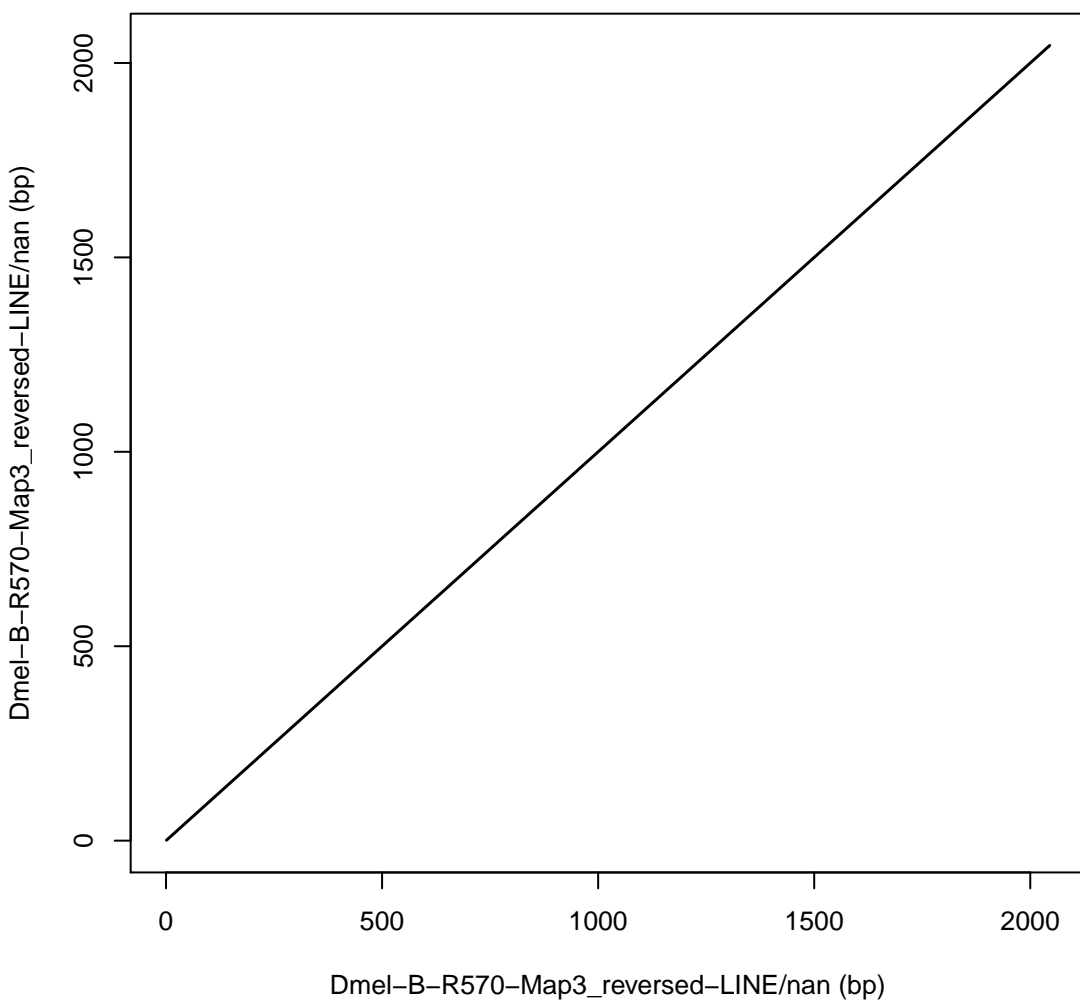
TE: Dmel-B-R570-Map3_reversed-LINE/nan
consensus size: 2045bp; fragments: 213; full length: 2 (≥ 1840.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

