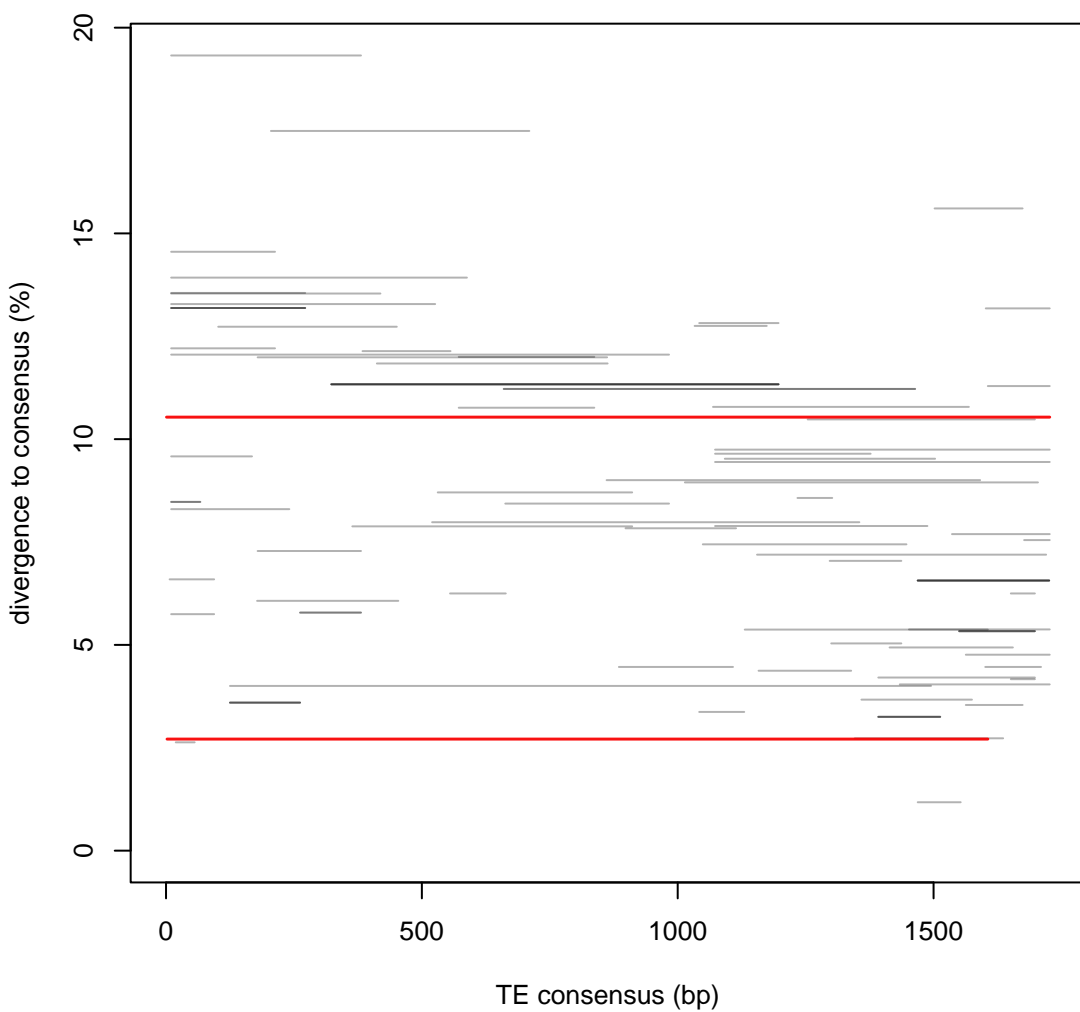
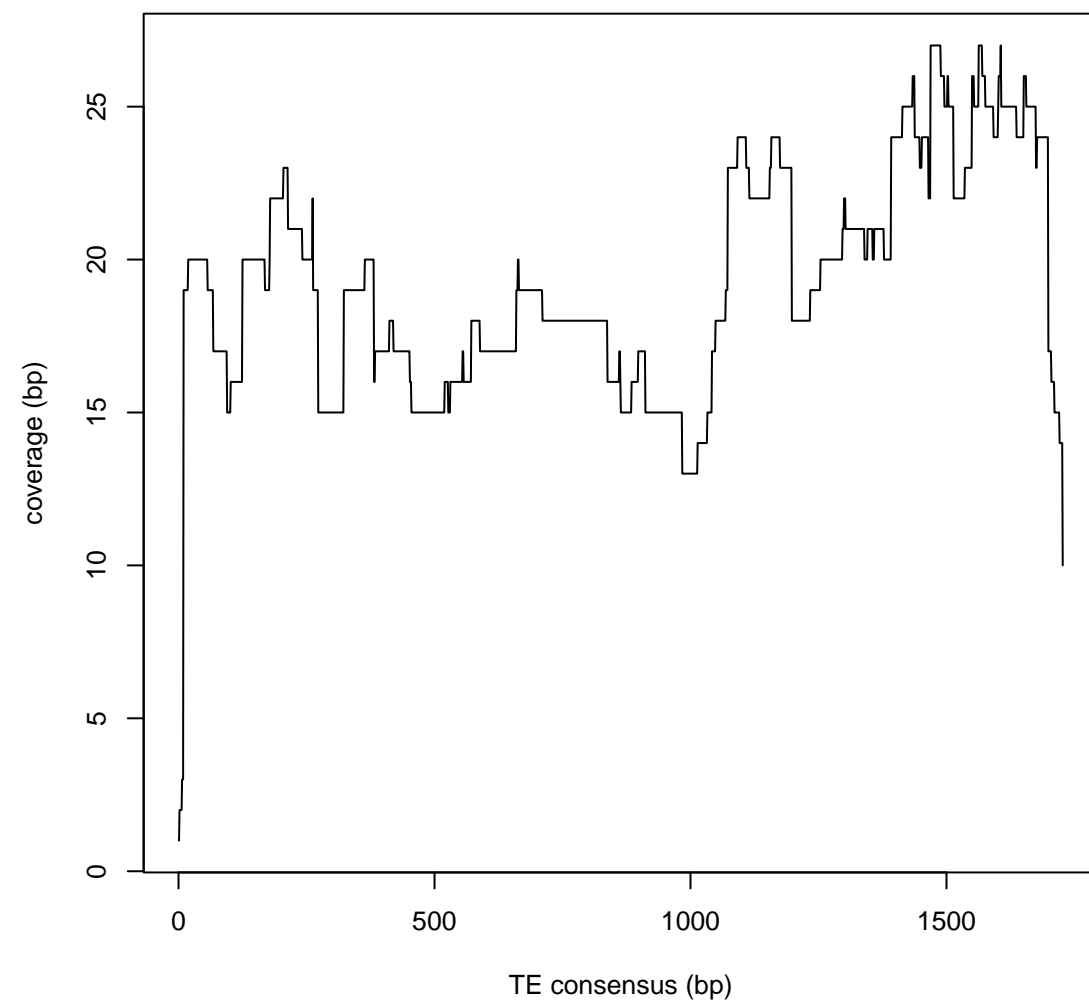


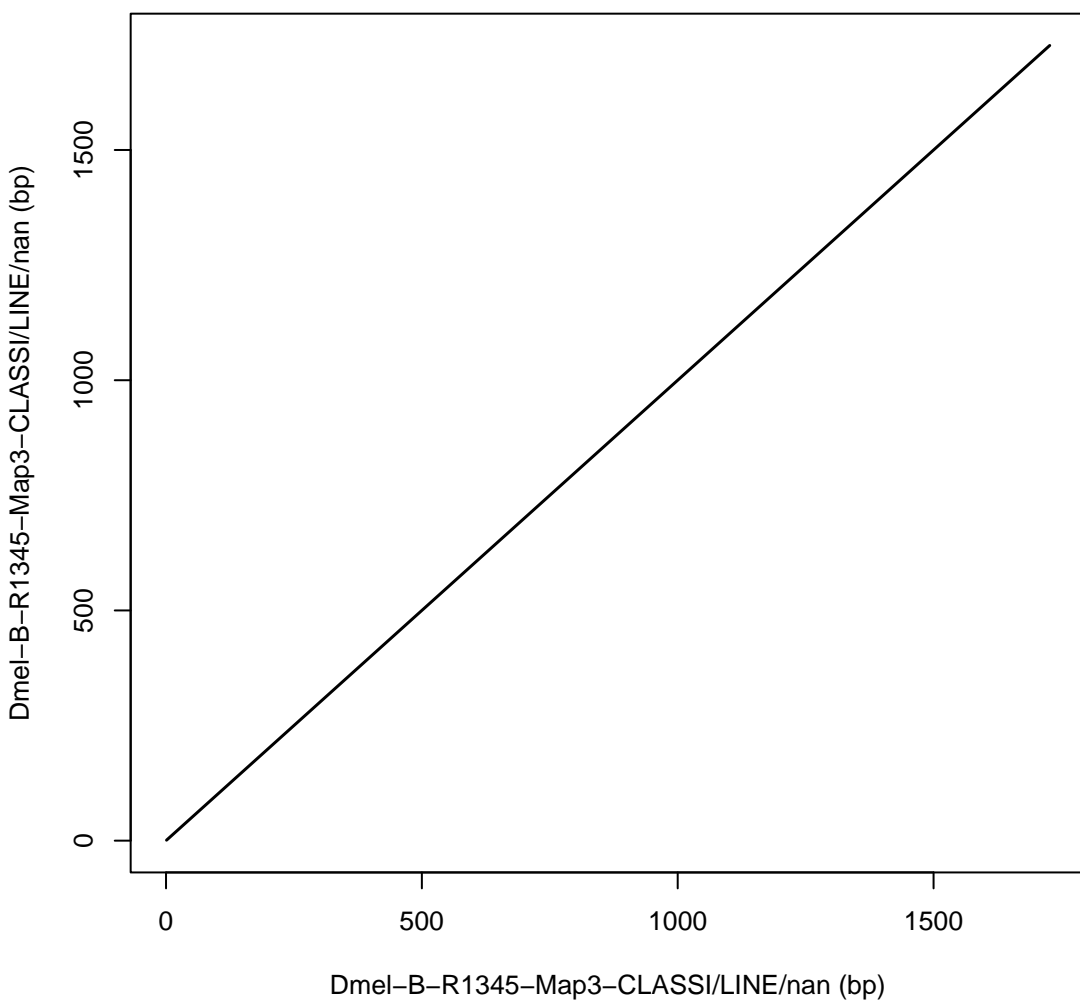
TE: Dmel-B-R1345-Map3-CLASSI/LINE/nan
consensus size: 1727bp; fragments: 94; full length: 2 (≥ 1554.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

