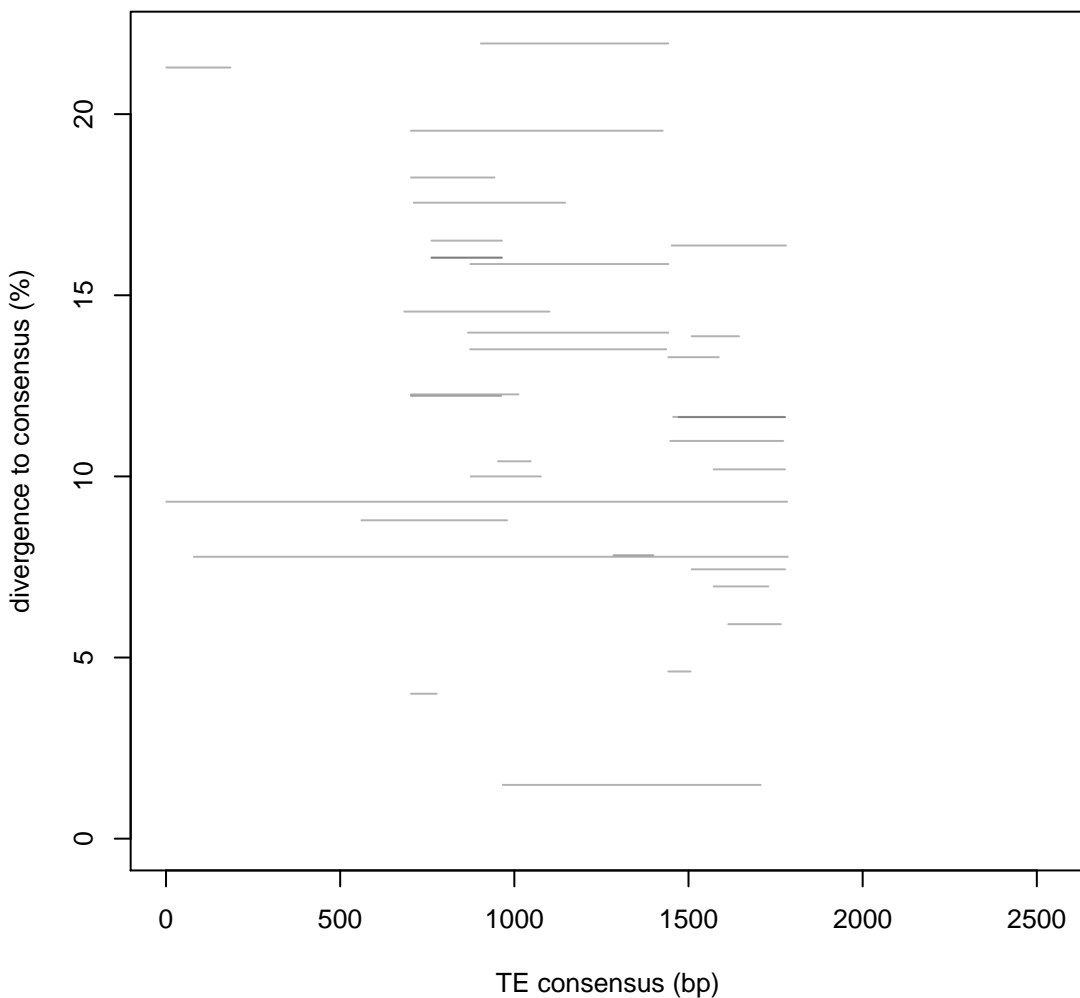
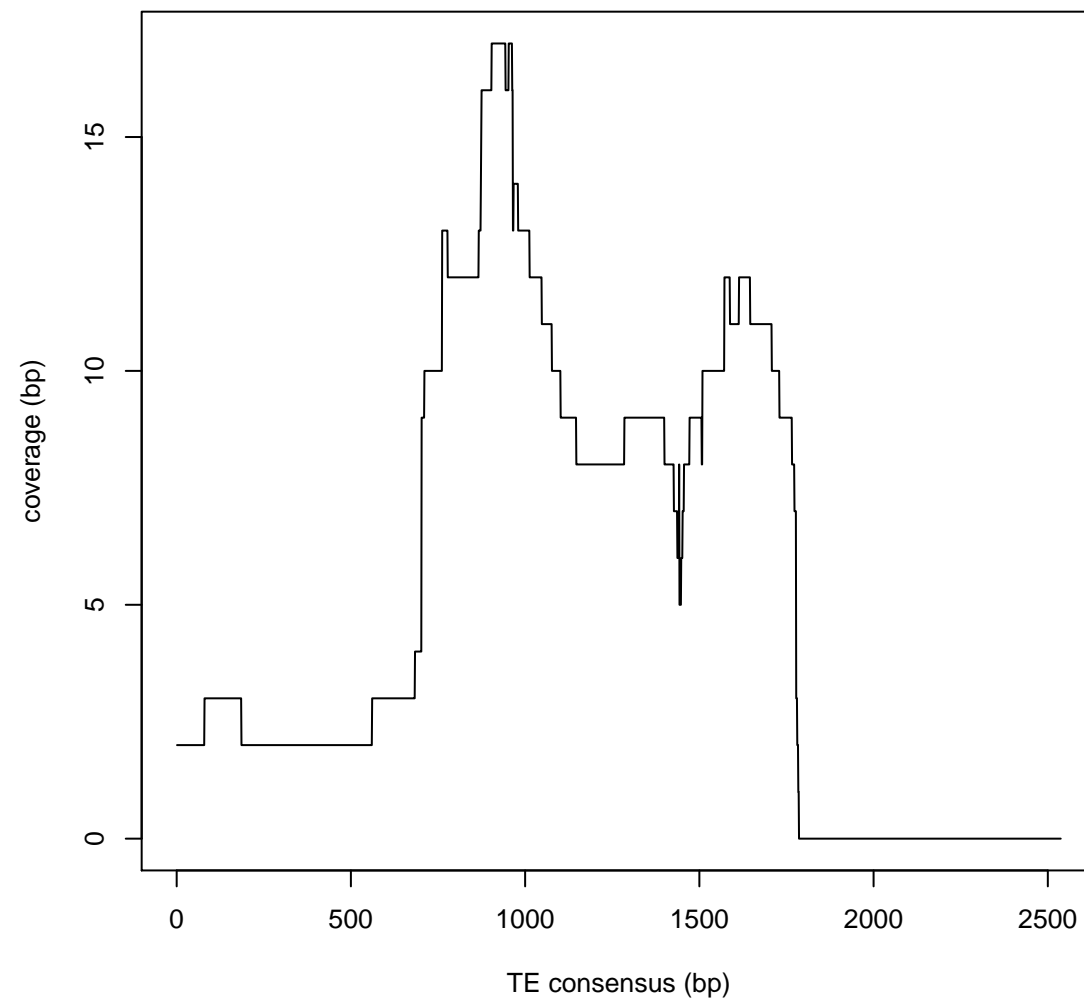


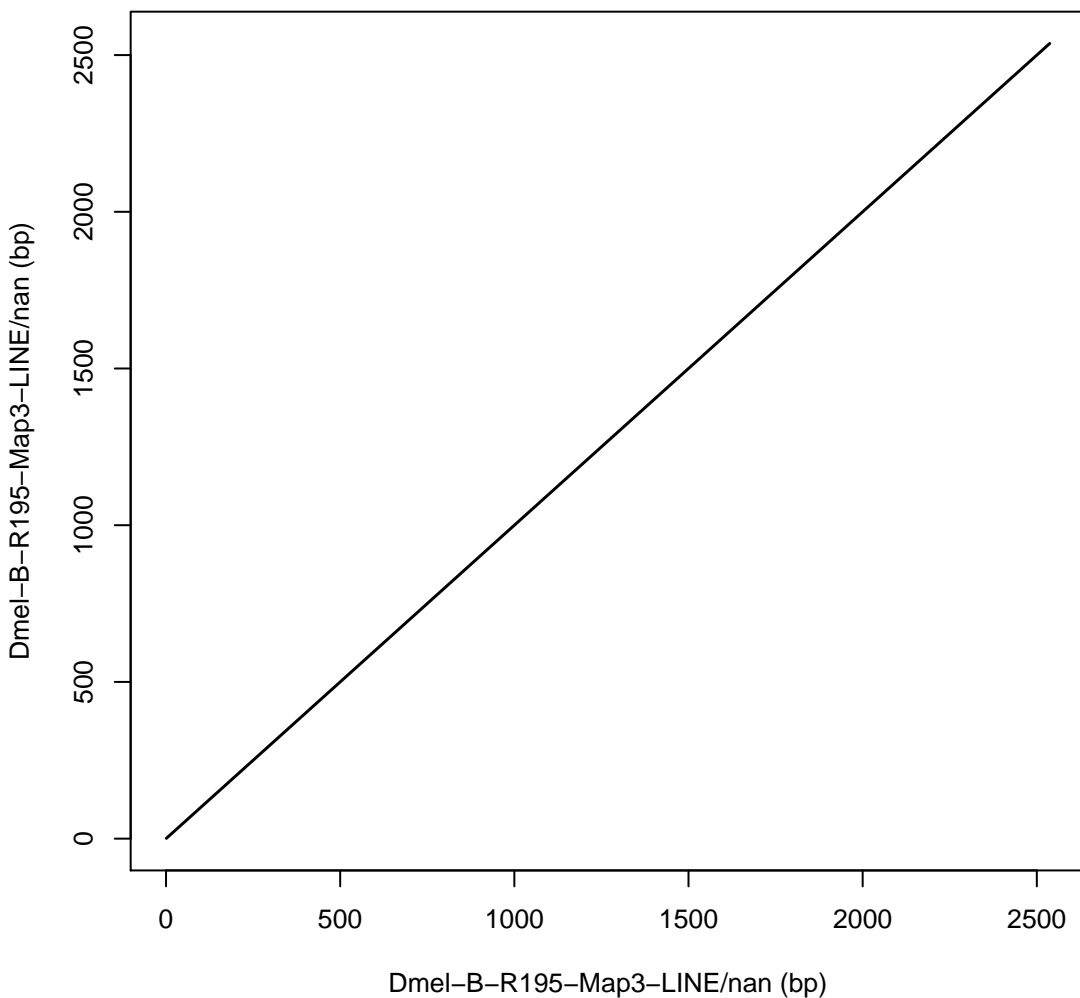
TE: Dmel-B-R195-Map3-LINE/nan
consensus size: 2537bp; fragments: 33; full length: 0 (>=2283.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

