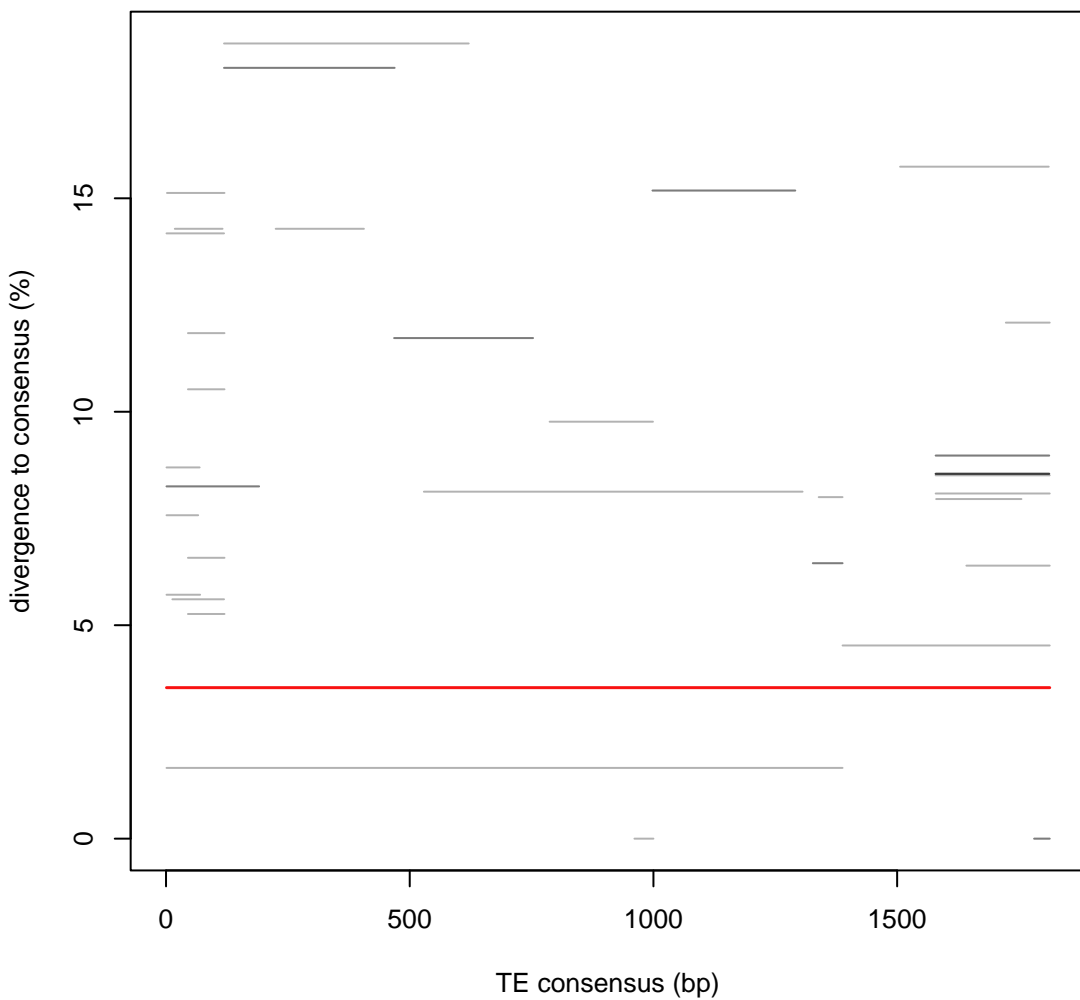
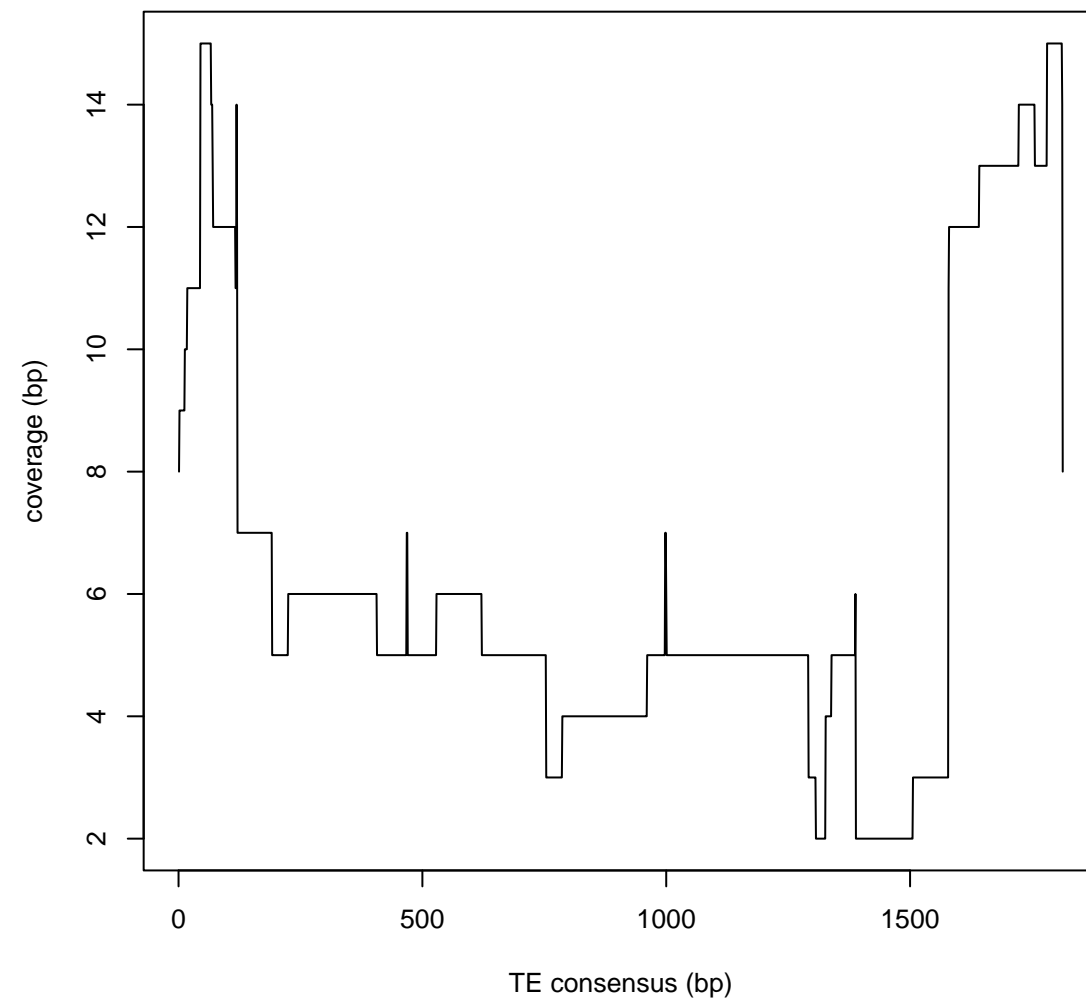


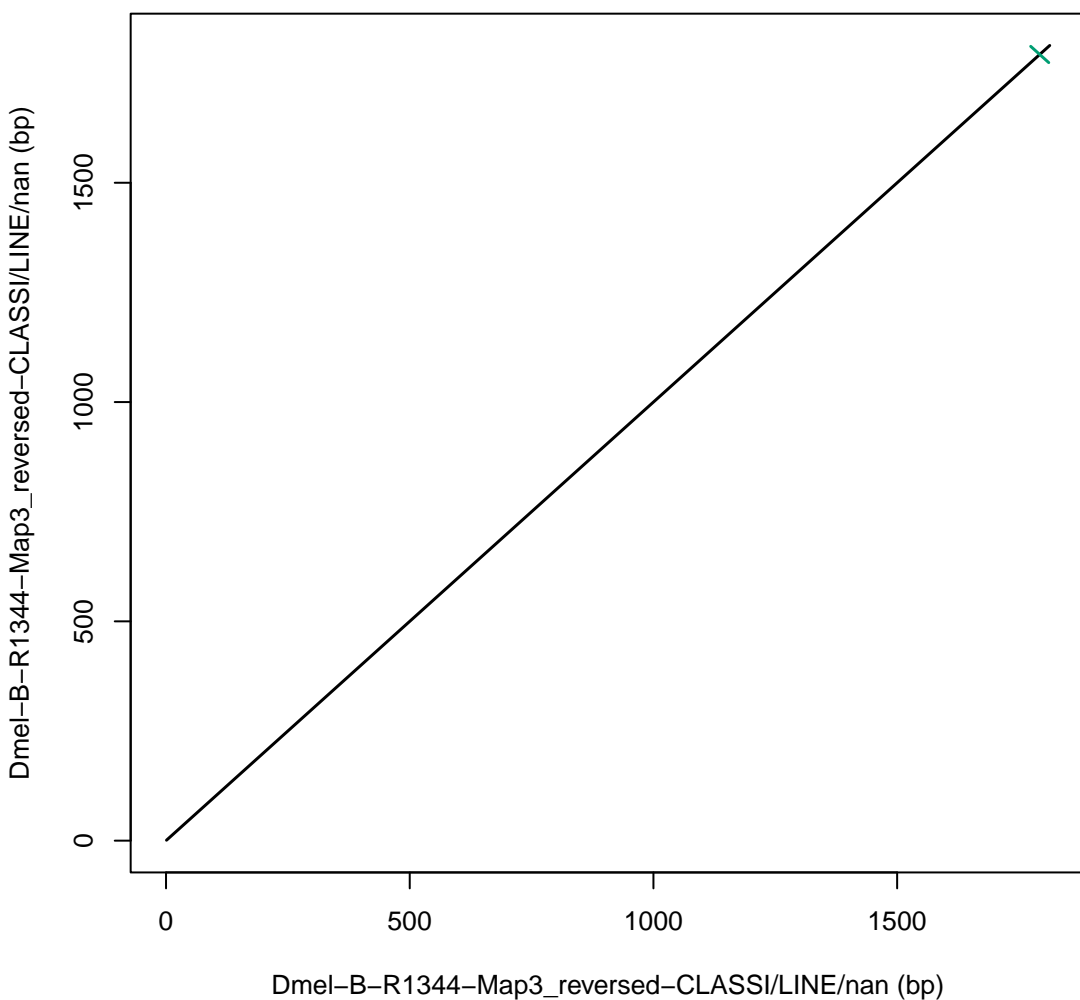
TE: Dmel-B-R1344-Map3_reversed-CLASSI/LINE/nan
consensus size: 1813bp; fragments: 44; full length: 1 (>=1631.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

