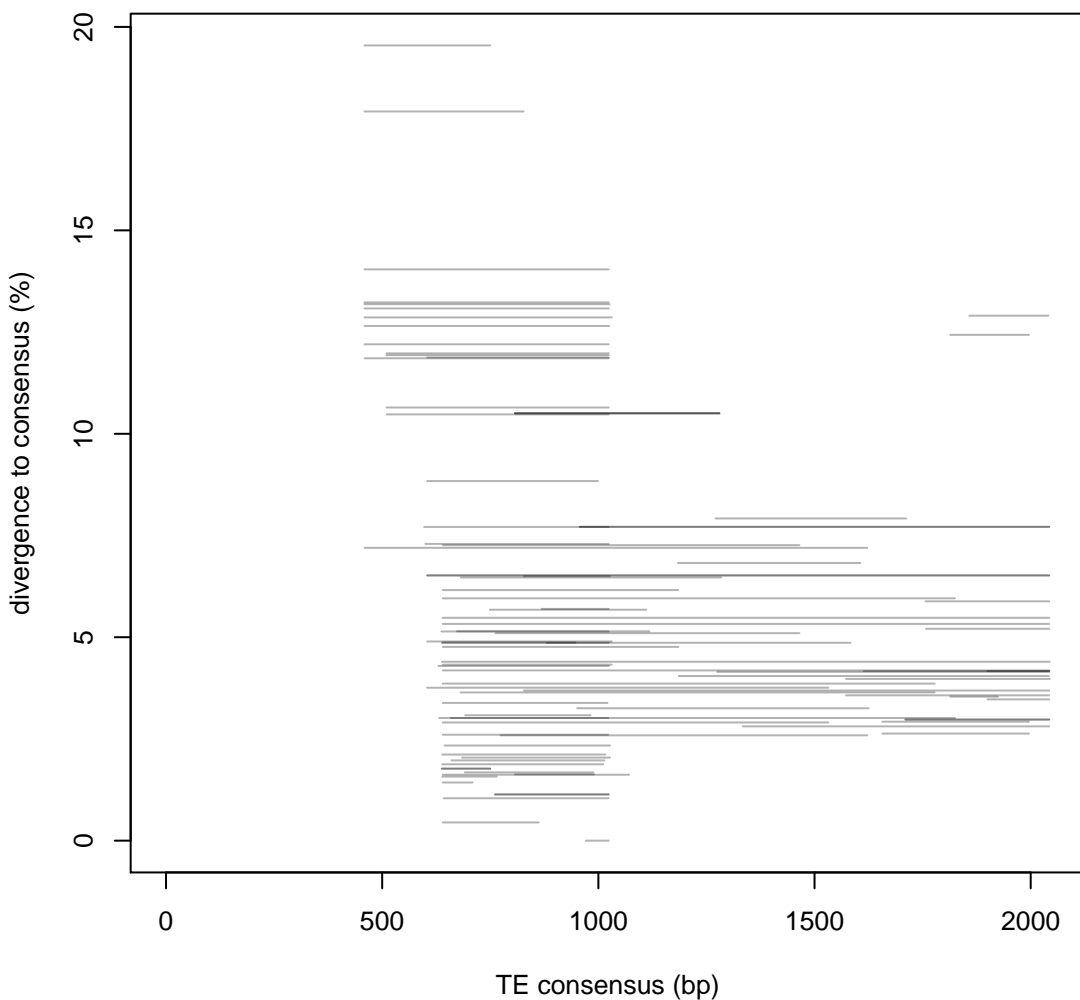
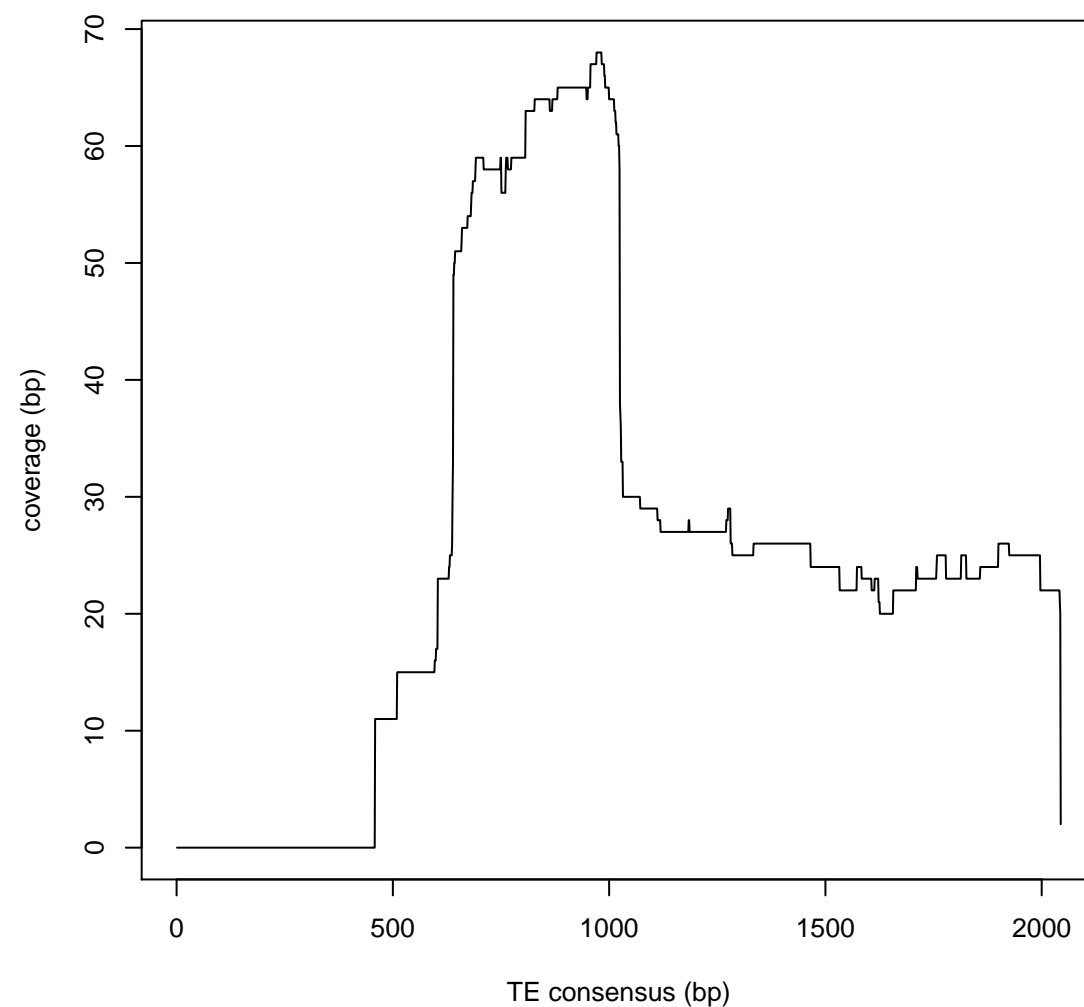


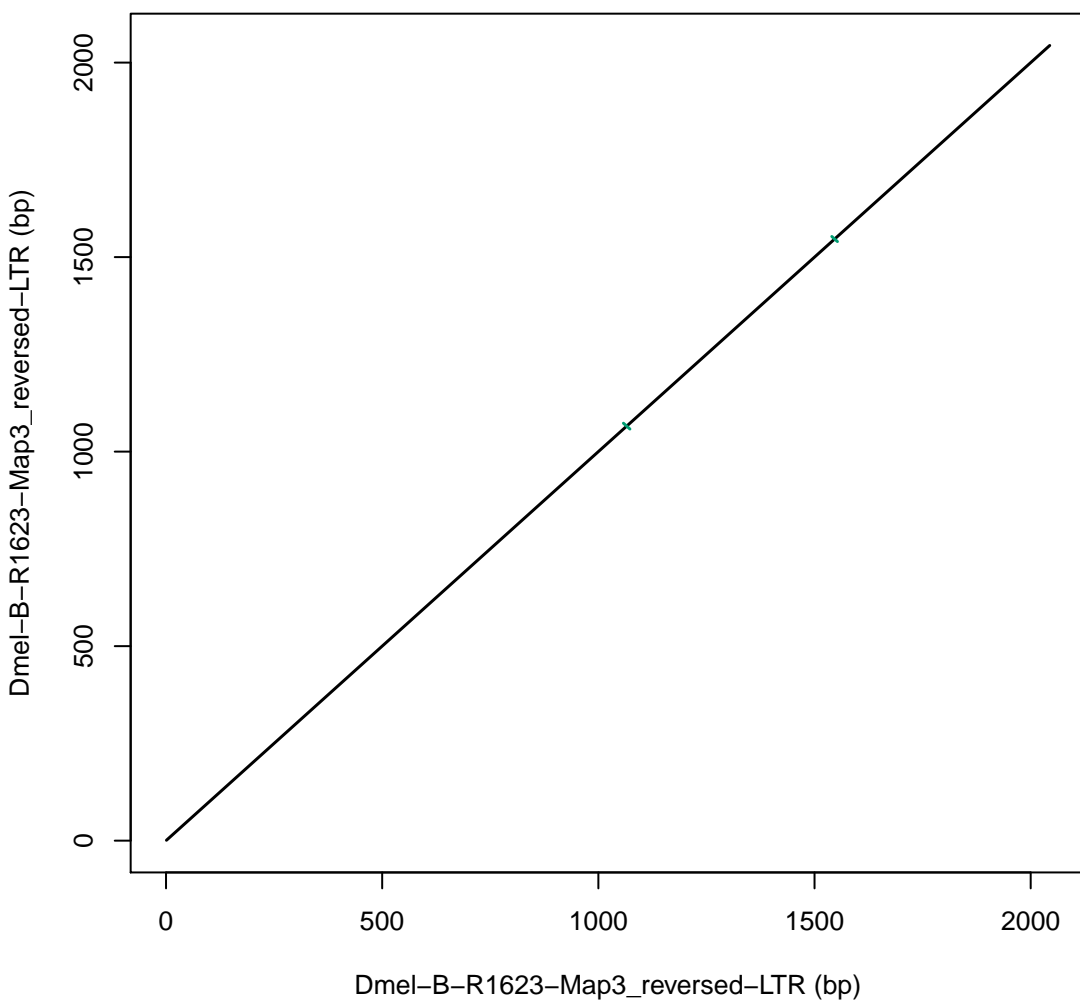
TE: Dmel-B-R1623-Map3_reversed-LTR
consensus size: 2044bp; fragments: 95; full length: 0 (>=1839.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

