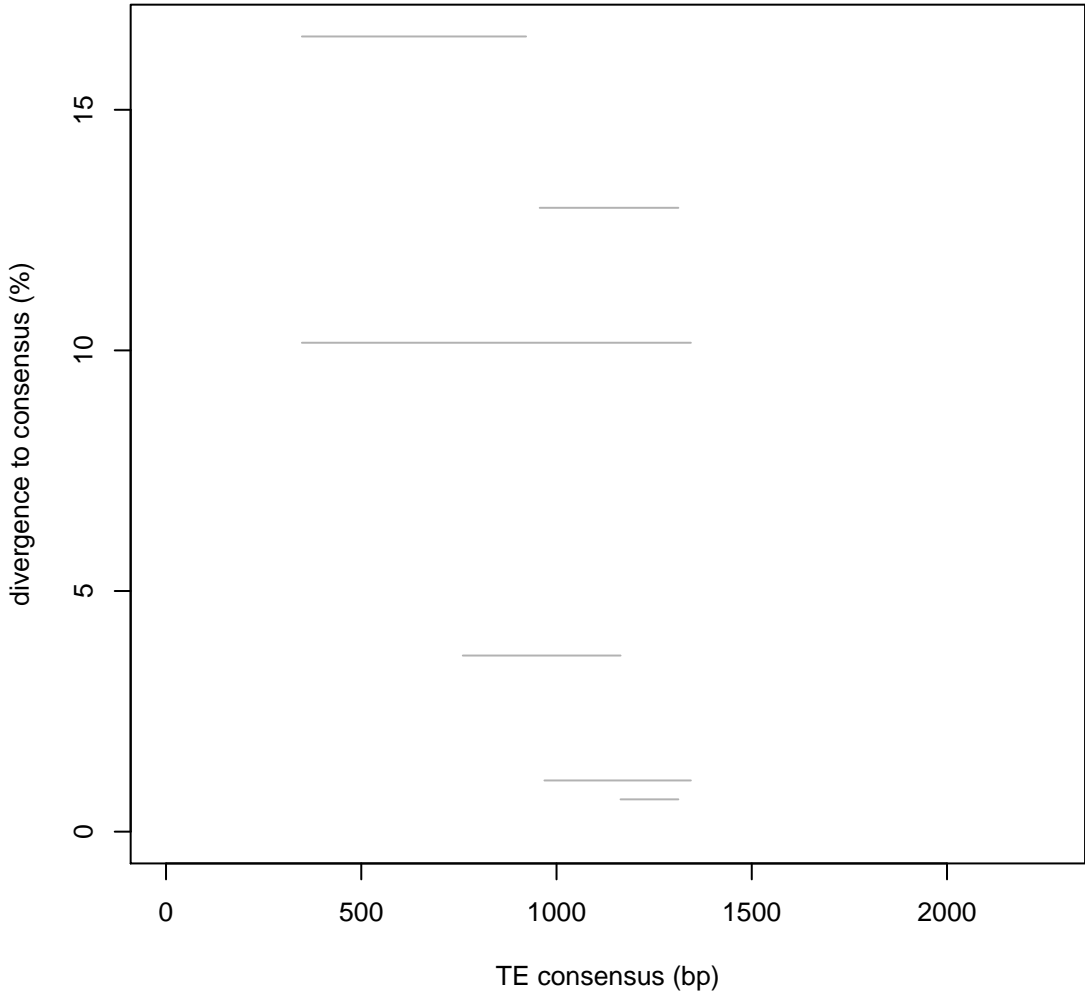
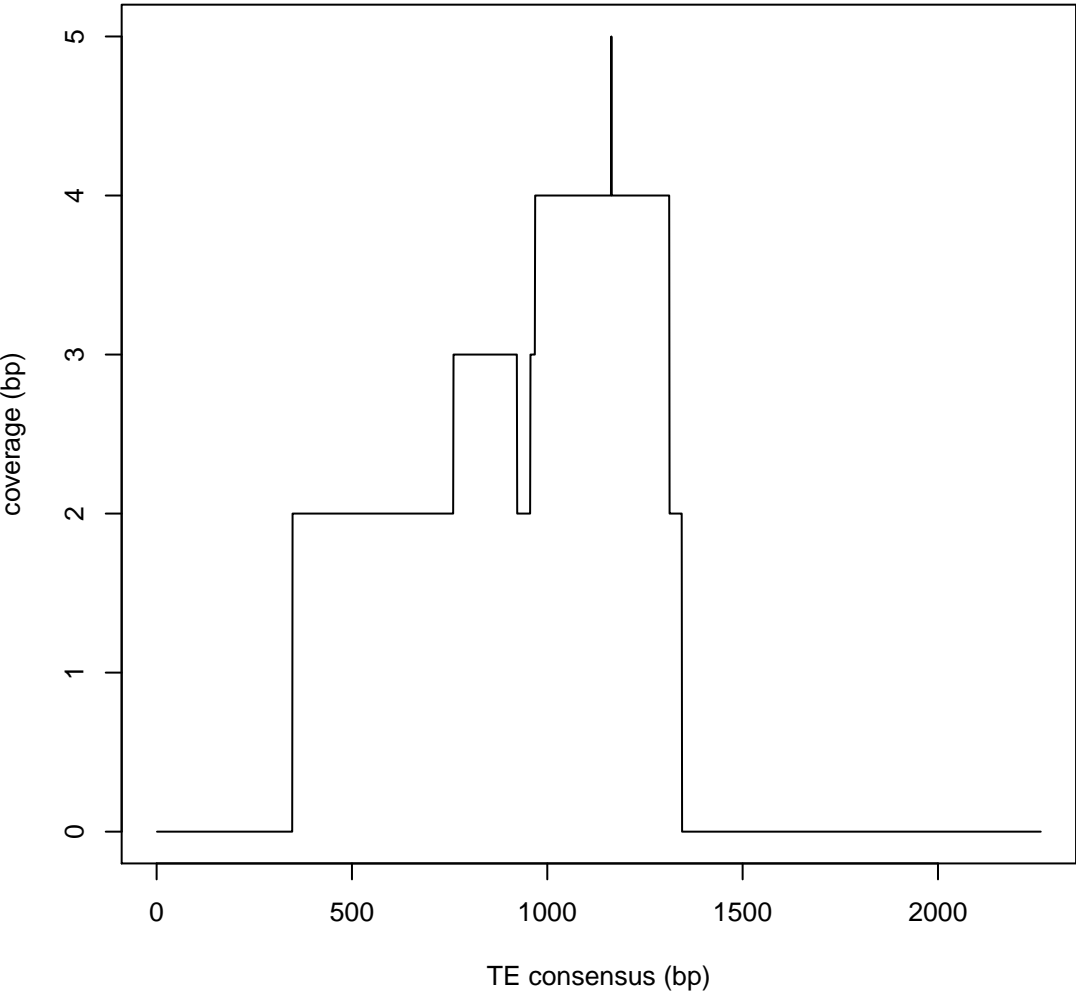


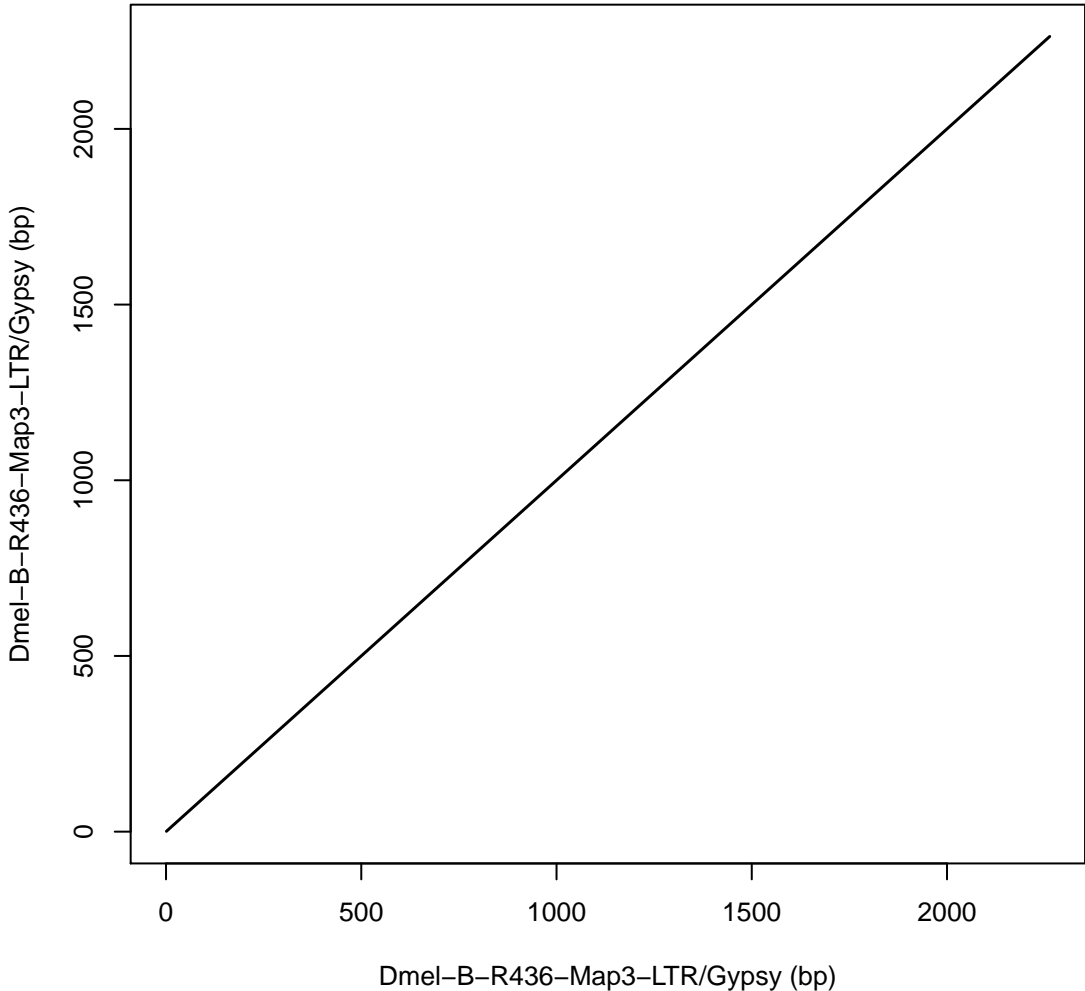
TE: Dmel-B-R436-Map3-LTR/Gypsy
consensus size: 2263bp; fragments: 6; full length: 0 (>=2036.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

