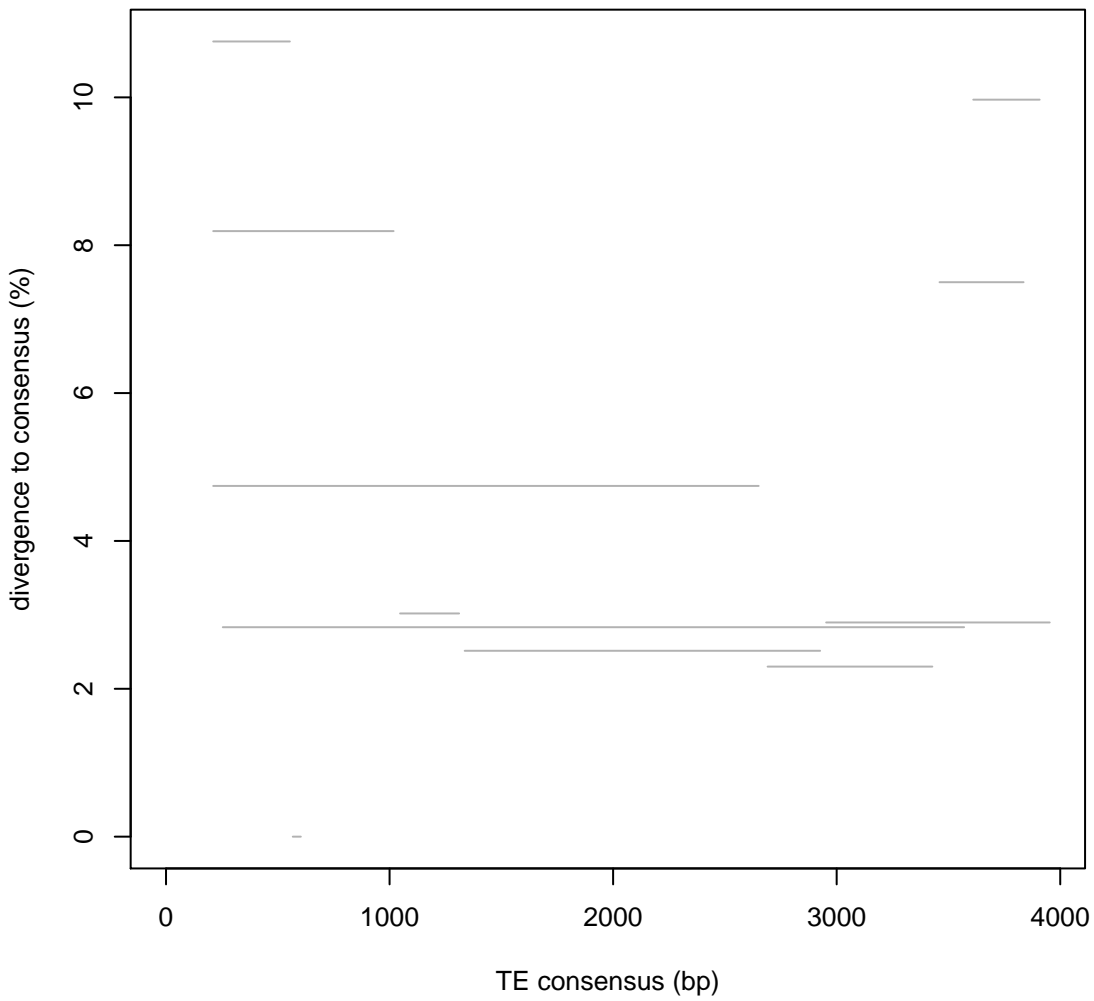
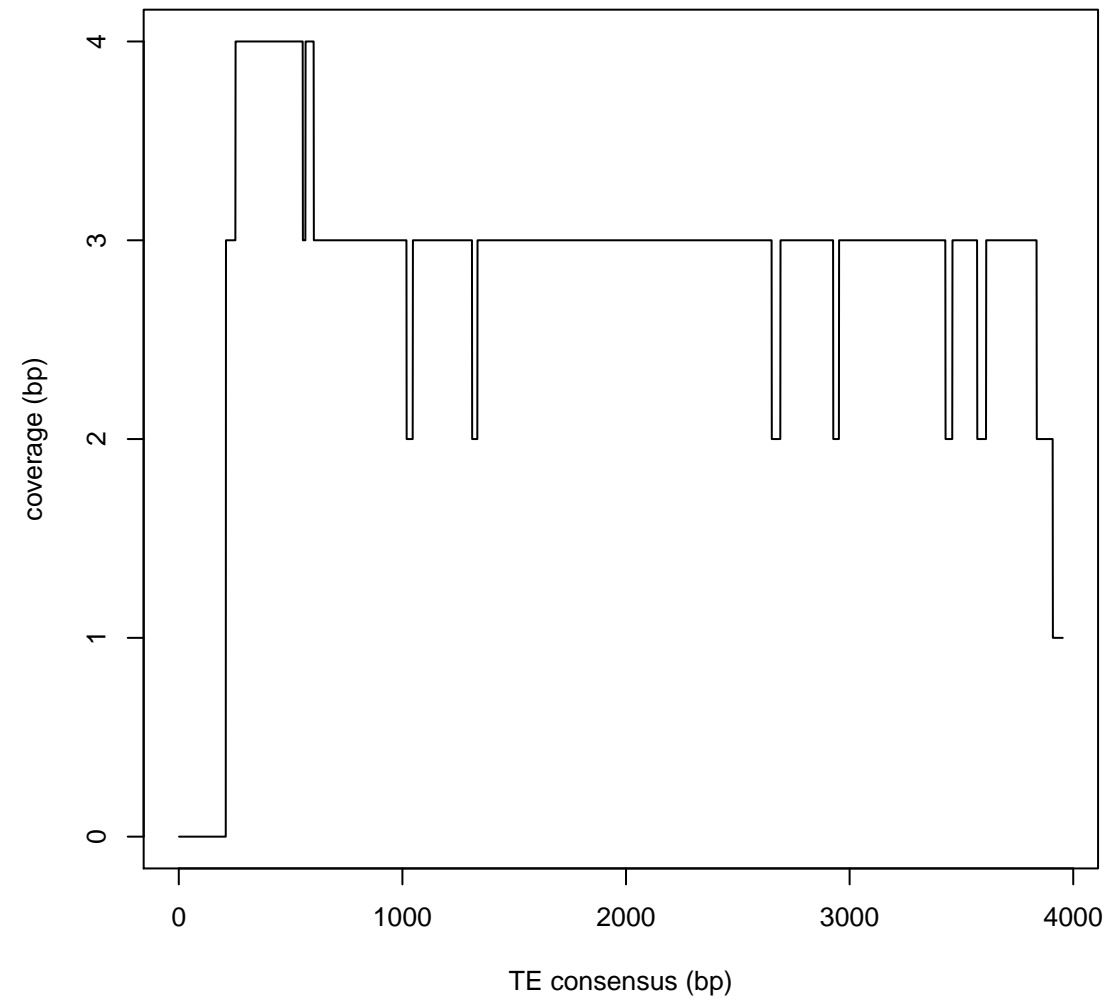


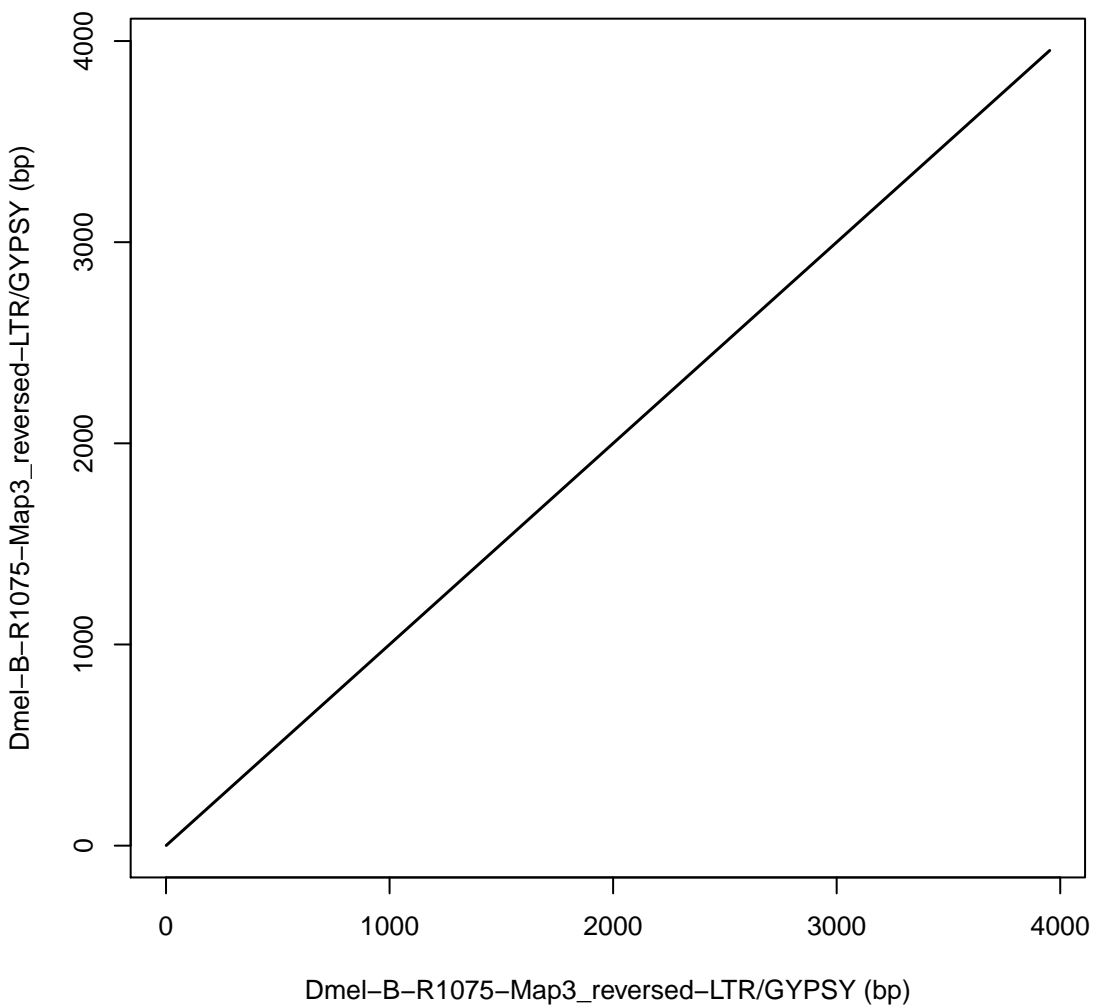
TE: Dmel-B-R1075-Map3_reversed-LTR/Gypsy
consensus size: 3953bp; fragments: 11; full length: 0 (>=3557.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

