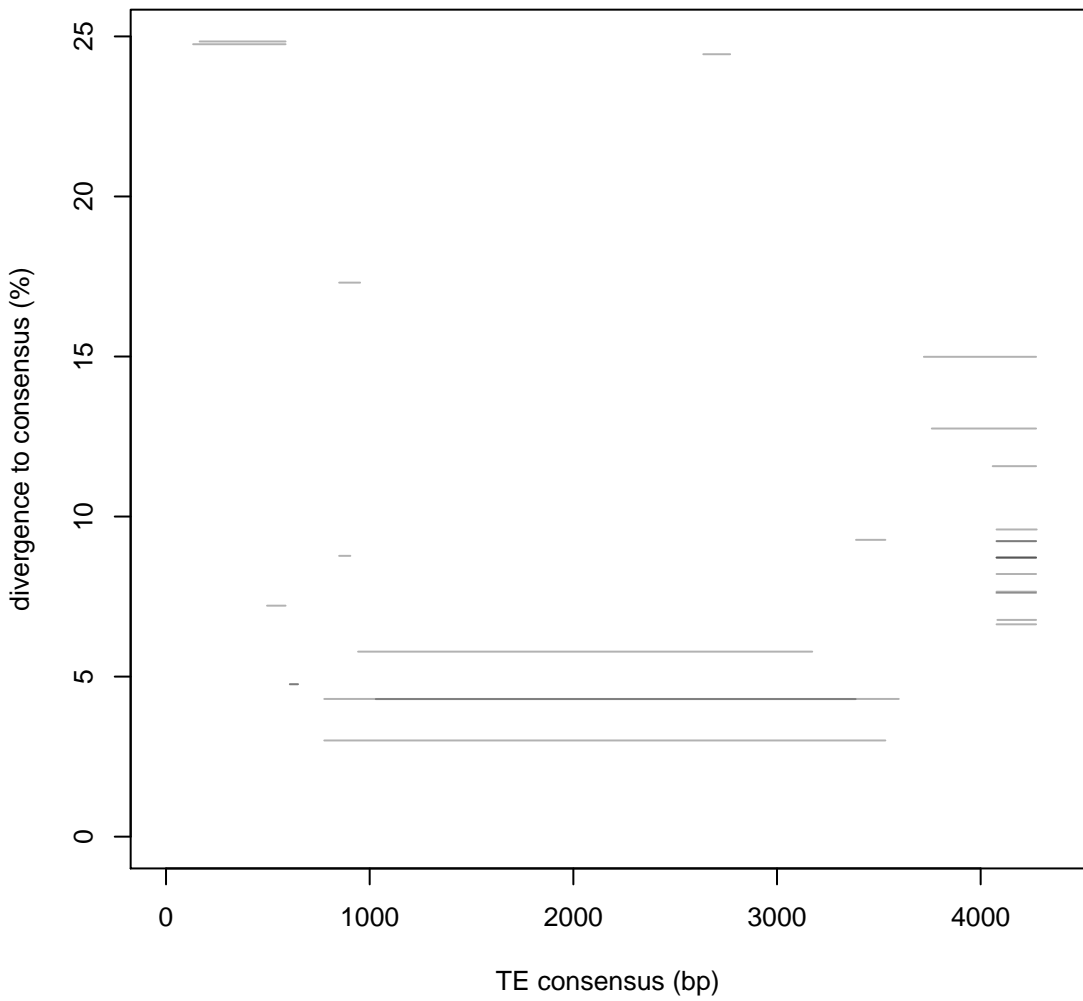
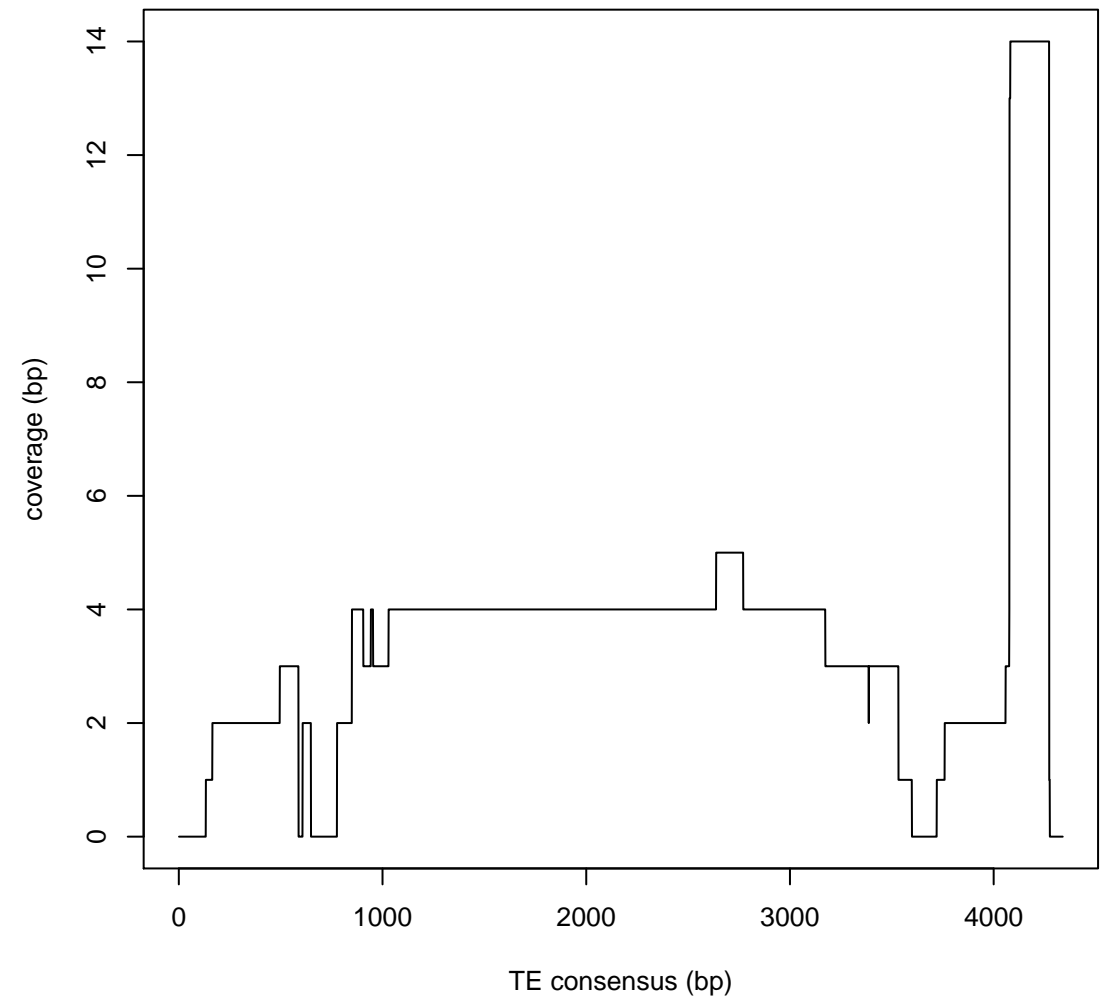


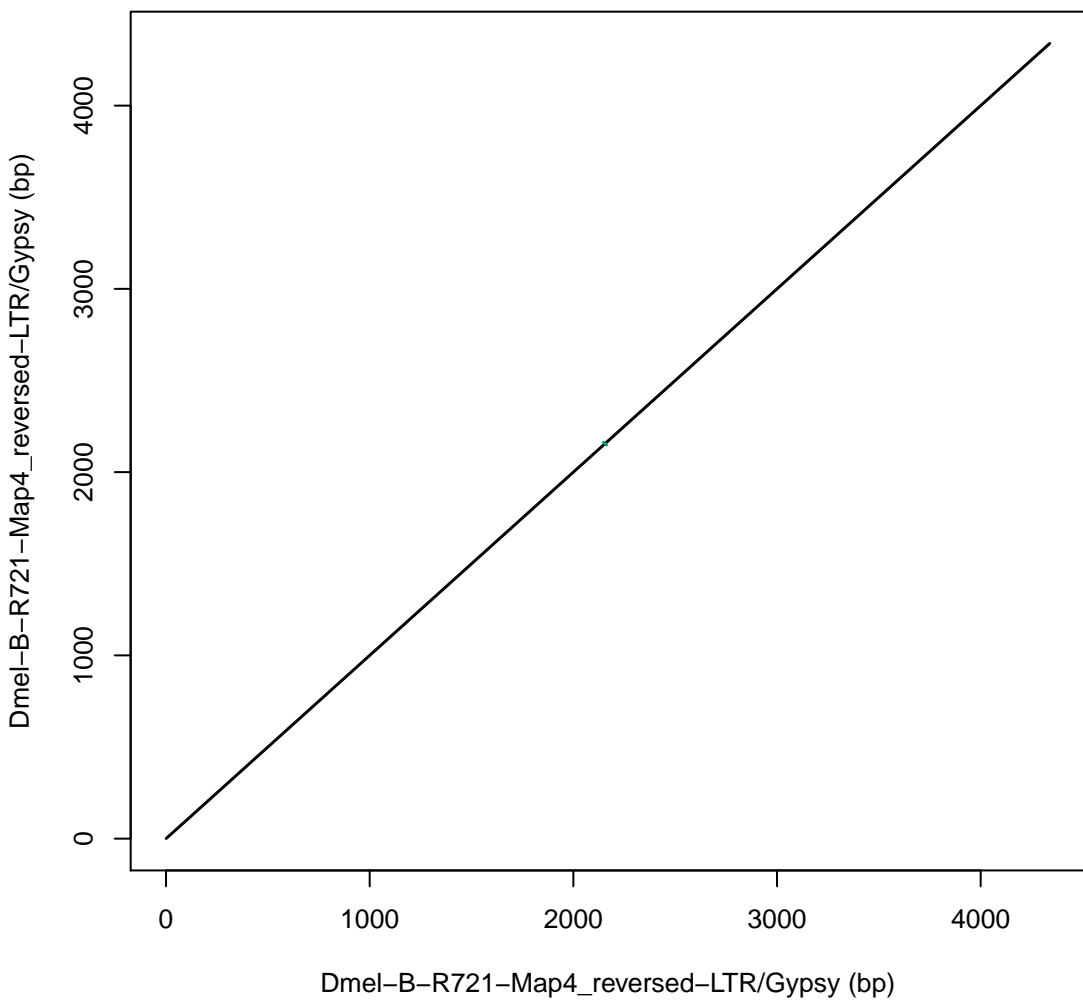
TE: Dmel-B-R721-Map4_reversed-LTR/Gypsy
consensus size: 4339bp; fragments: 27; full length: 0 (>=3905.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

