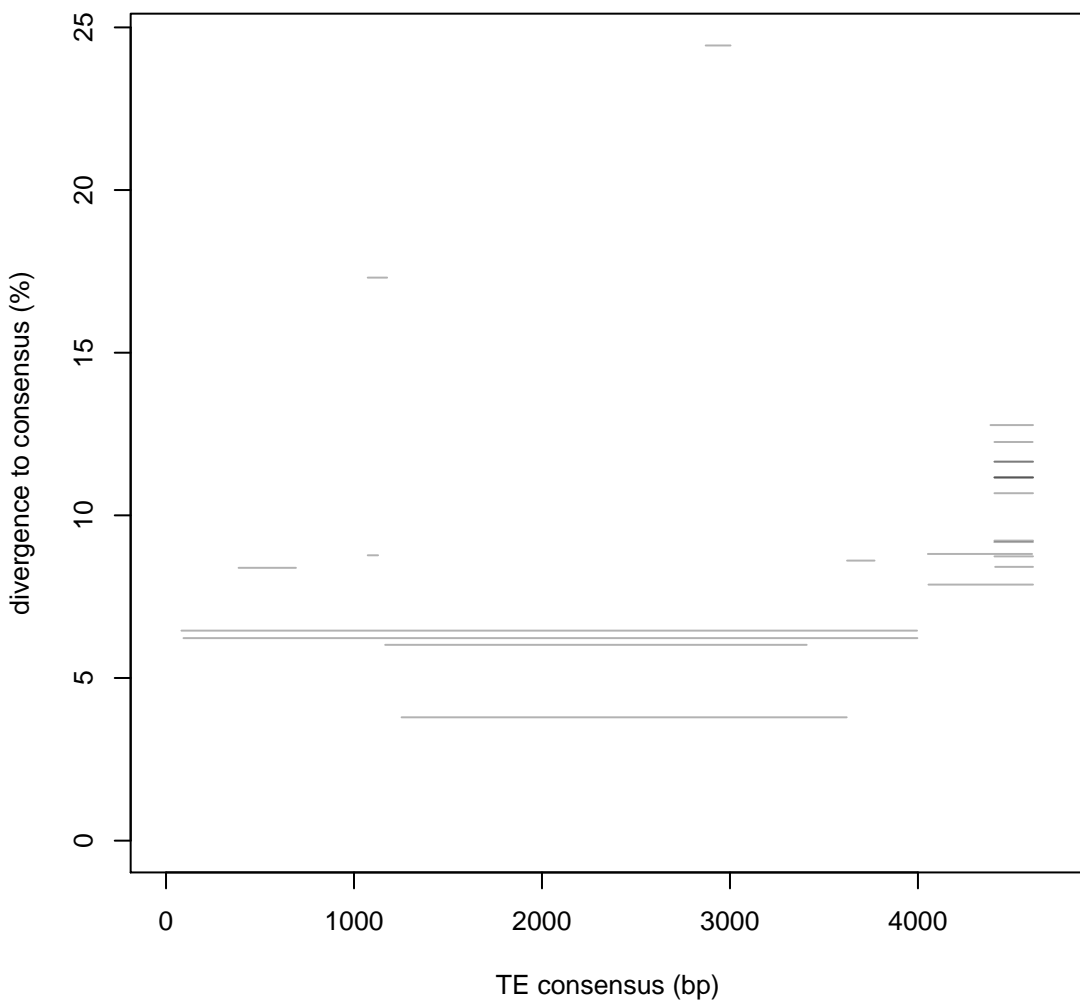
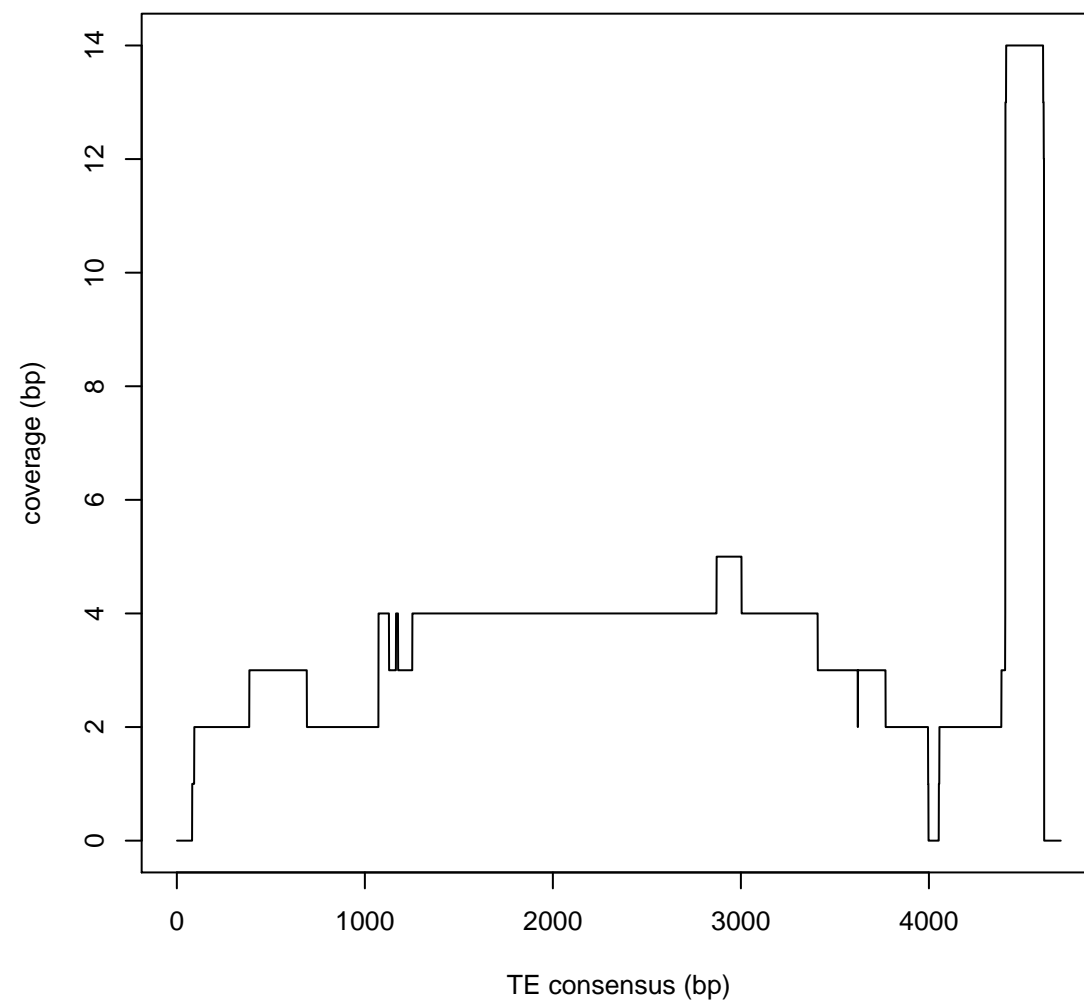


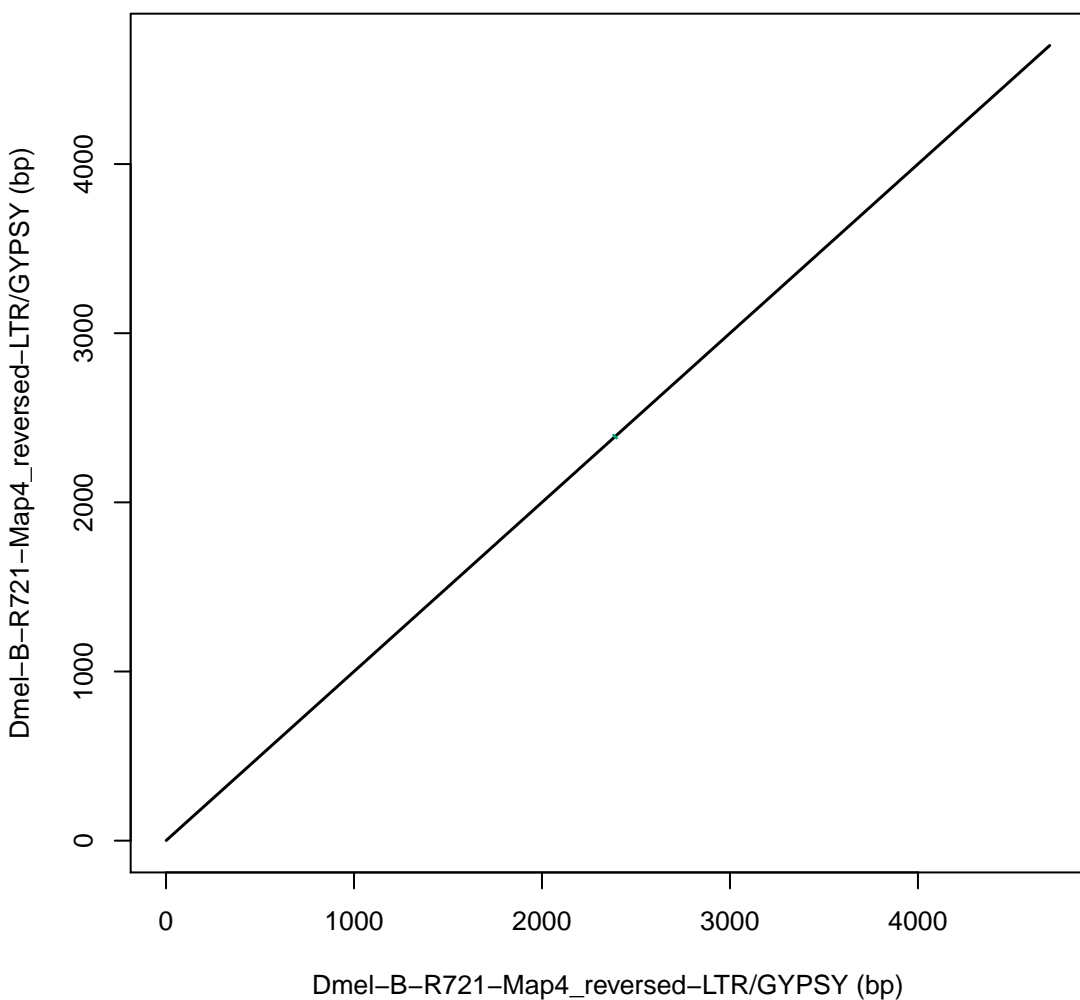
TE: Dmel-B-R721-Map4_reversed-LTR/Gypsy
consensus size: 4701bp; fragments: 23; full length: 0 (>=4230.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

