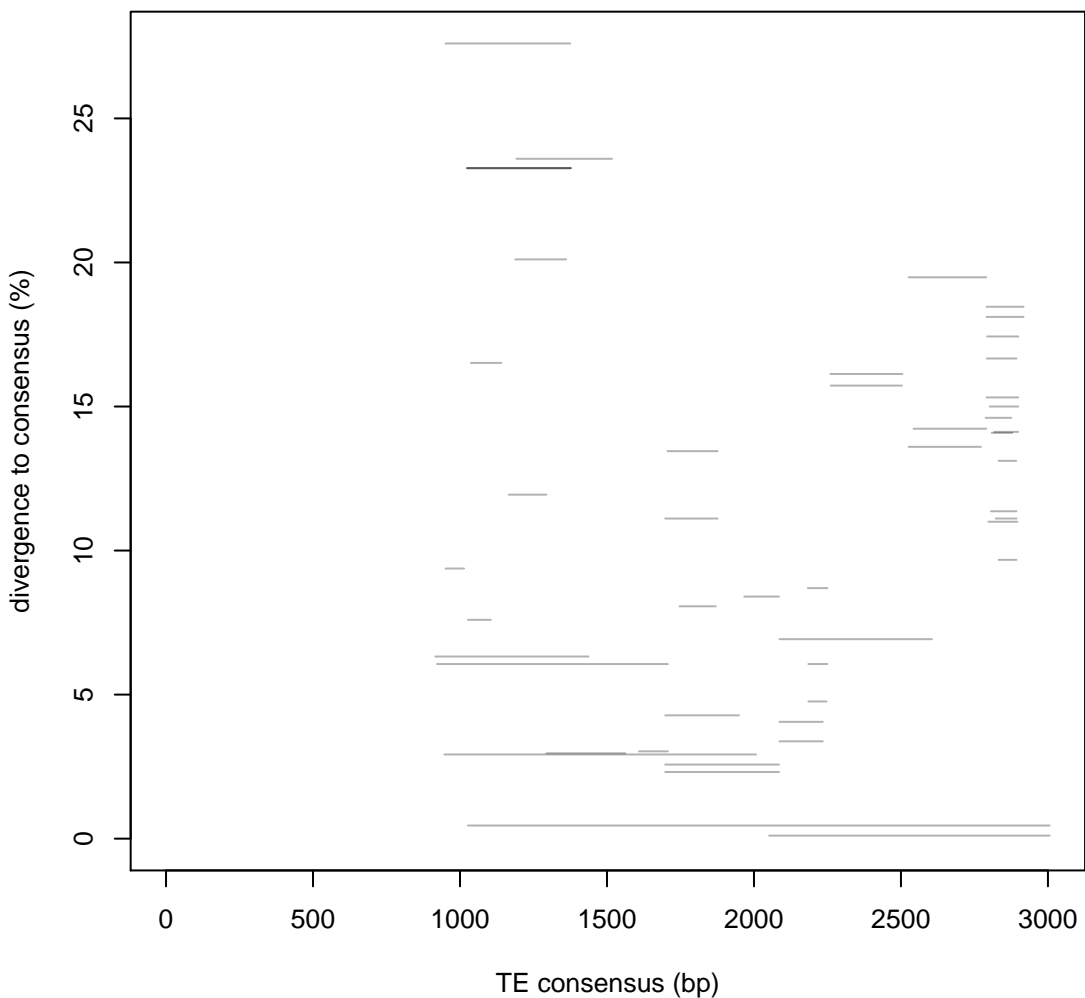
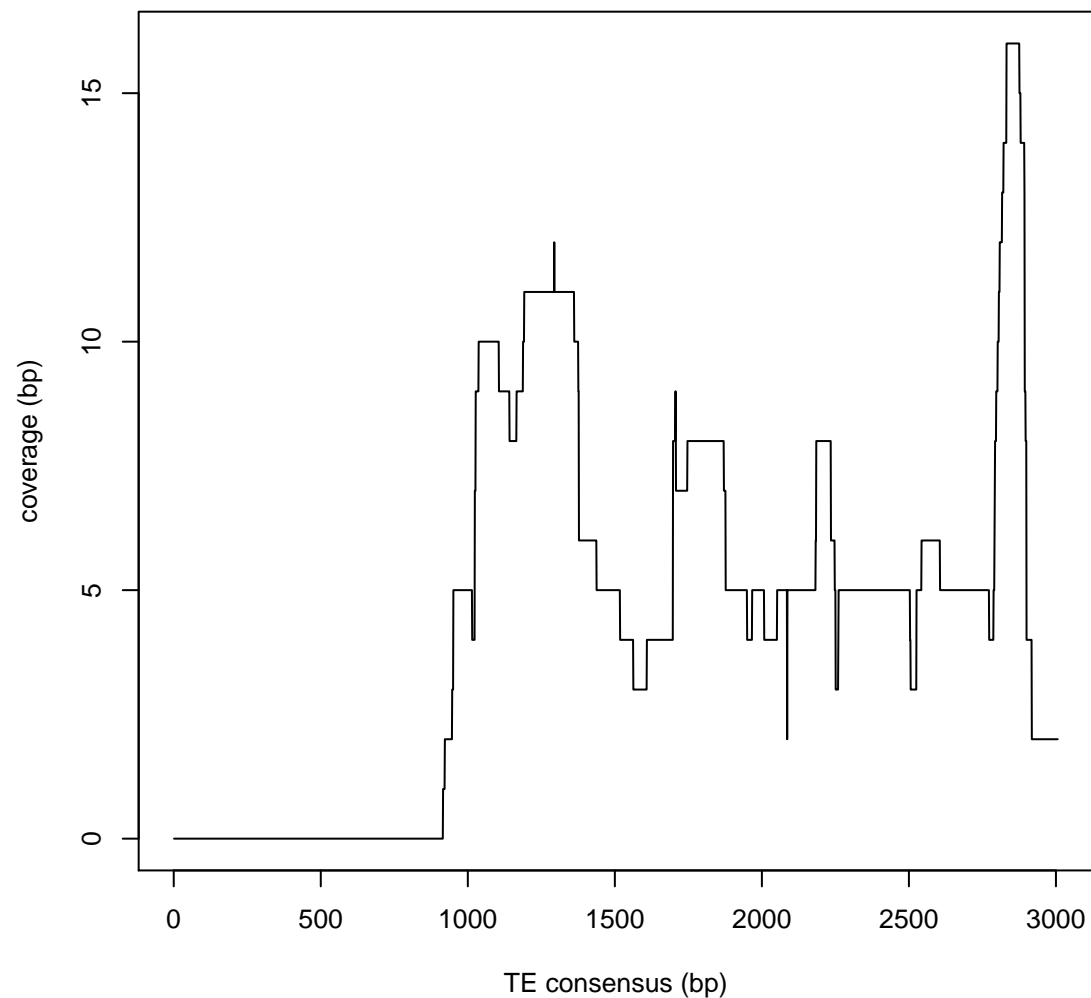


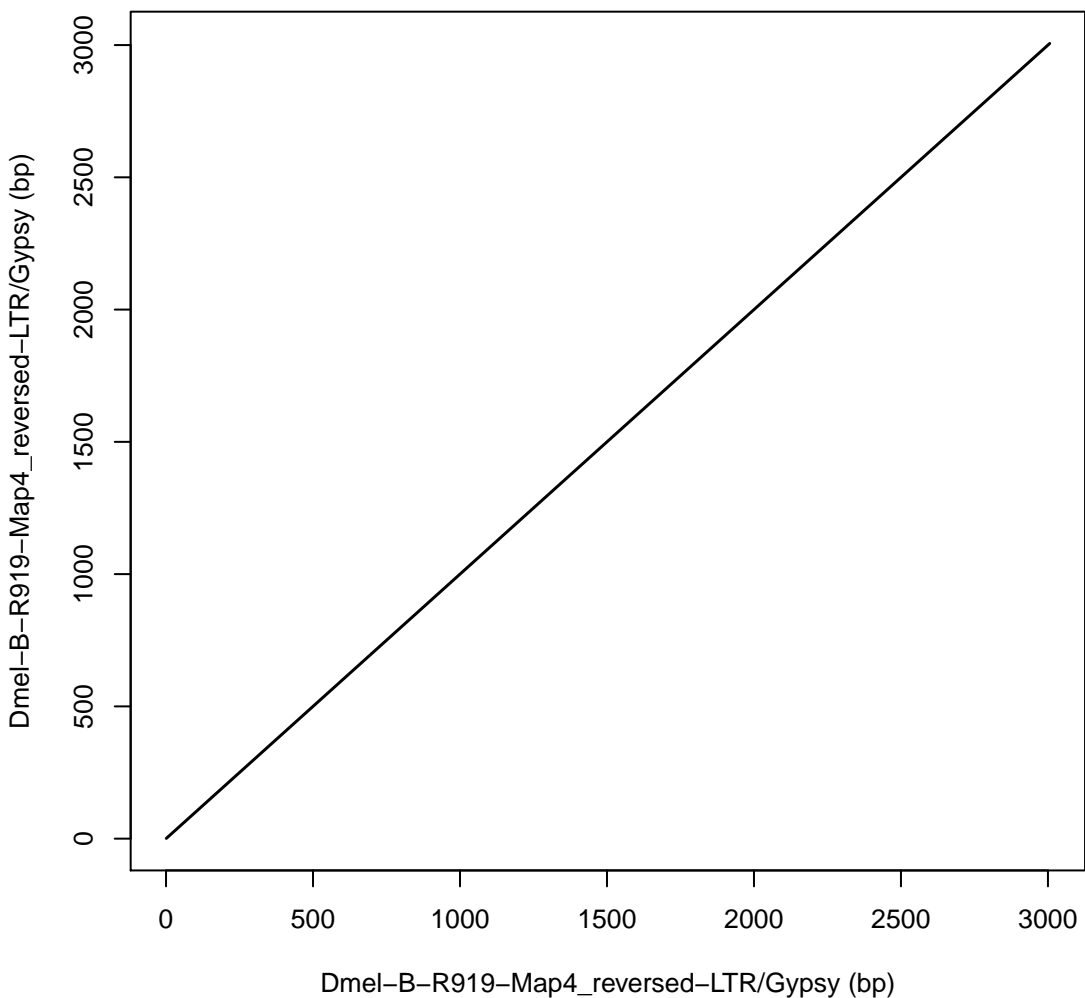
TE: Dmel-B-R919-Map4_reversed-LTR/Gypsy
consensus size: 3006bp; fragments: 49; full length: 0 (>=2705.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

