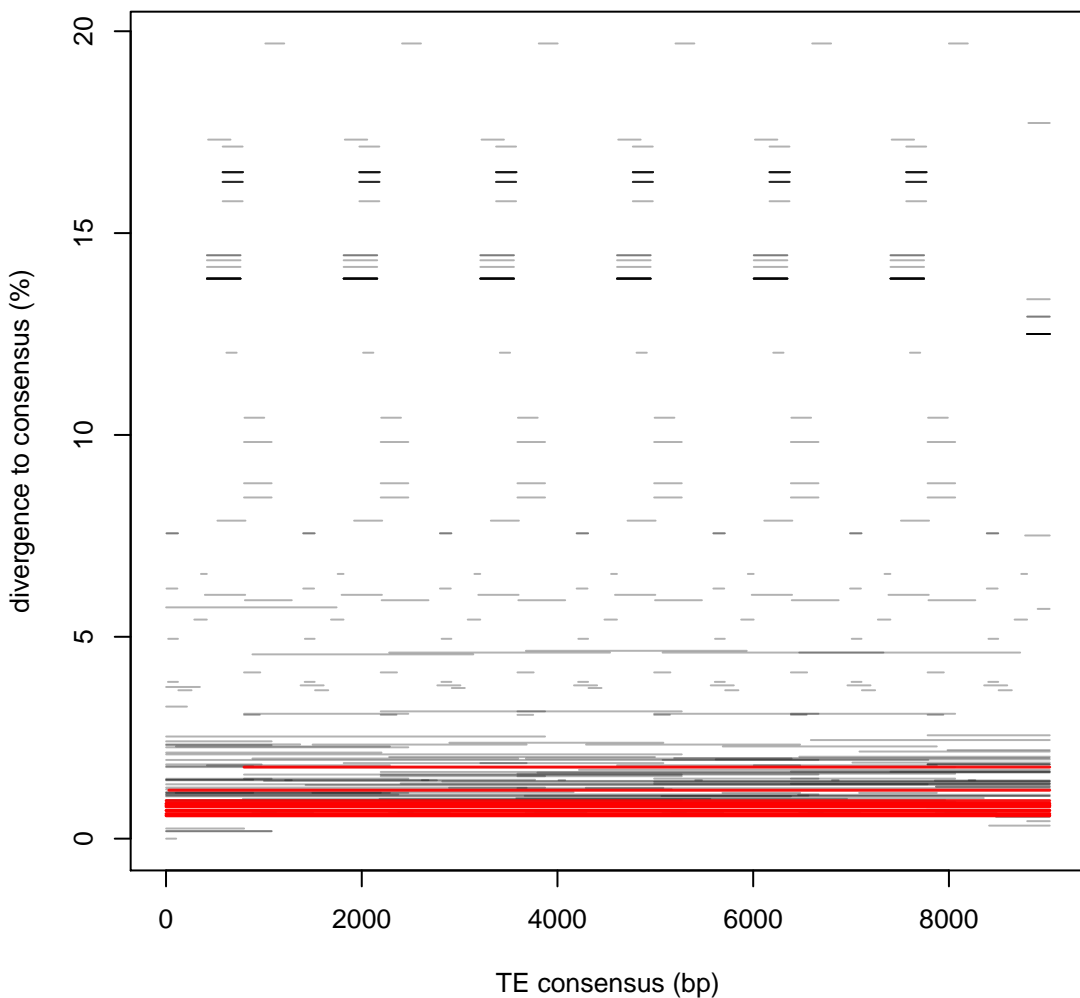
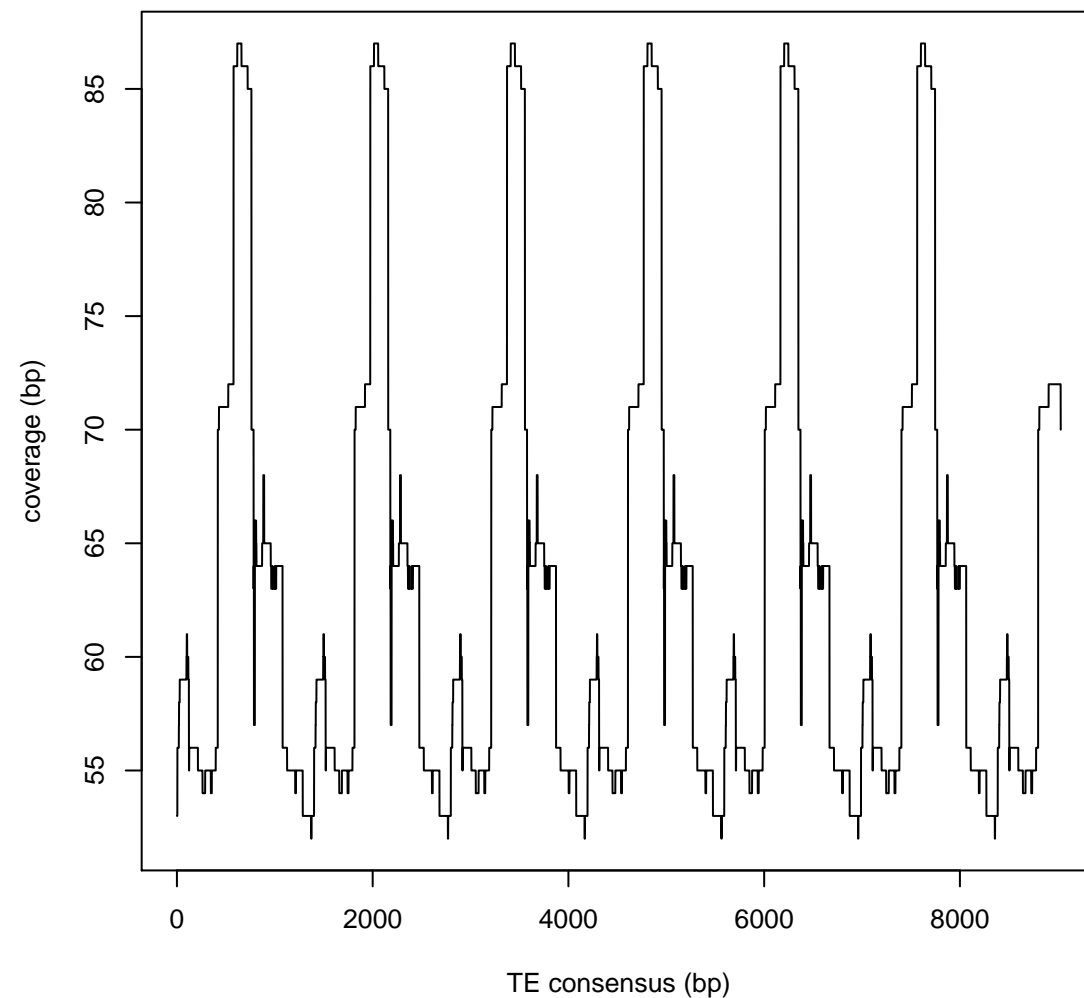


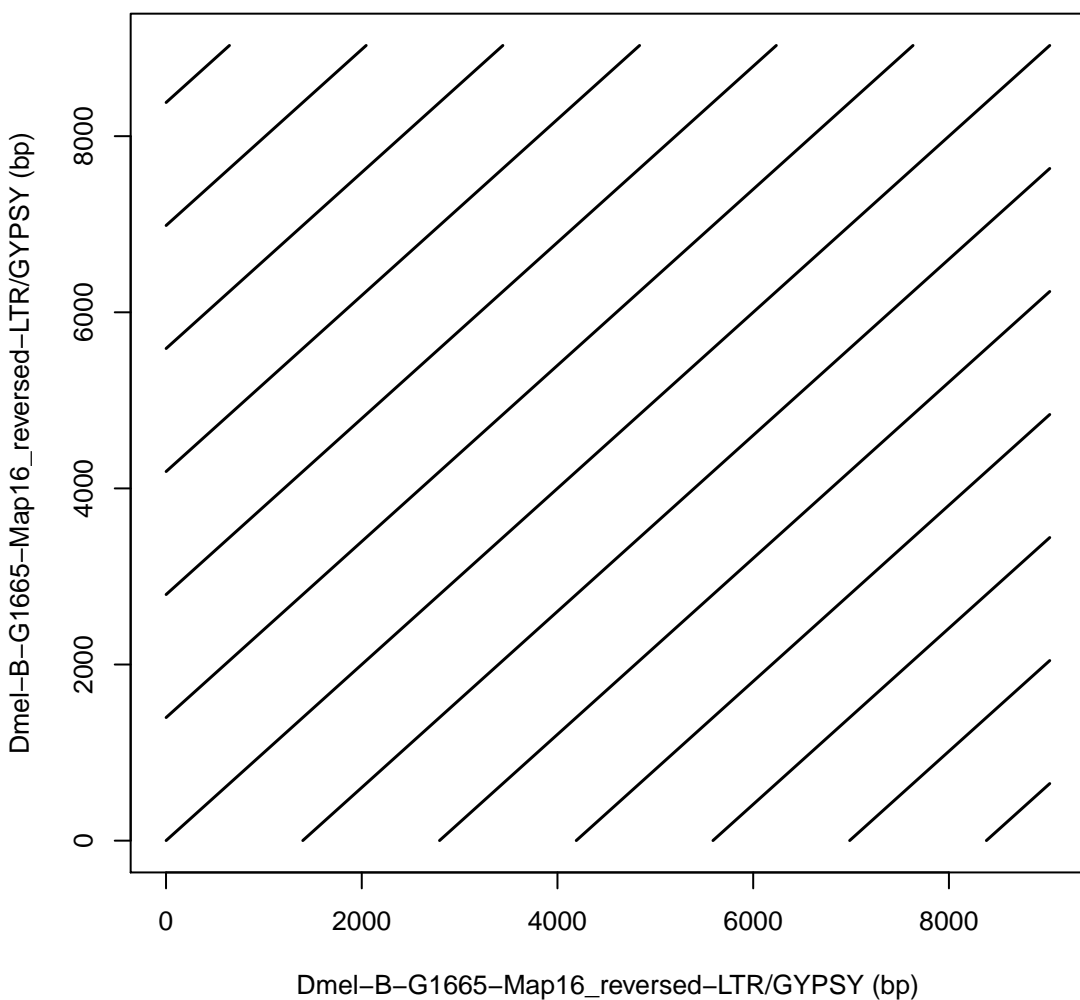
TE: Dmel-B-G1665-Map16_reversed-LTR/Gypsy
consensus size: 9030bp; fragments: 479; full length: 16 (>=8127bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

