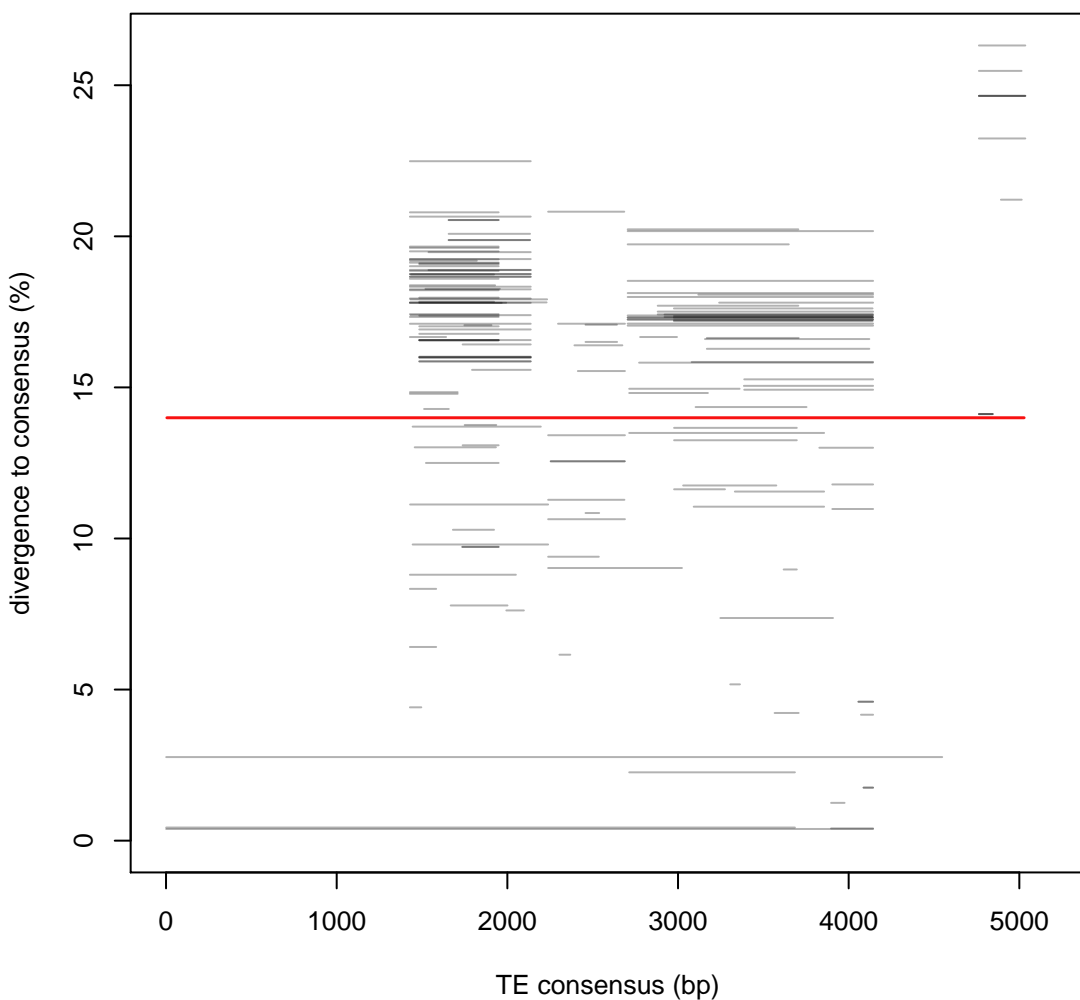
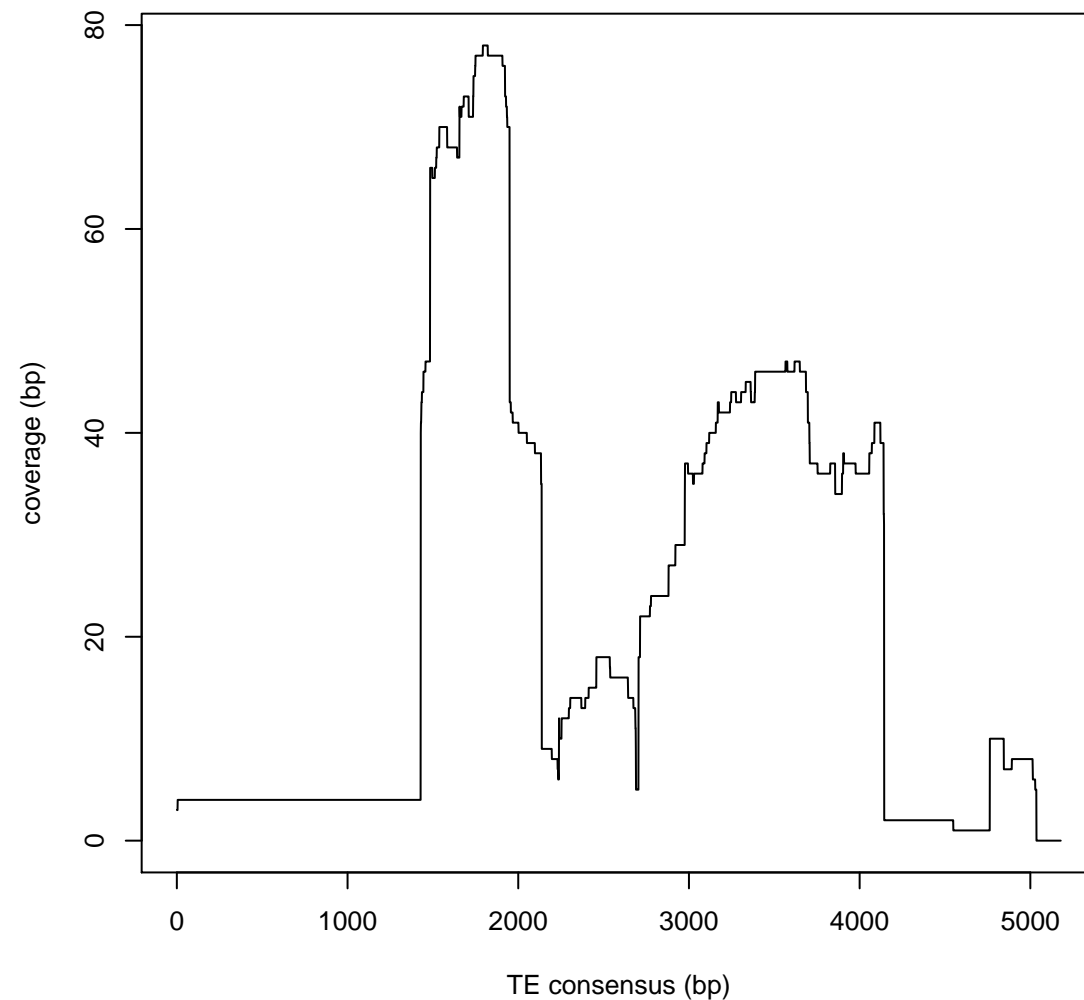


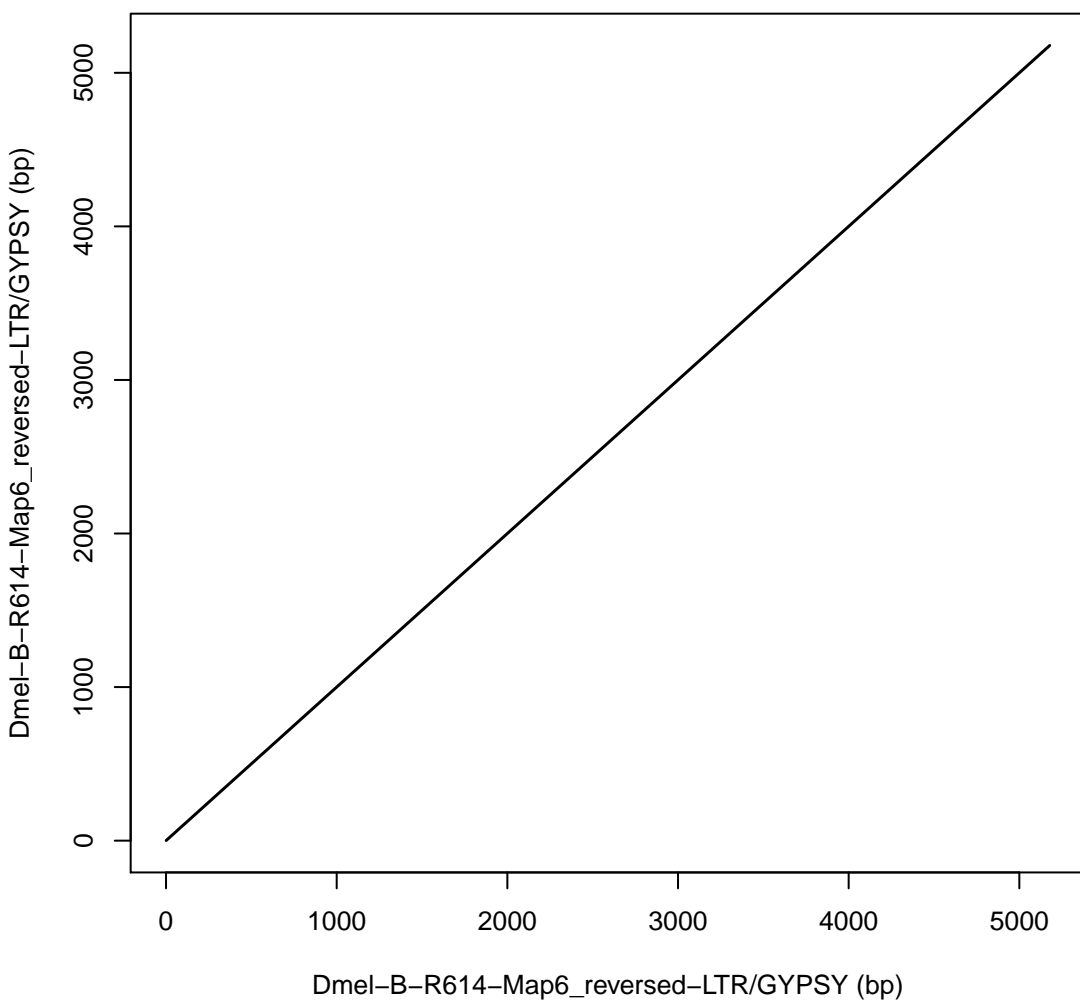
TE: Dmel-B-R614-Map6_reversed-LTR/GYPSY
consensus size: 5178bp; fragments: 170; full length: 1 (>=4660.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

