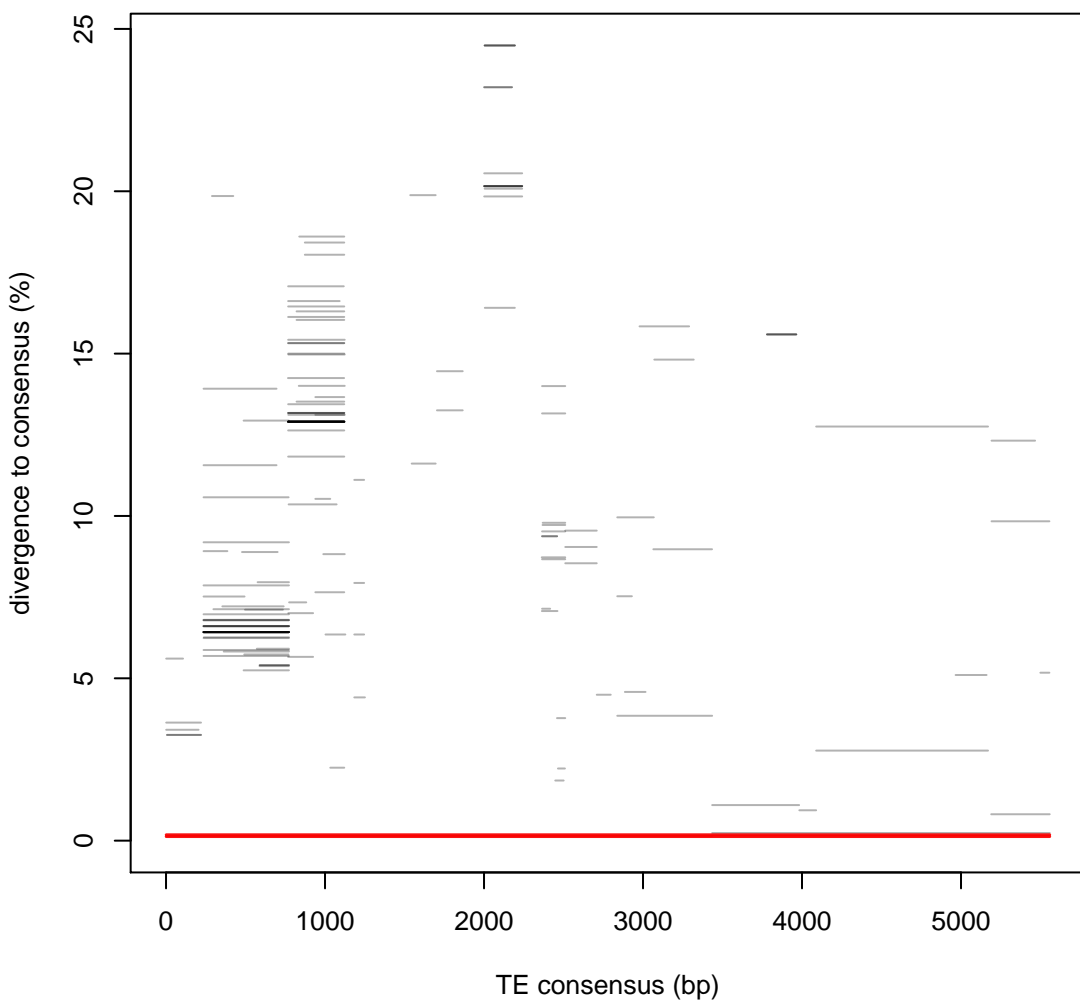
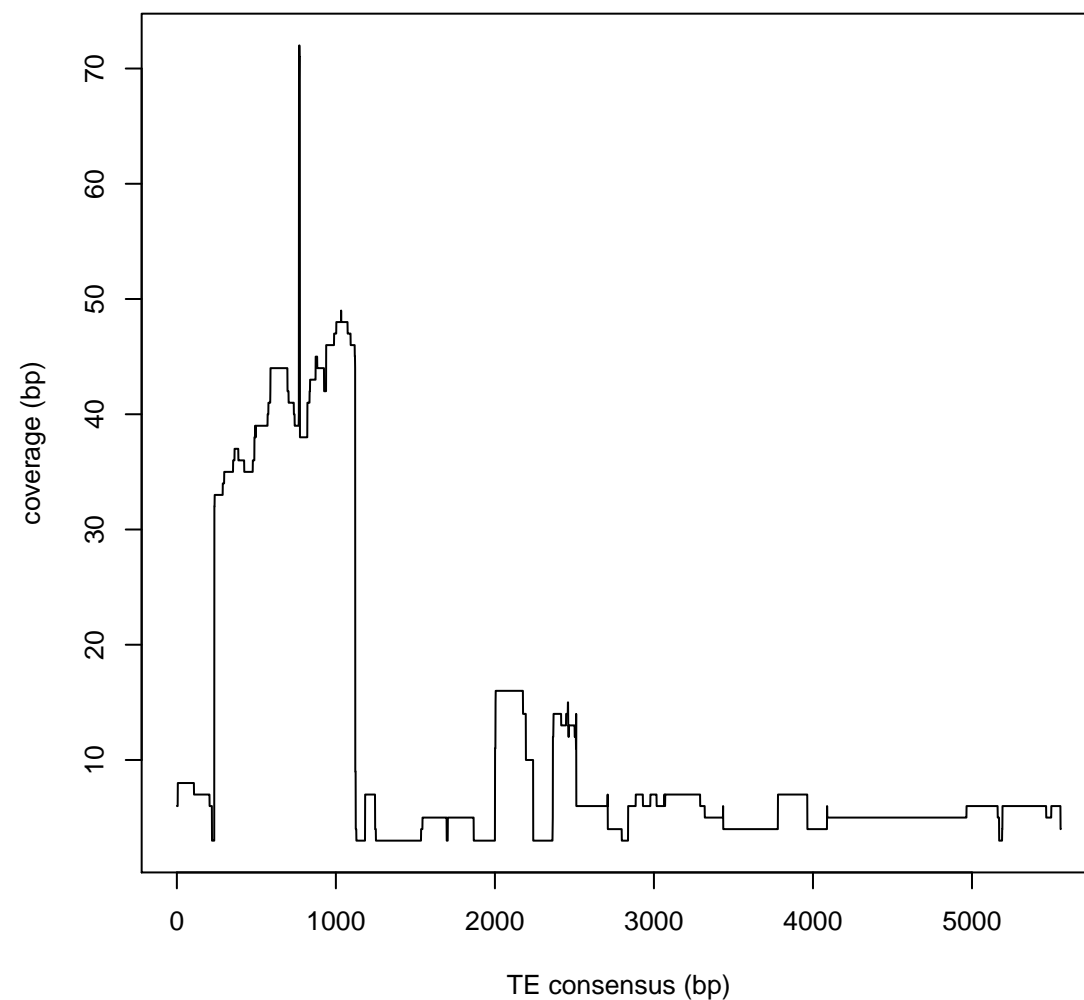


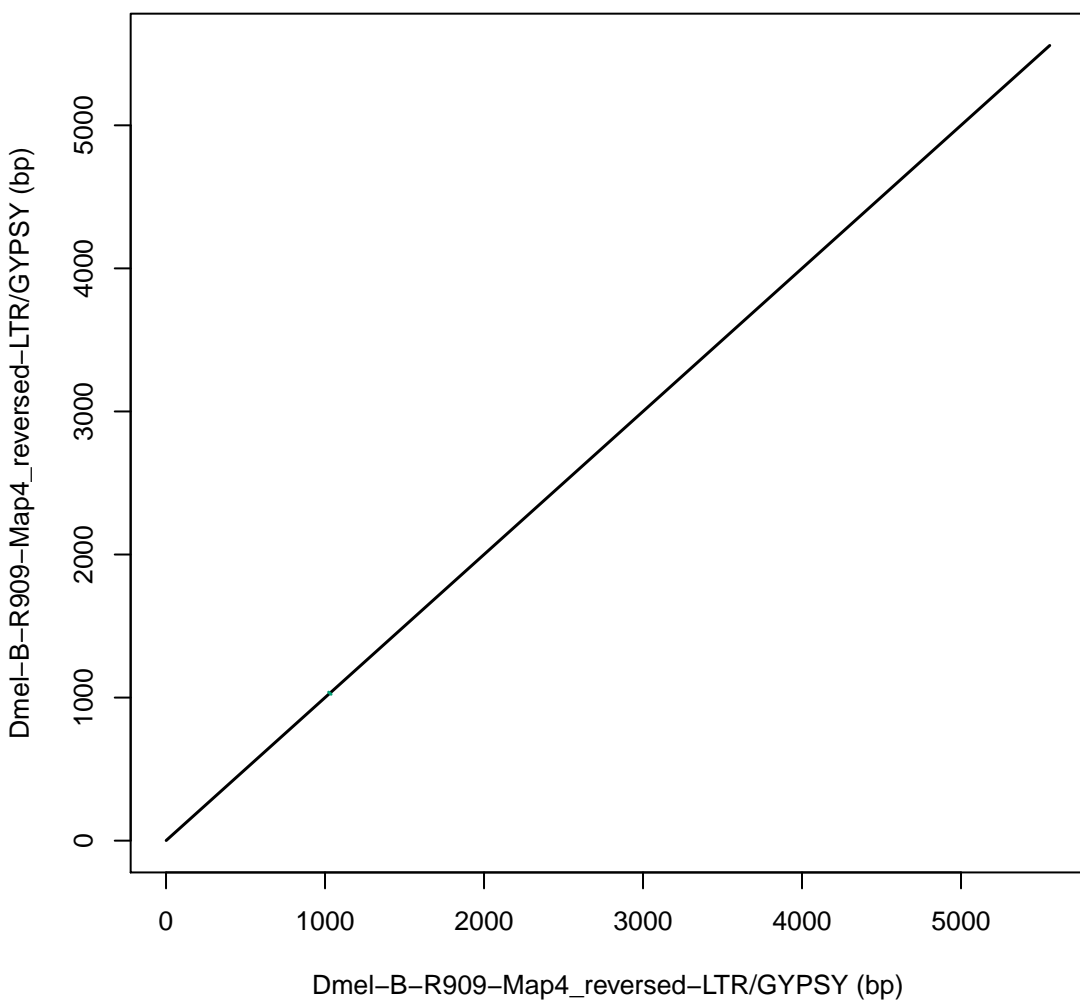
TE: Dmel-B-R909-Map4_reversed-LTR/GYPSY
consensus size: 5558bp; fragments: 160; full length: 2 (>=5002.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

