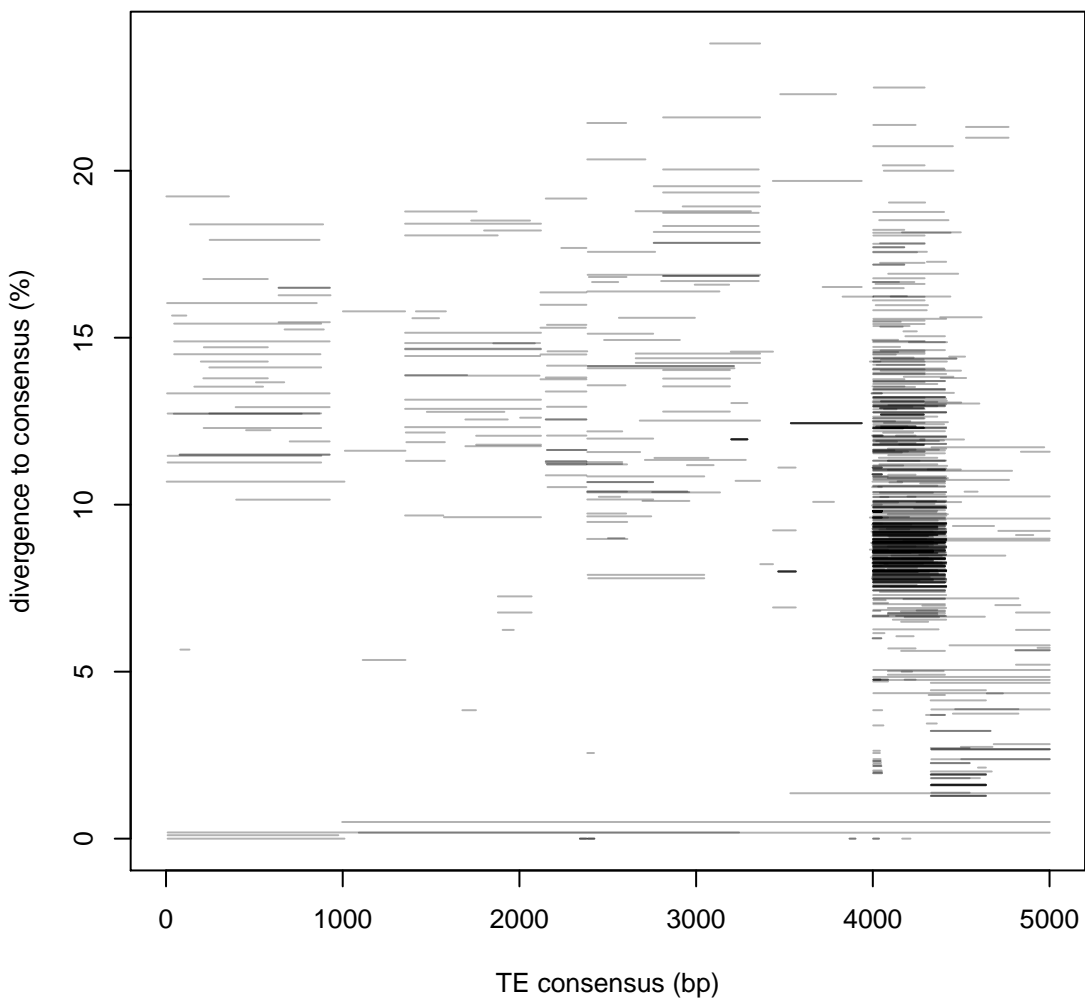
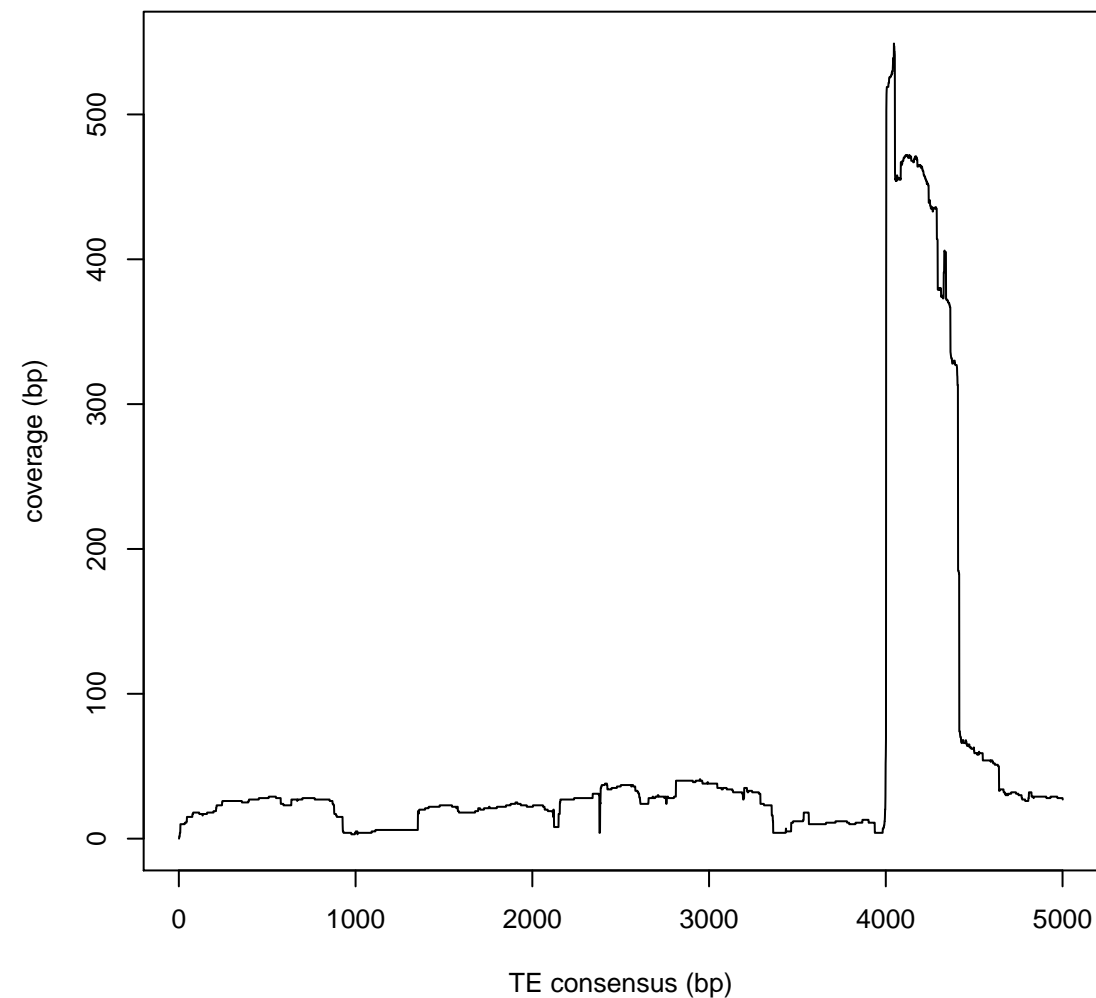


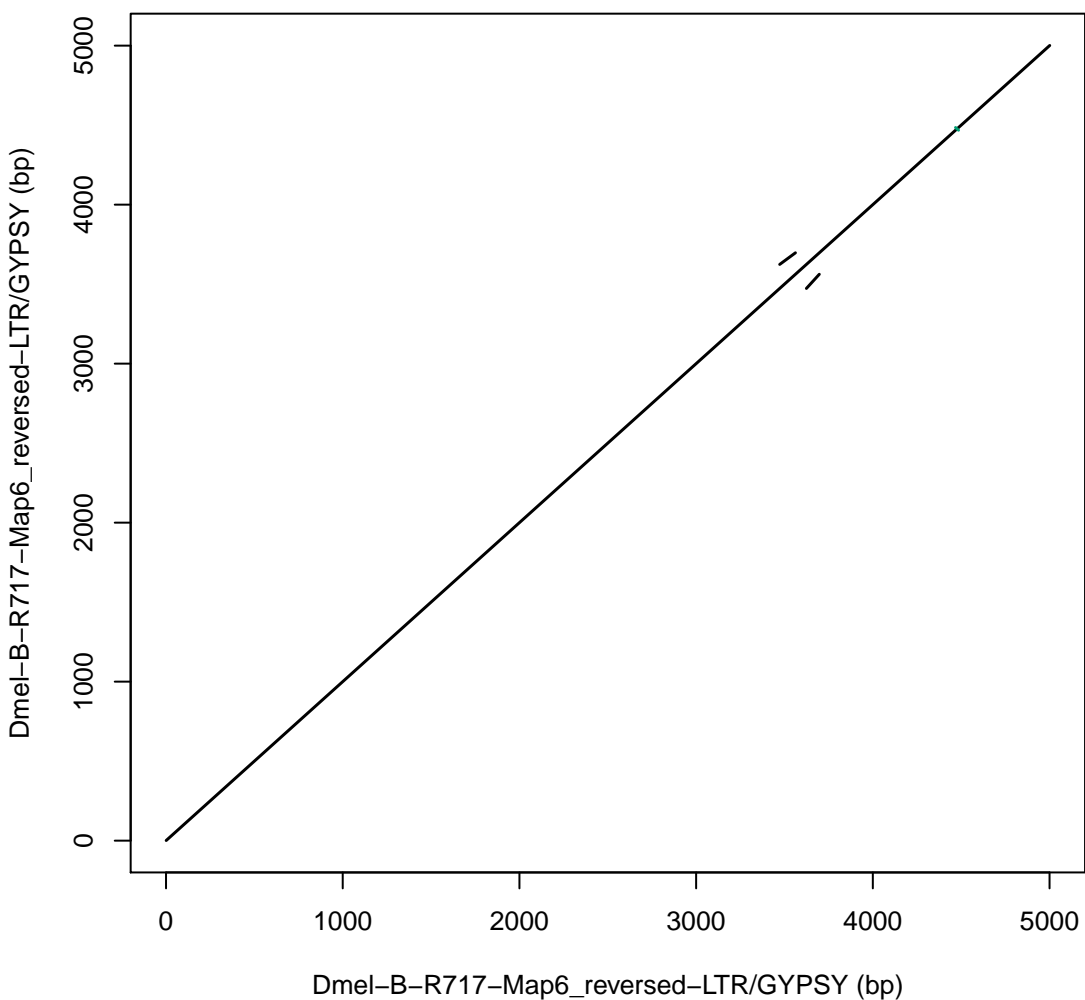
TE: Dmel-B-R717-Map6_reversed-LTR/GYPSY
consensus size: 5001bp; fragments: 887; full length: 0 (>=4500.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

