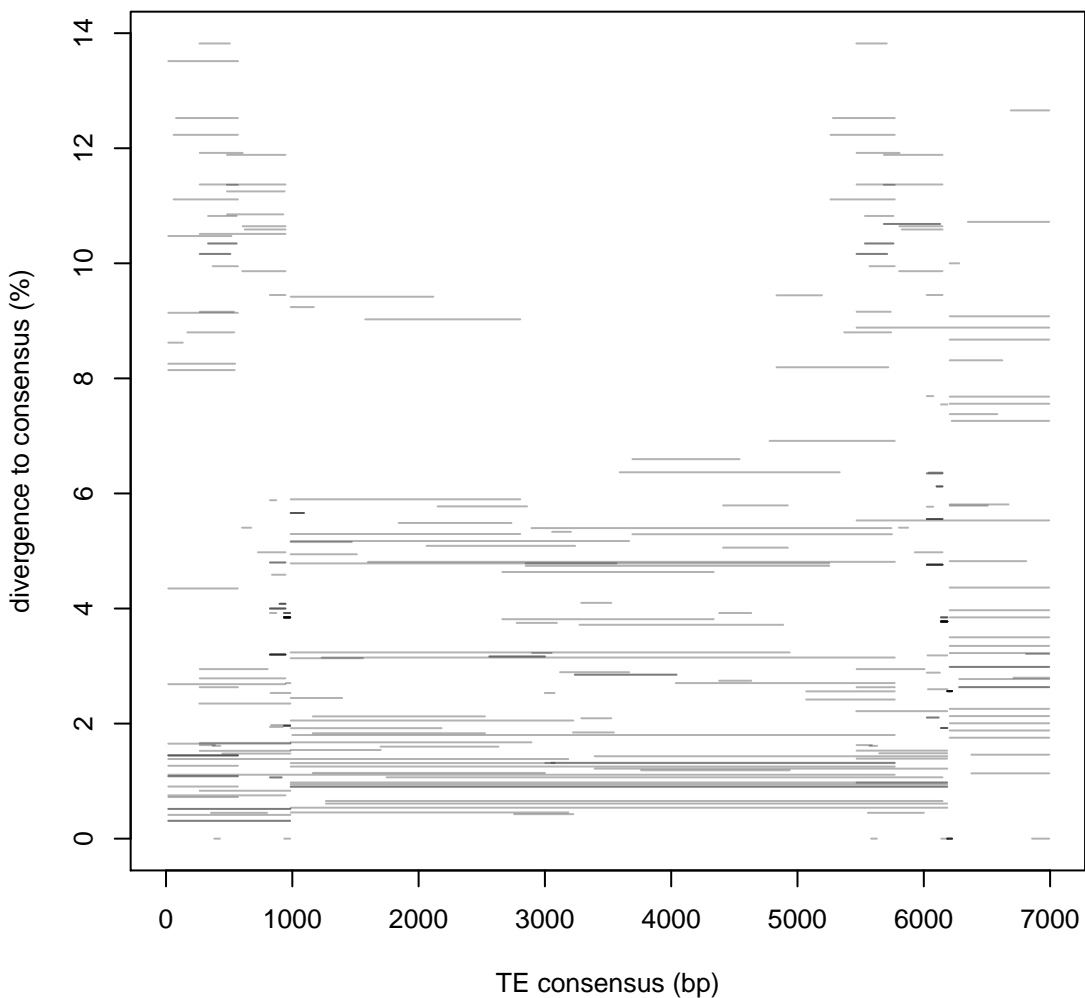
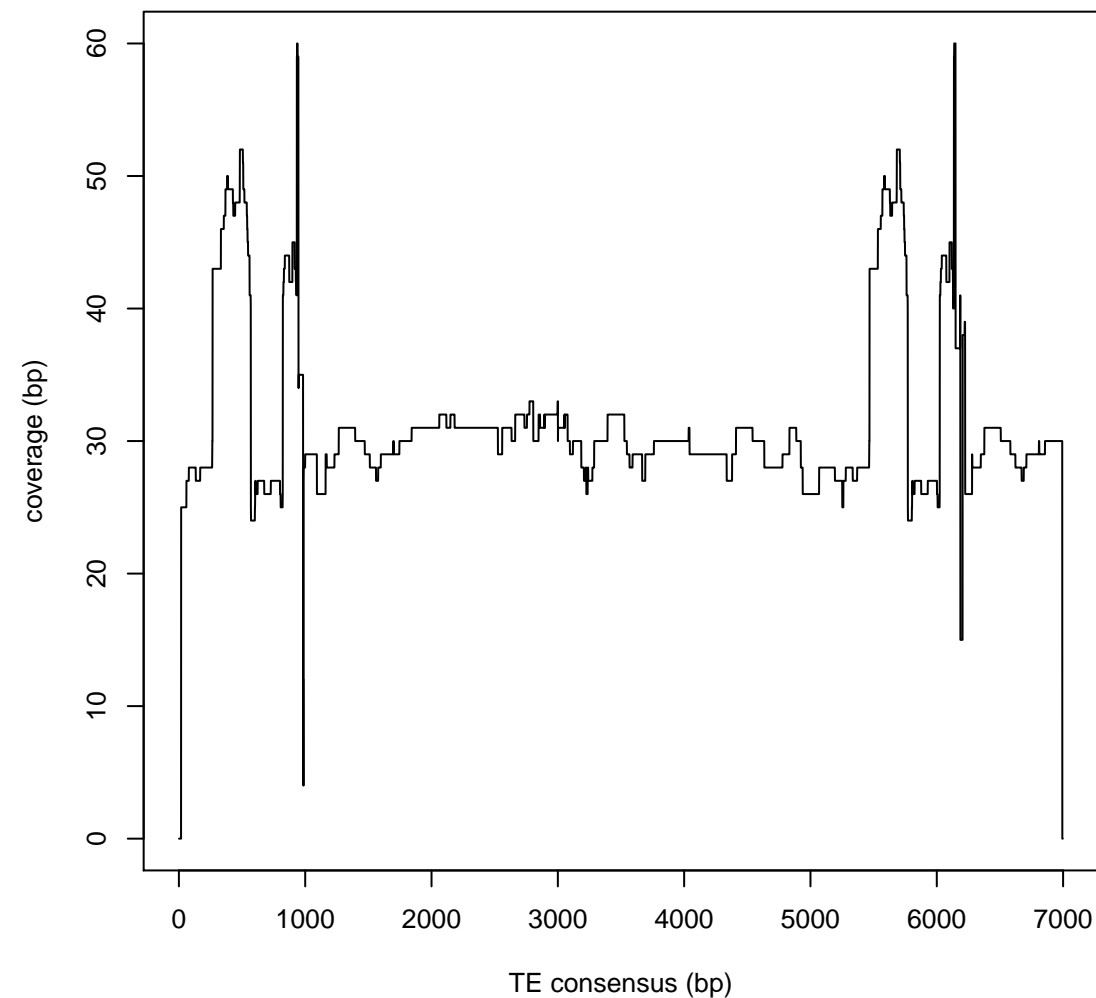


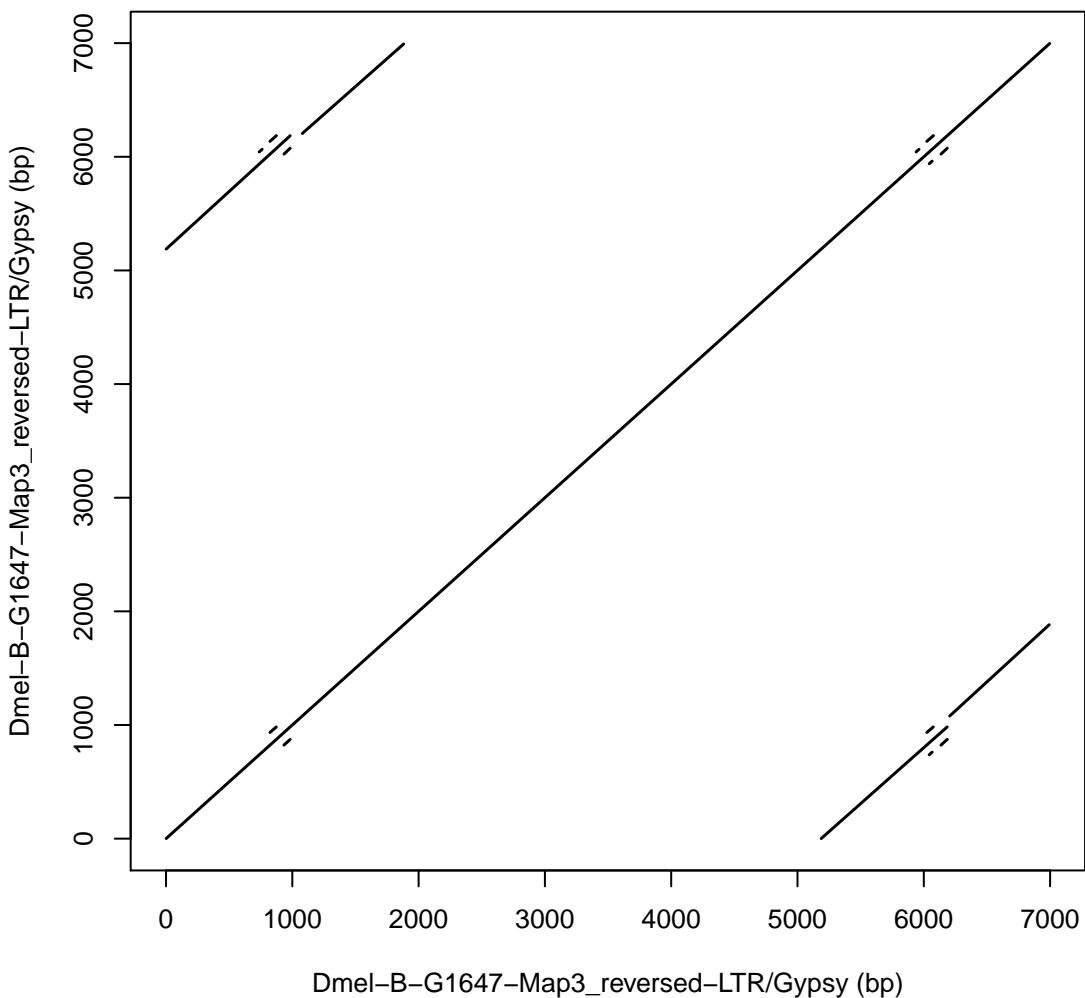
TE: Dmel-B-G1647-Map3_reversed-LTR/Gypsy
consensus size: 6997bp; fragments: 305; full length: 0 (>=6297.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

