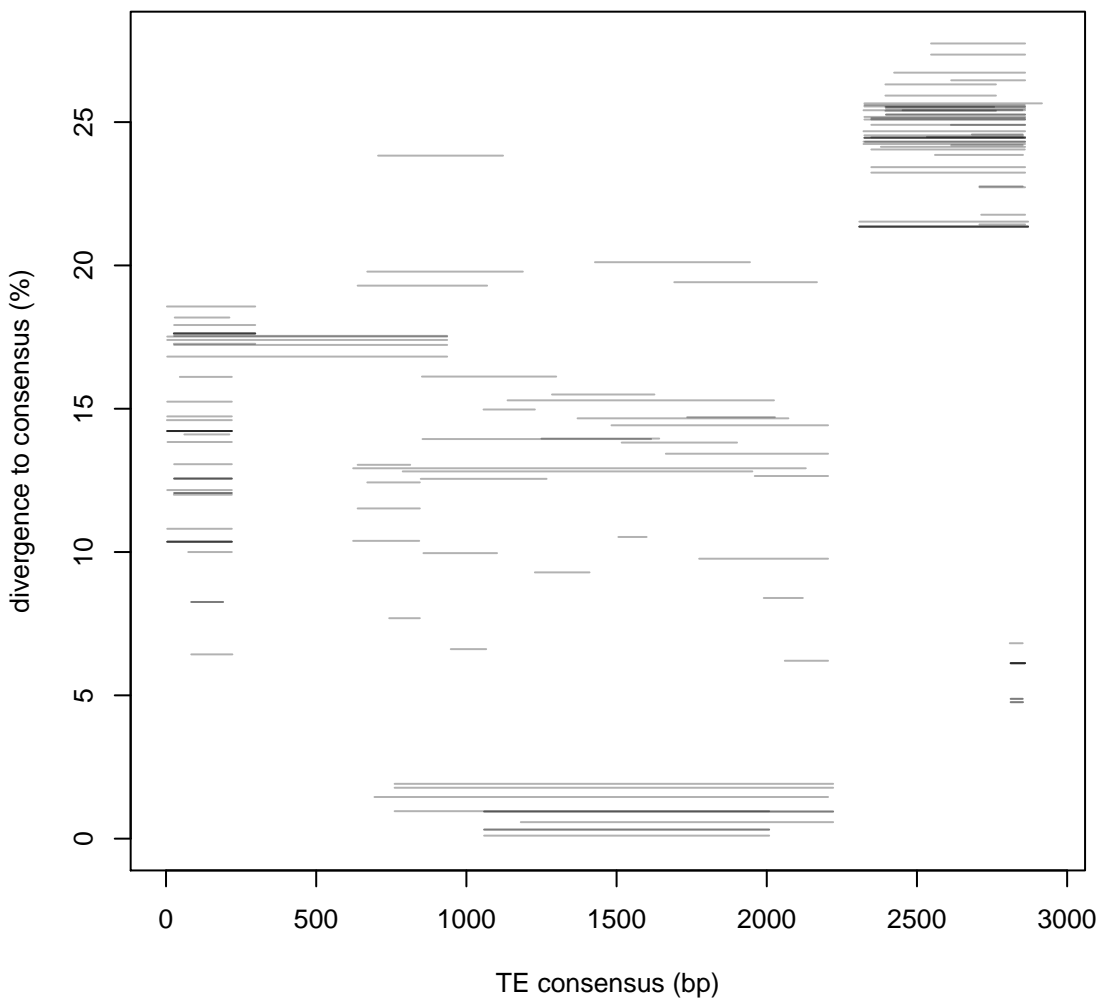
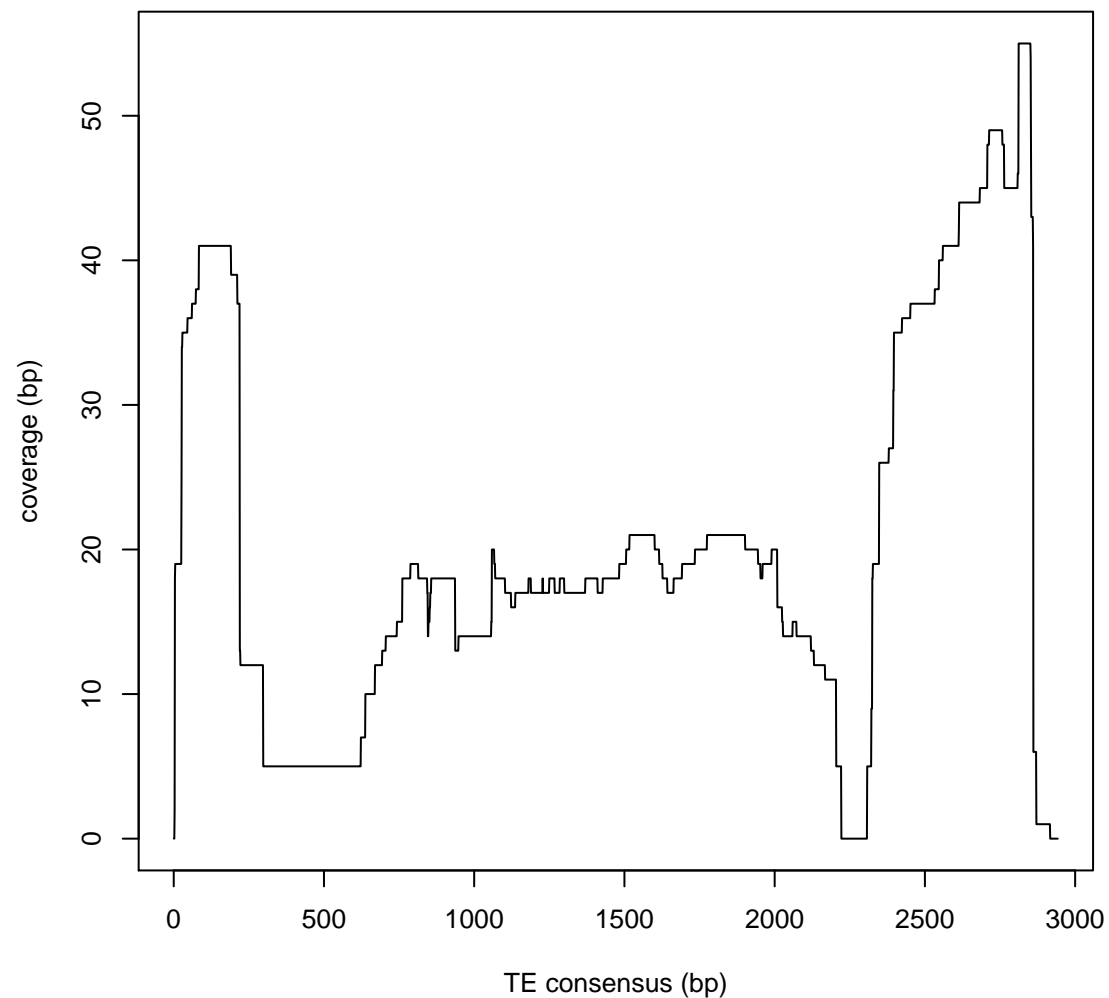


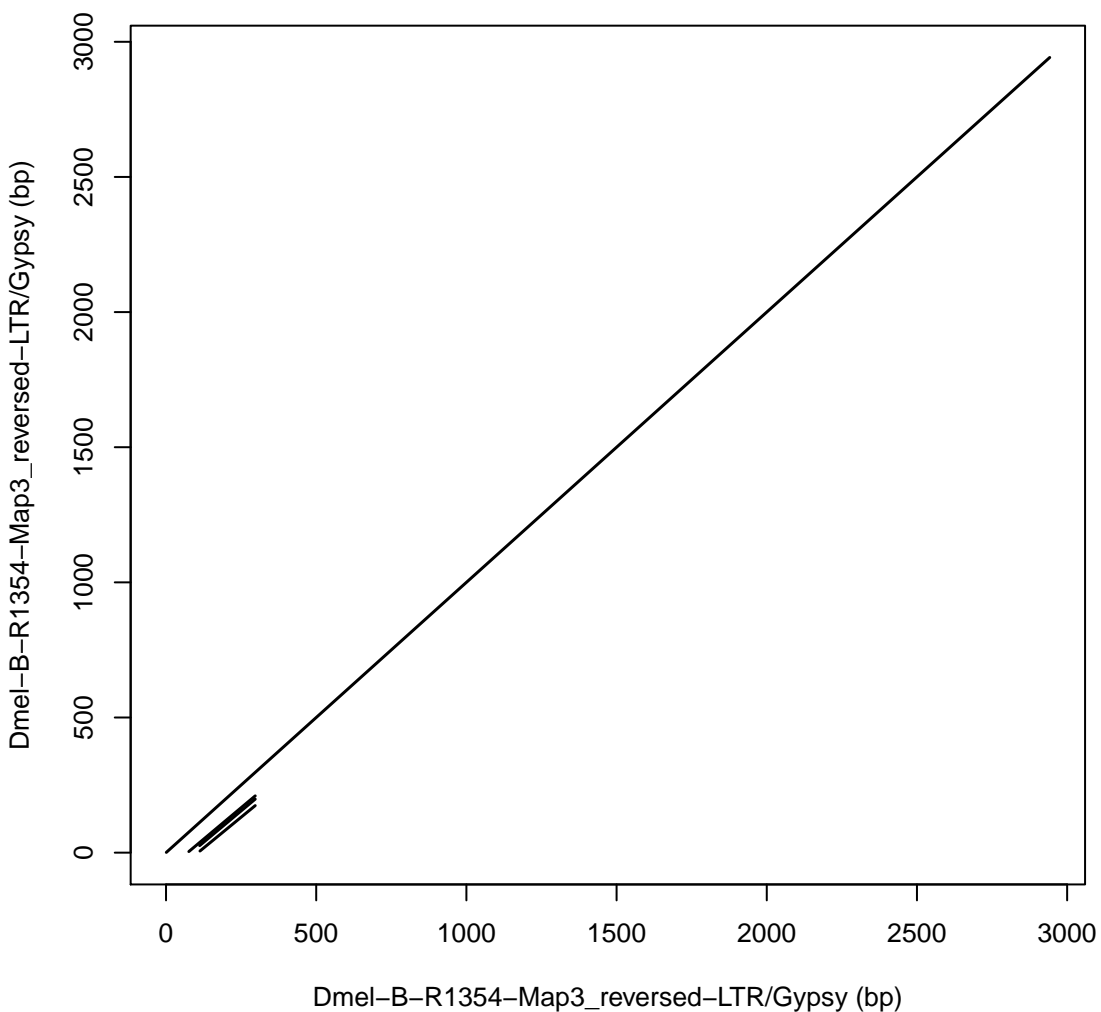
TE: Dmel-B-R1354-Map3_reversed-LTR/Gypsy
consensus size: 2942bp; fragments: 142; full length: 0 (>=2647.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

