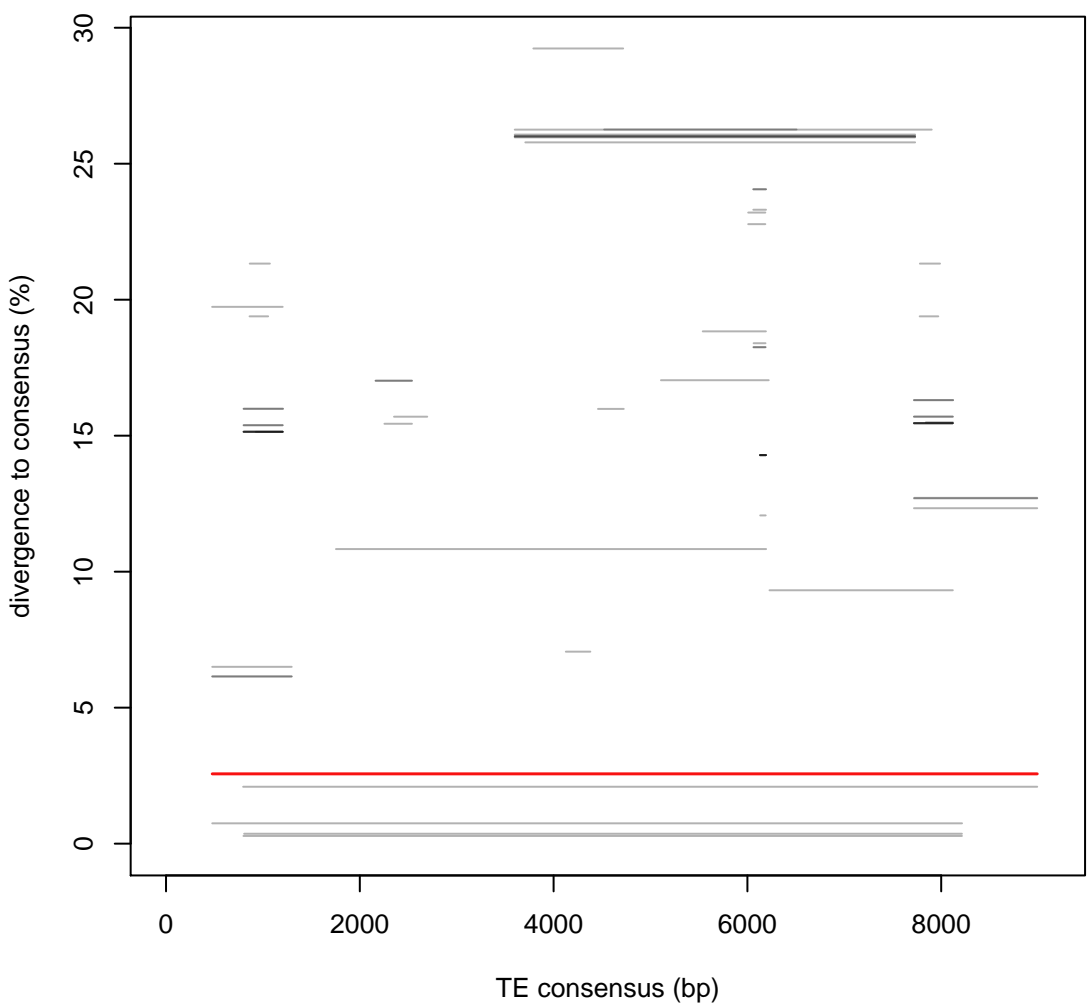
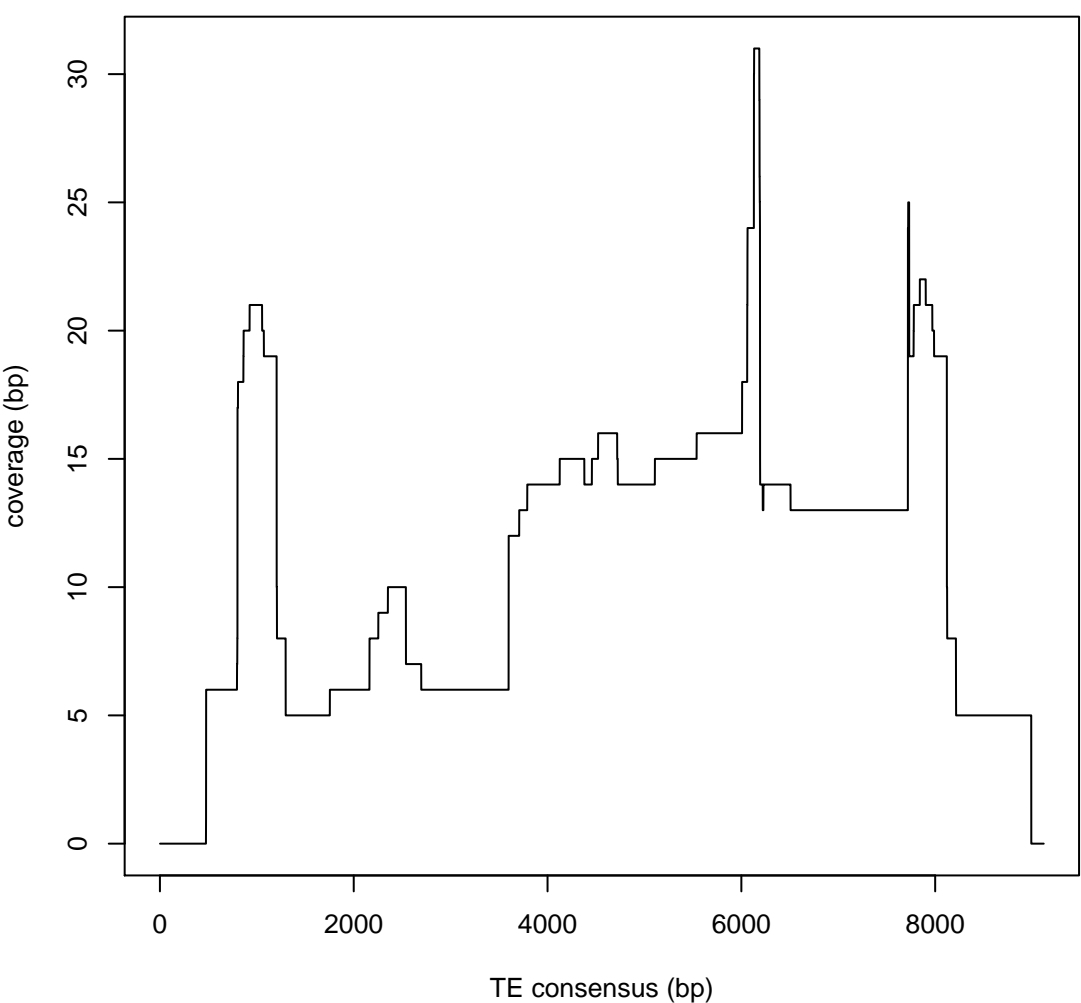


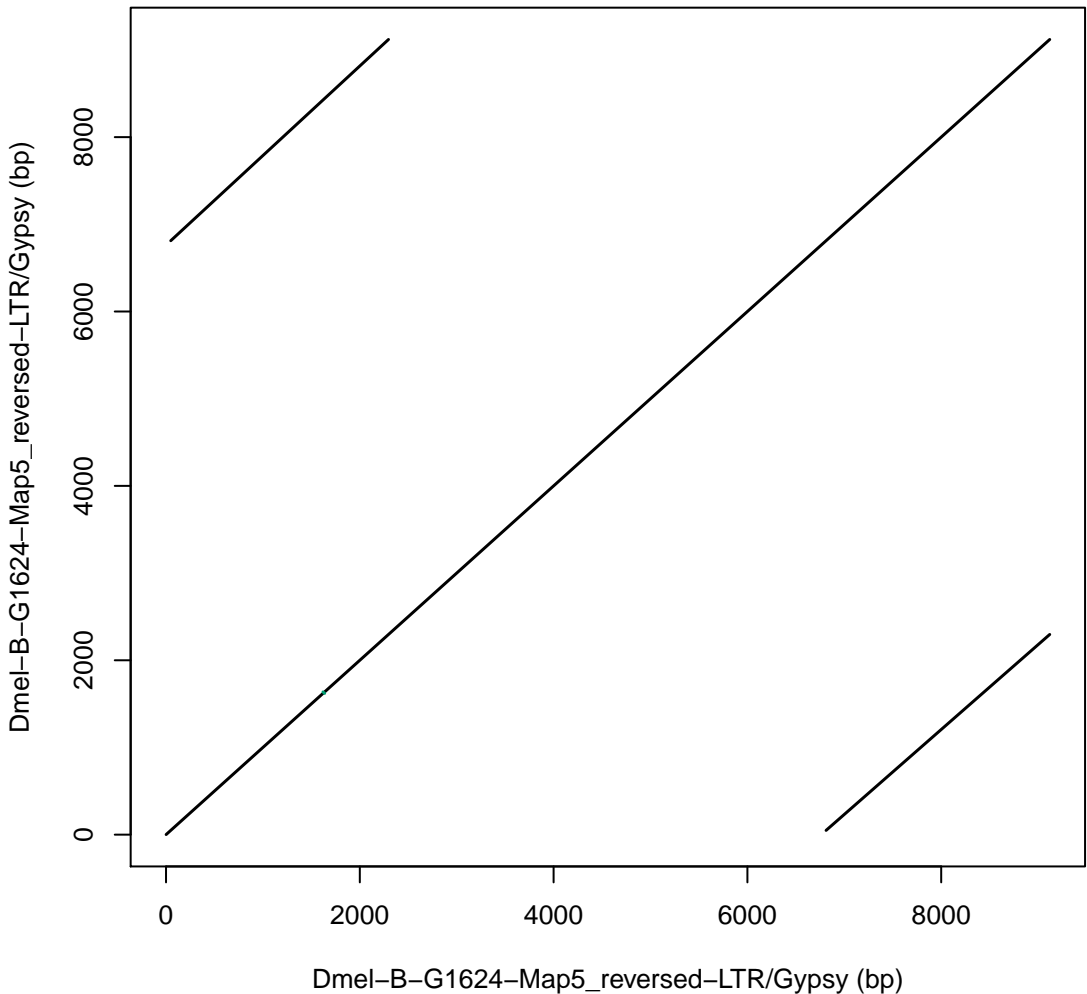
TE: Dmel-B-G1624-Map5_reversed-LTR/Gypsy
consensus size: 9120bp; fragments: 70; full length: 1 (>=8208bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

