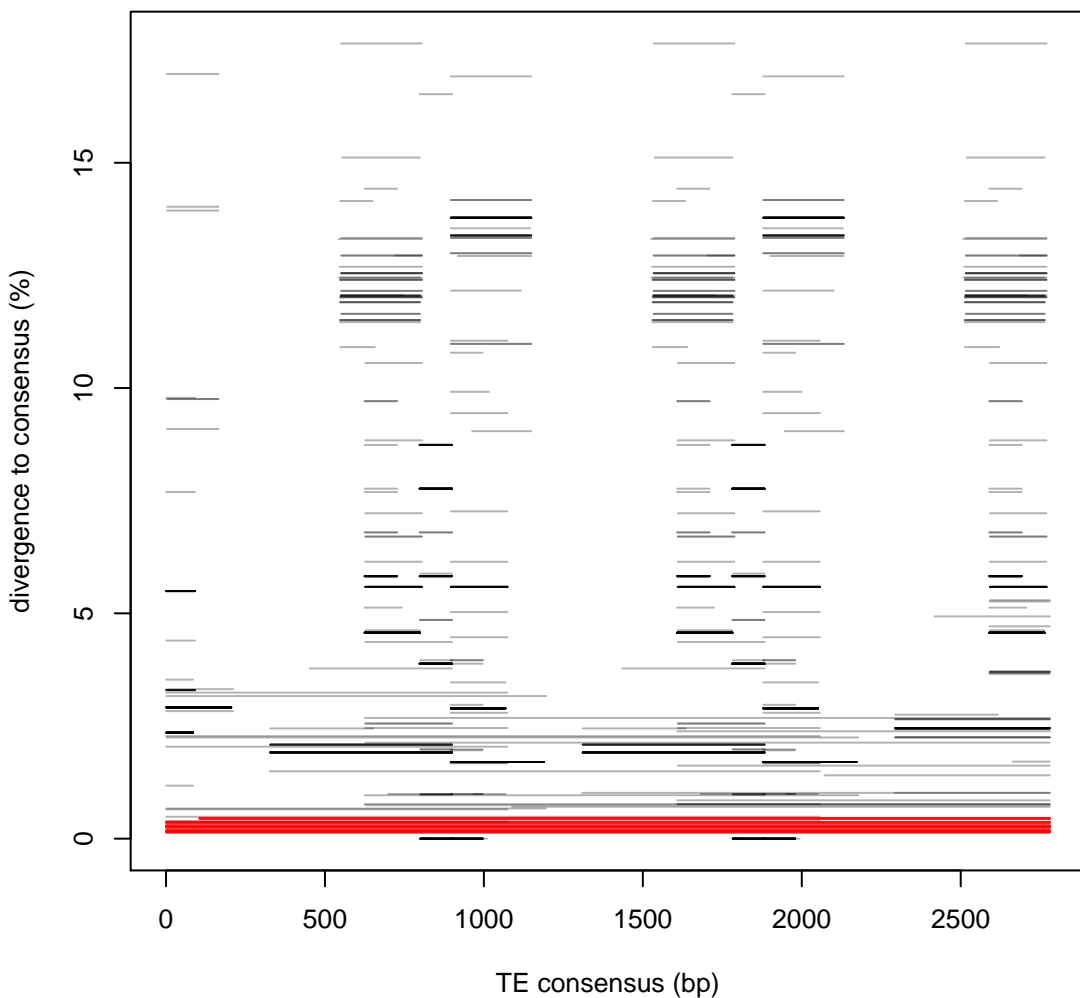
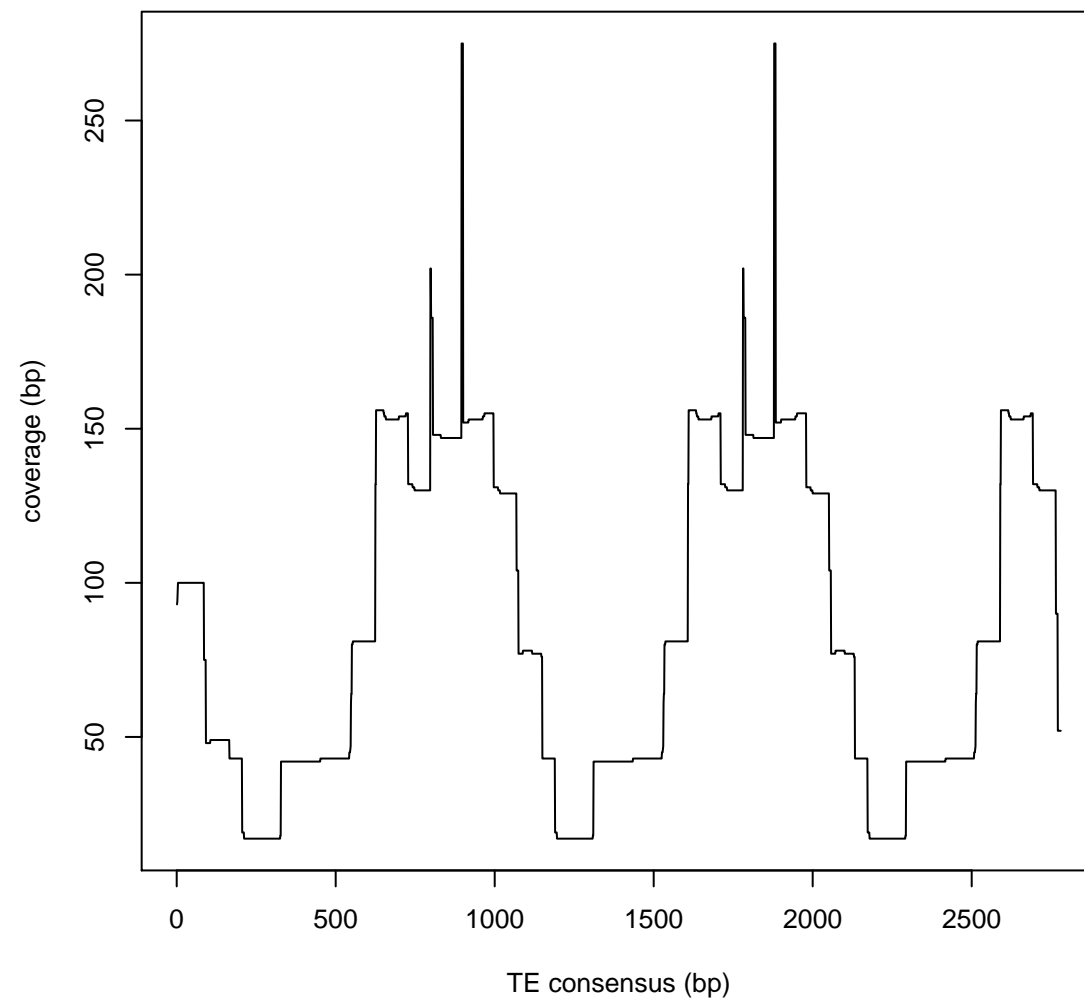


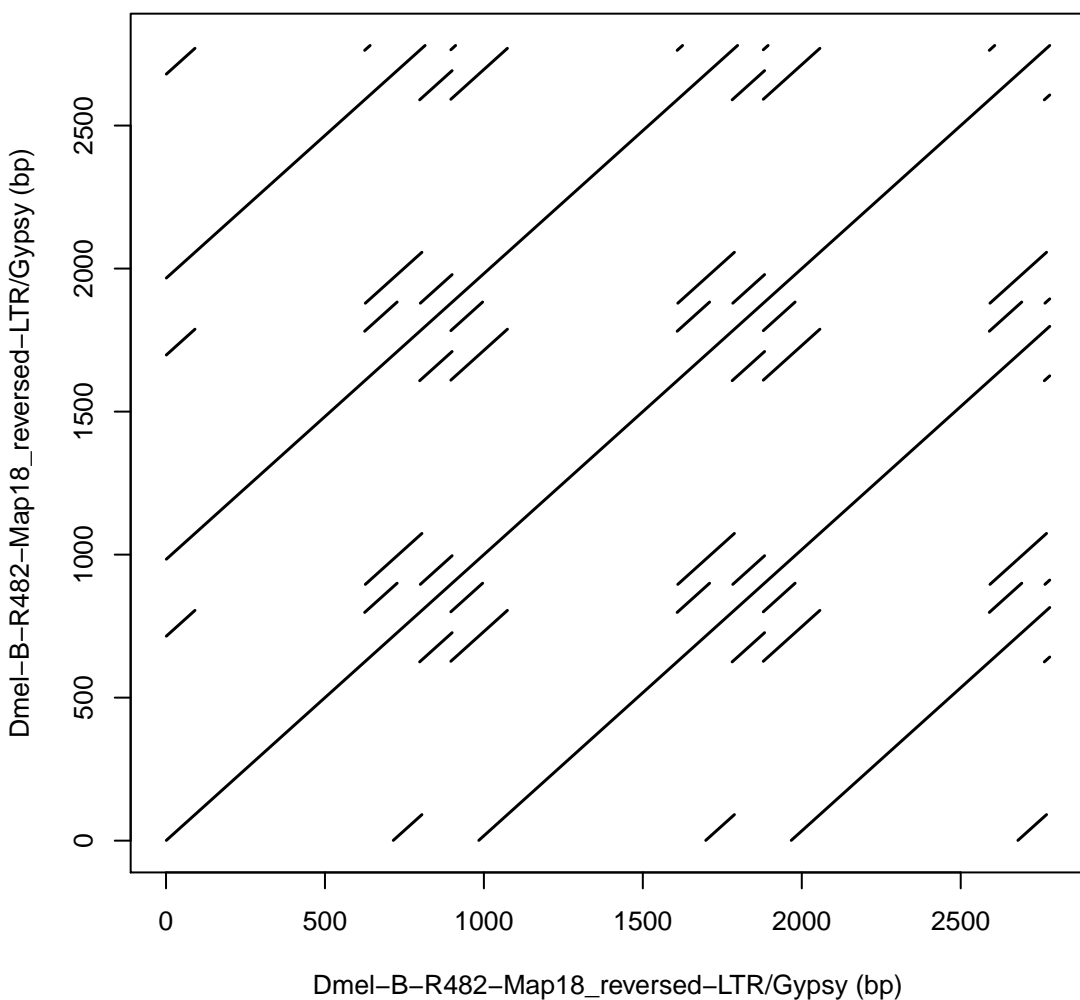
TE: Dmel-B-R482-Map18_reversed-LTR/Gypsy
consensus size: 2780bp; fragments: 980; full length: 6 (≥ 2502 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

