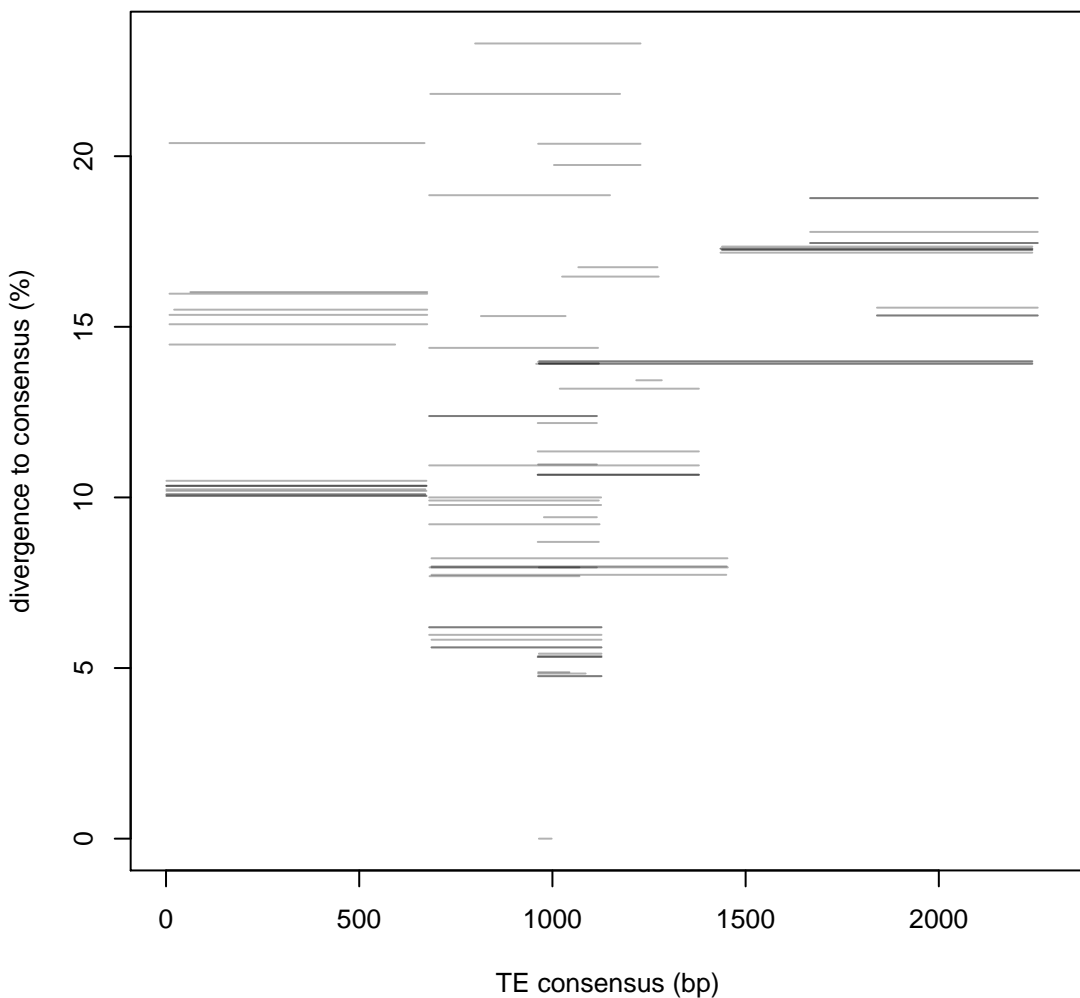
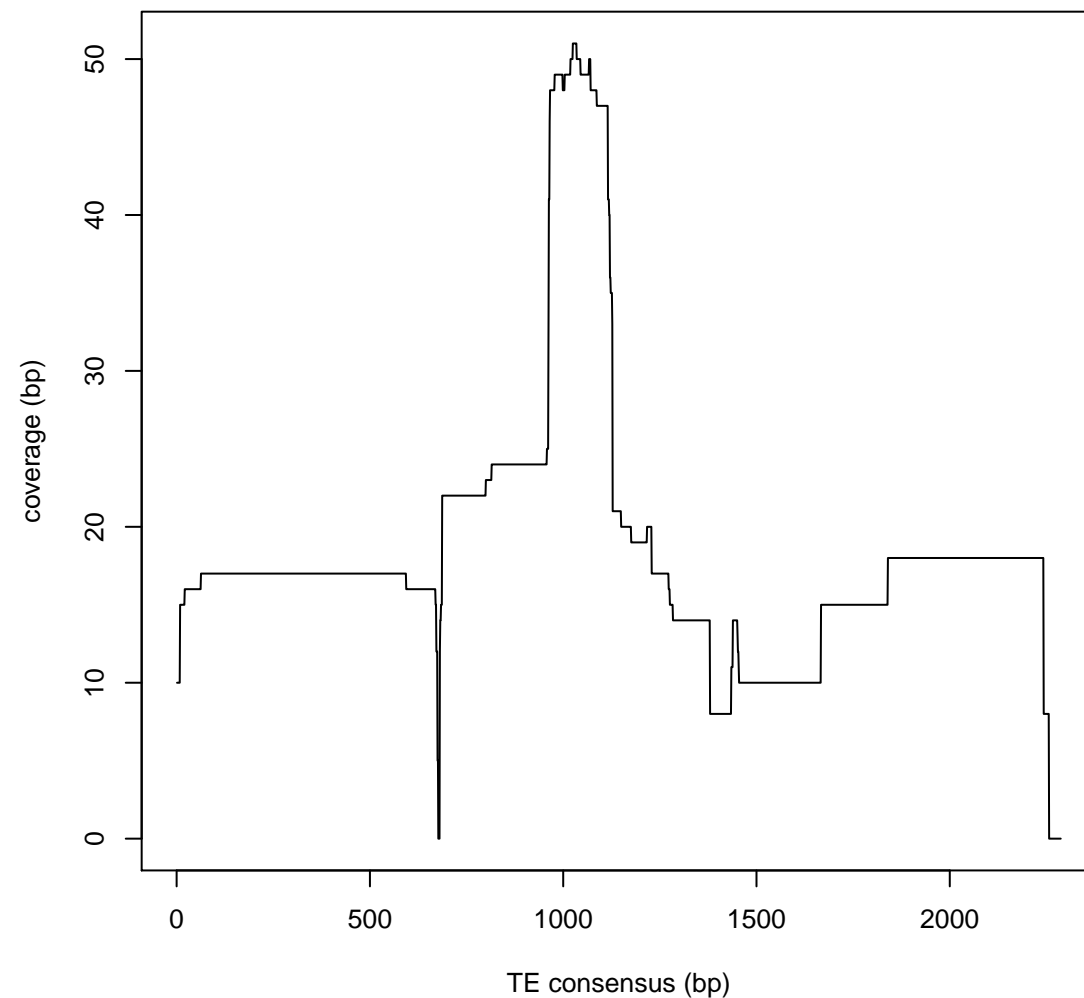


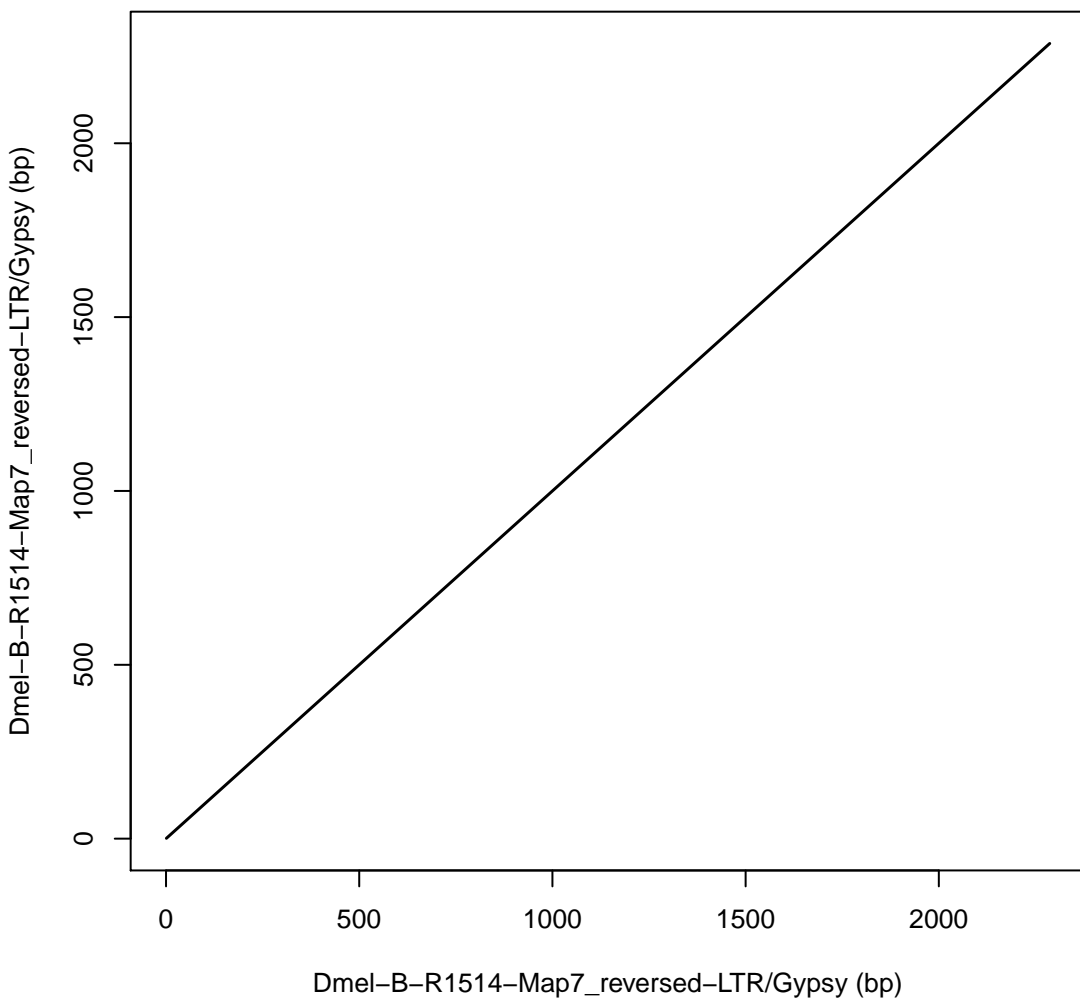
TE: Dmel-B-R1514-Map7\_reversed-LTR/Gypsy  
consensus size: 2287bp; fragments: 85; full length: 0 (>=2058.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

