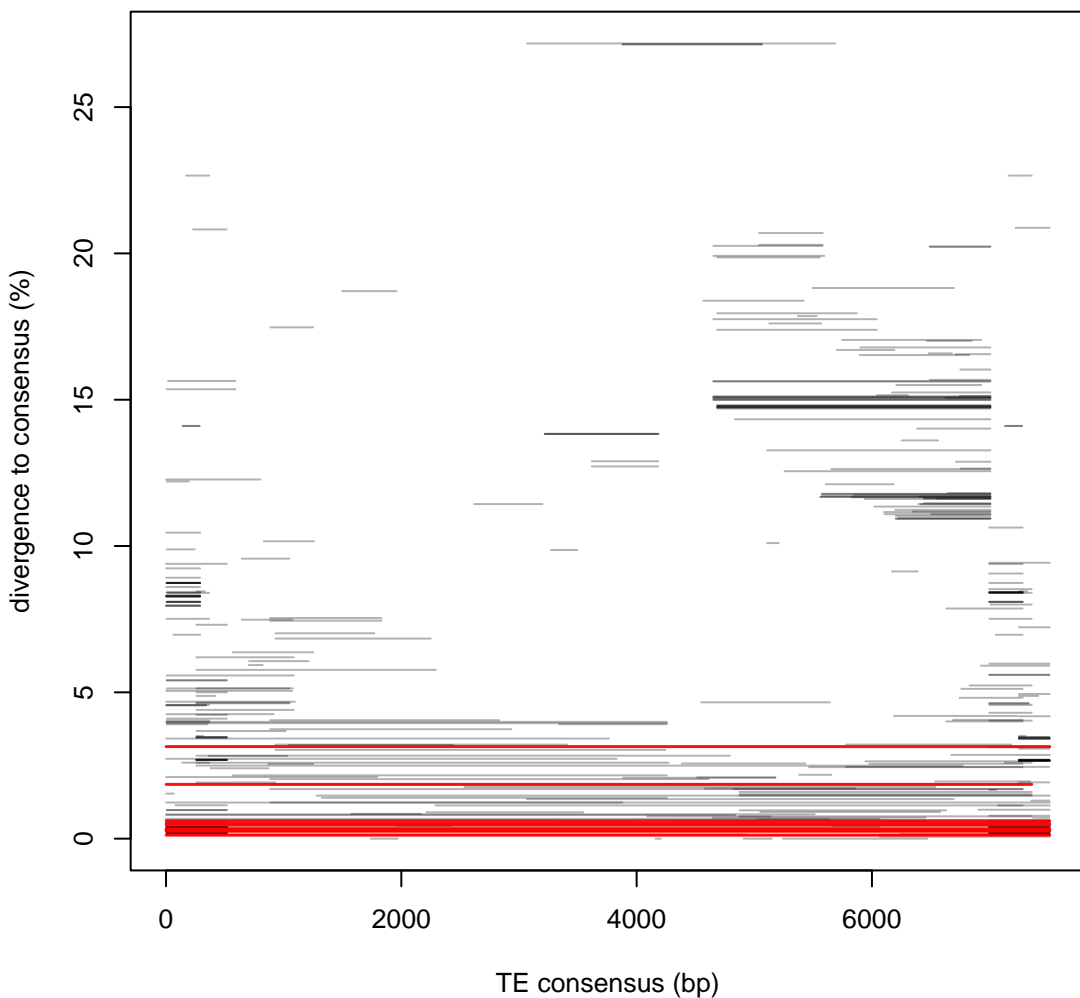
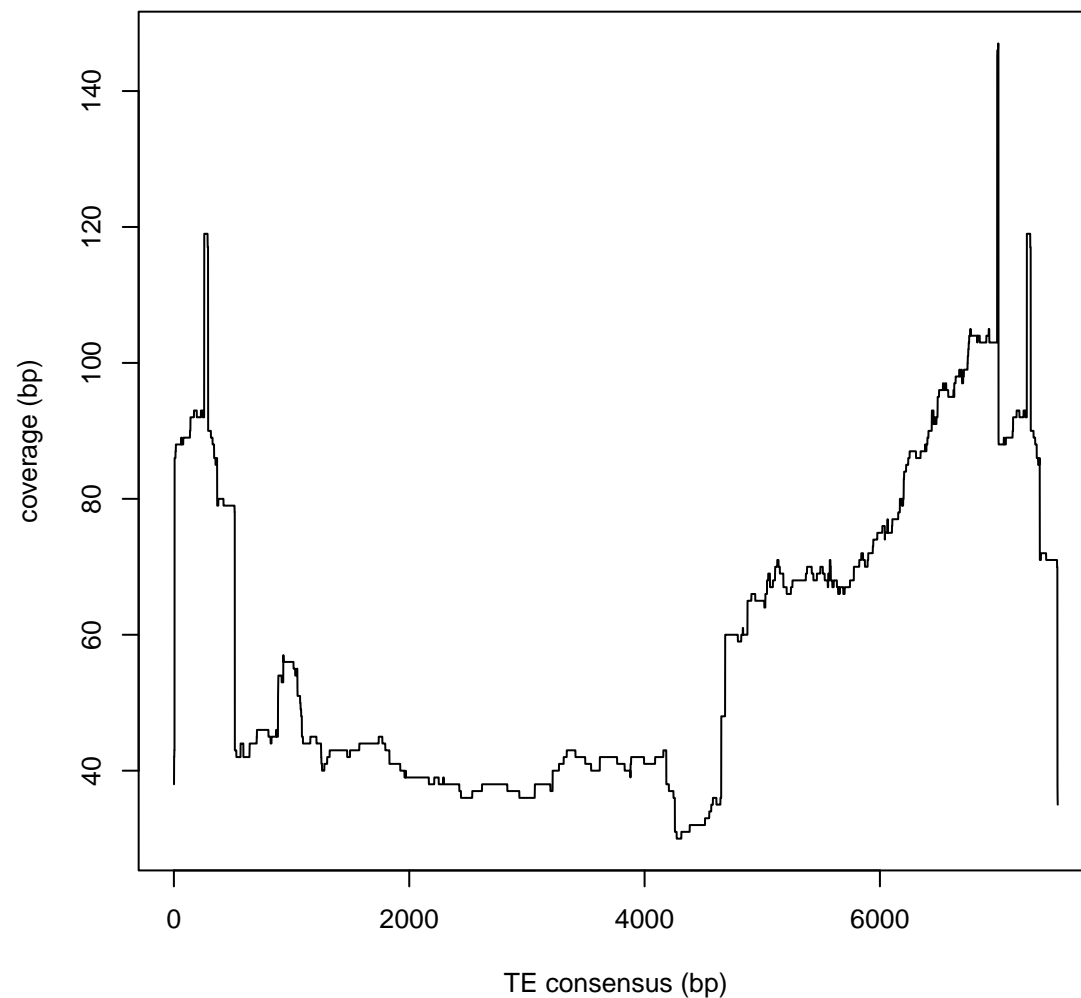


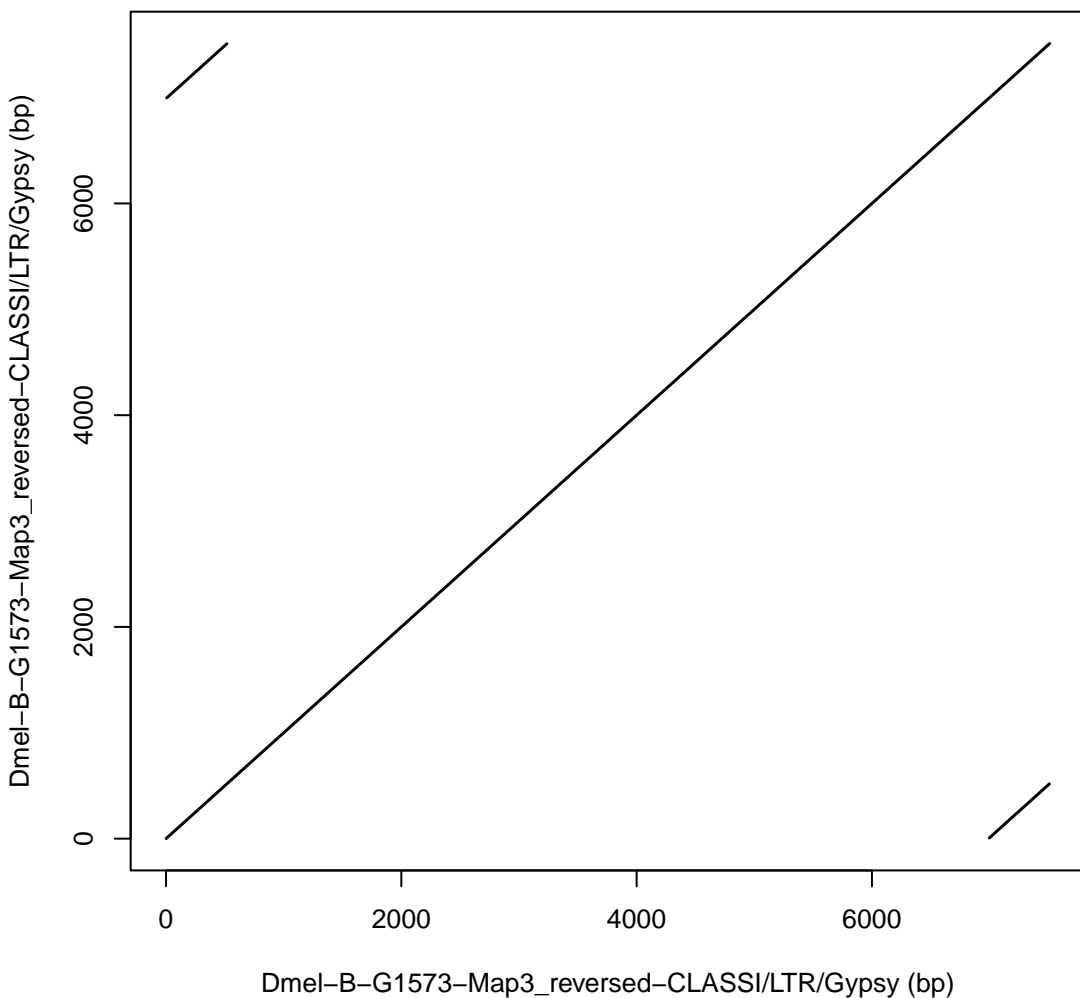
TE: Dmel-B-G1573-Map3_reversed-CLASSI/LTR/Gypsy
consensus size: 7511bp; fragments: 378; full length: 10 (>=6759.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

