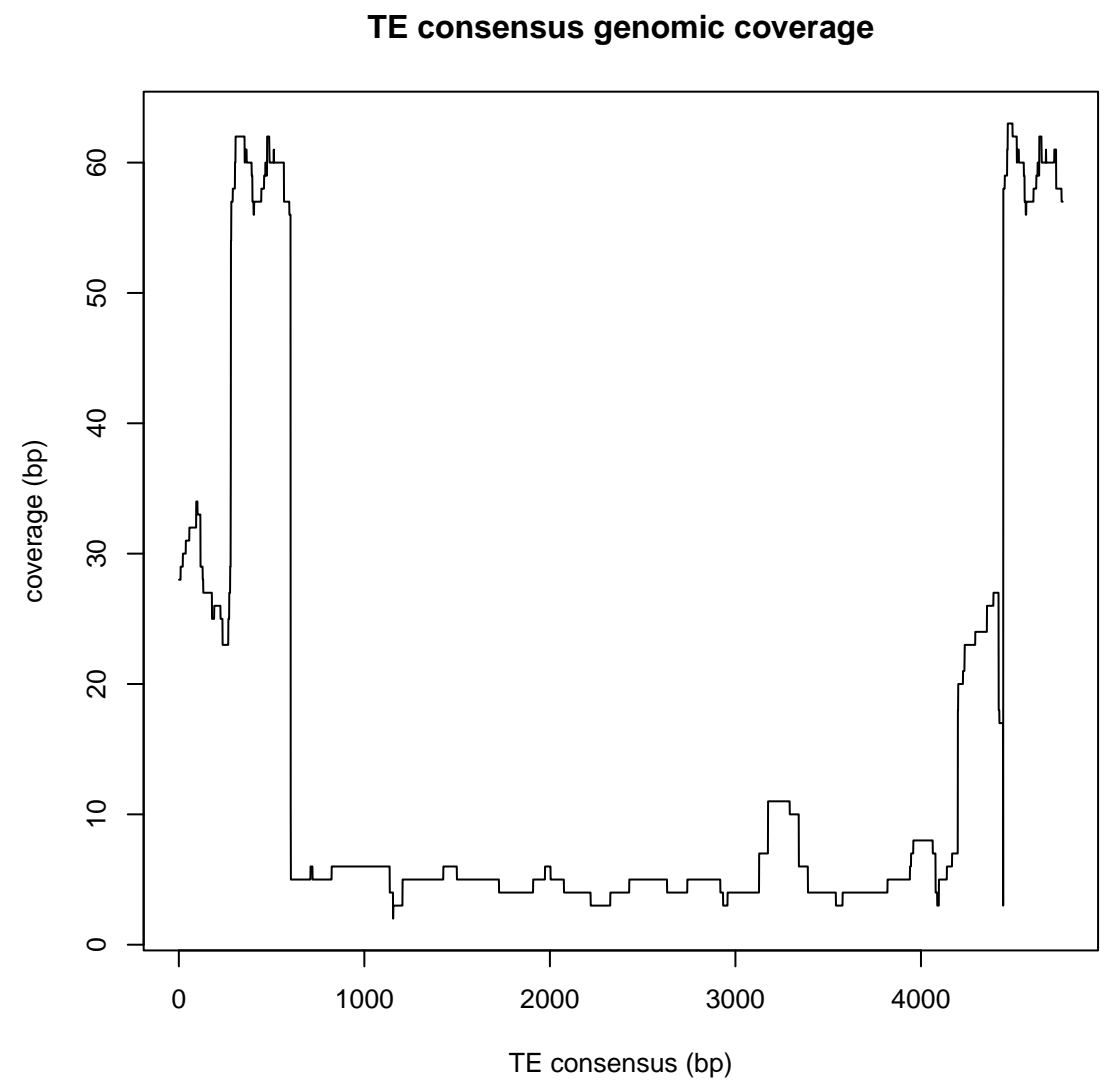
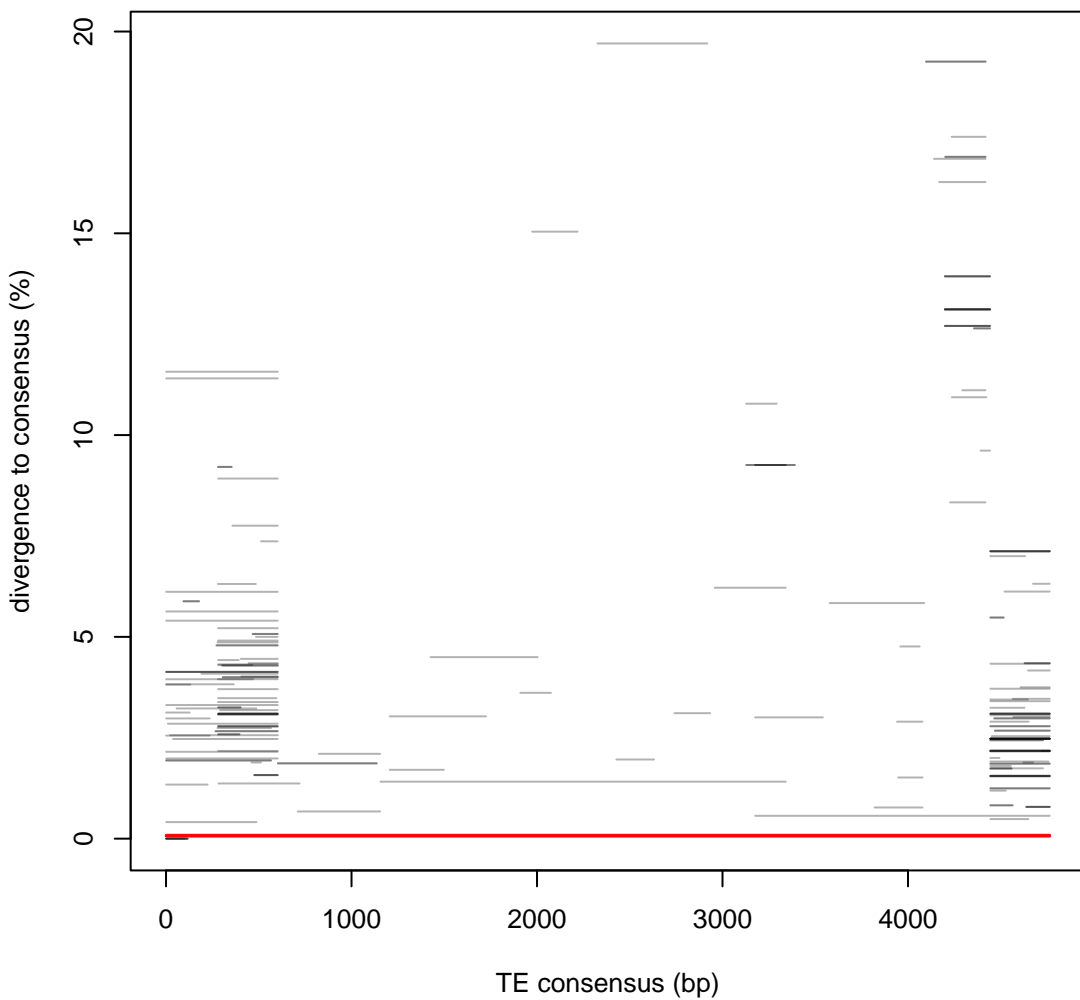
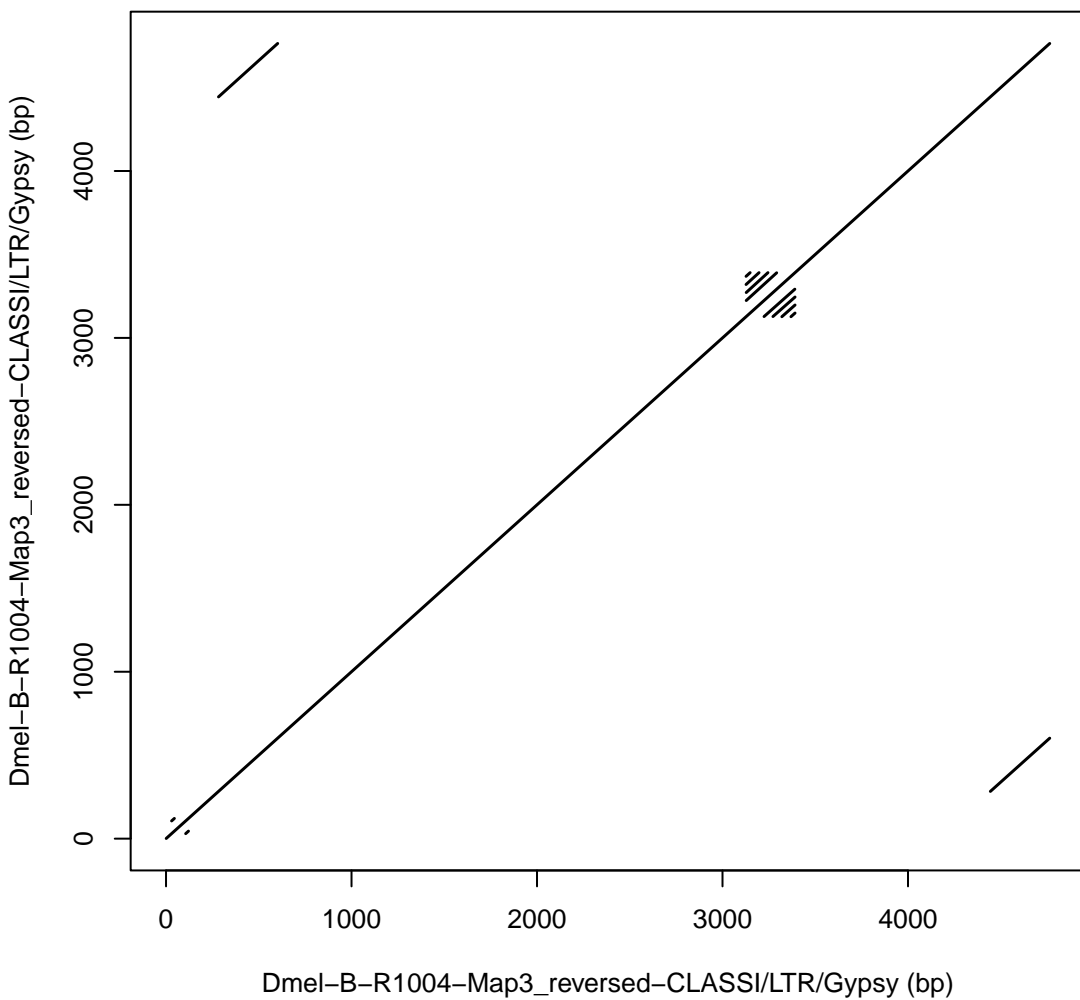


TE: Dmel-B-R1004-Map3_reversed-CLASSI/LTR/Gypsy
consensus size: 4764bp; fragments: 209; full length: 2 (>=4287.6bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

