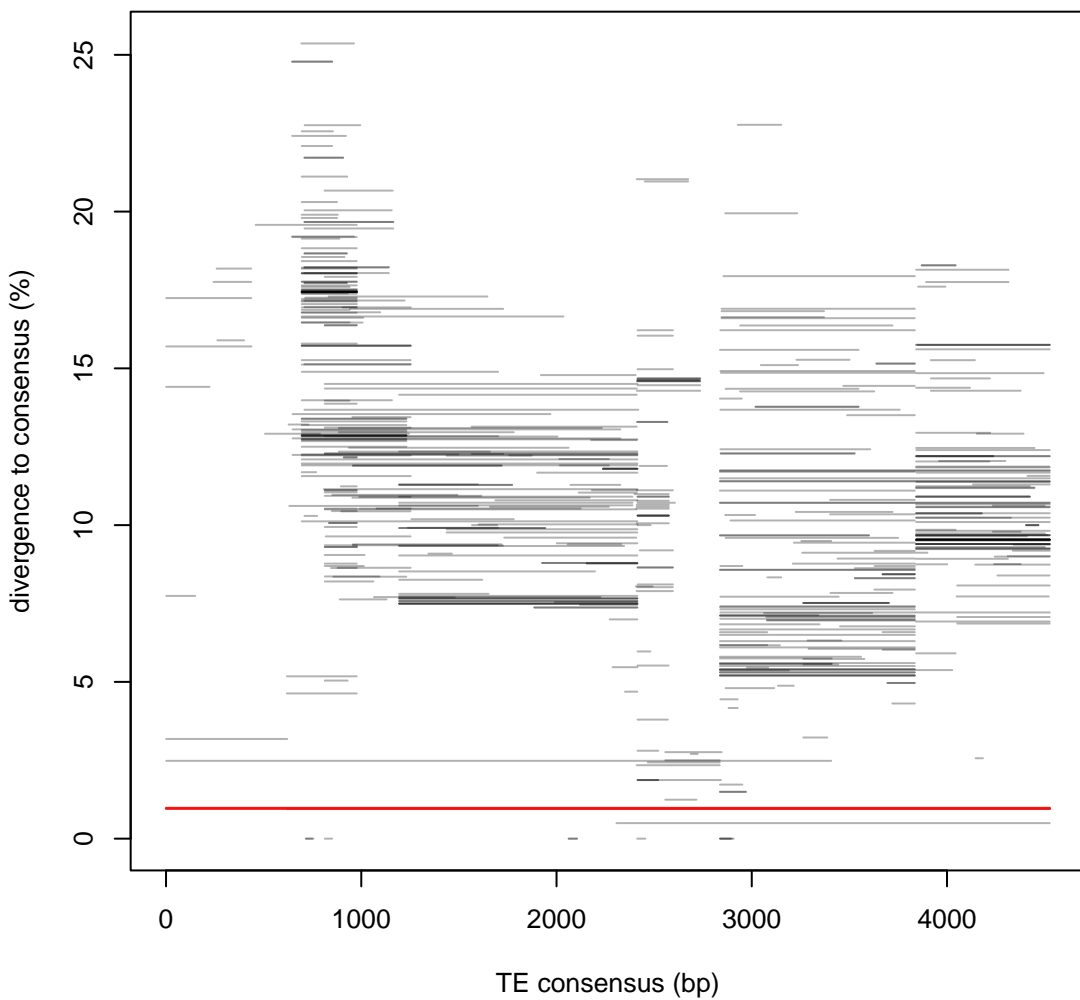
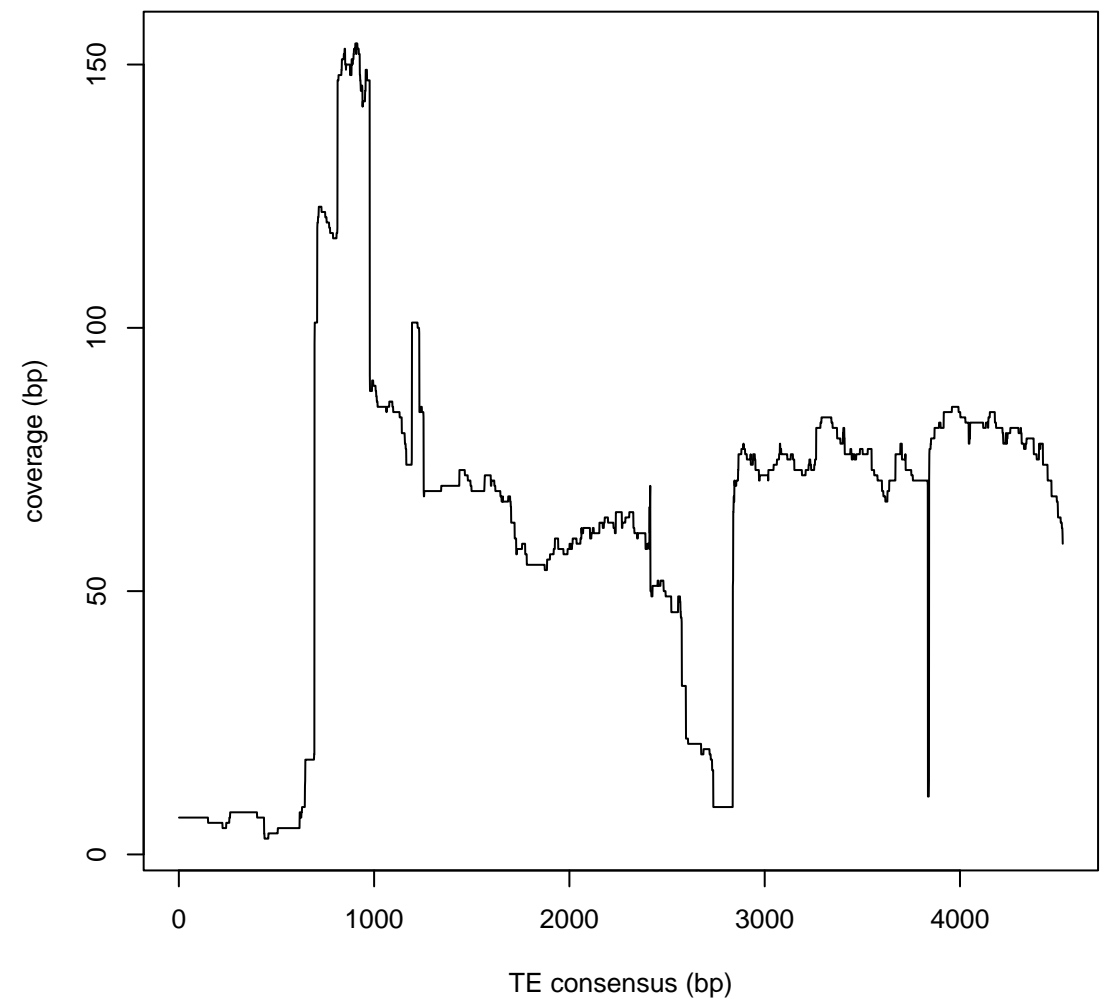


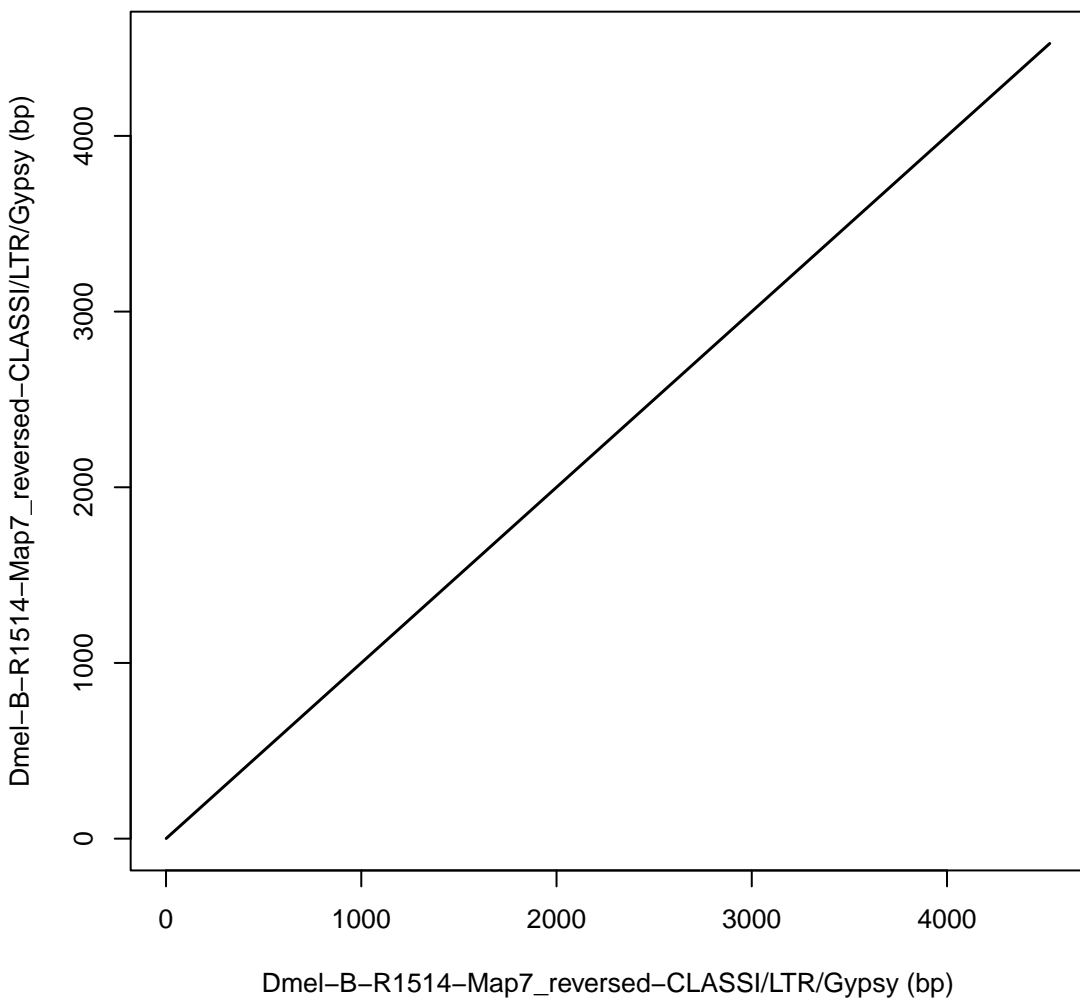
TE: Dmel-B-R1514-Map7\_reversed-CLASSI/LTR/Gypsy  
consensus size: 4526bp; fragments: 532; full length: 1 (>=4073.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

