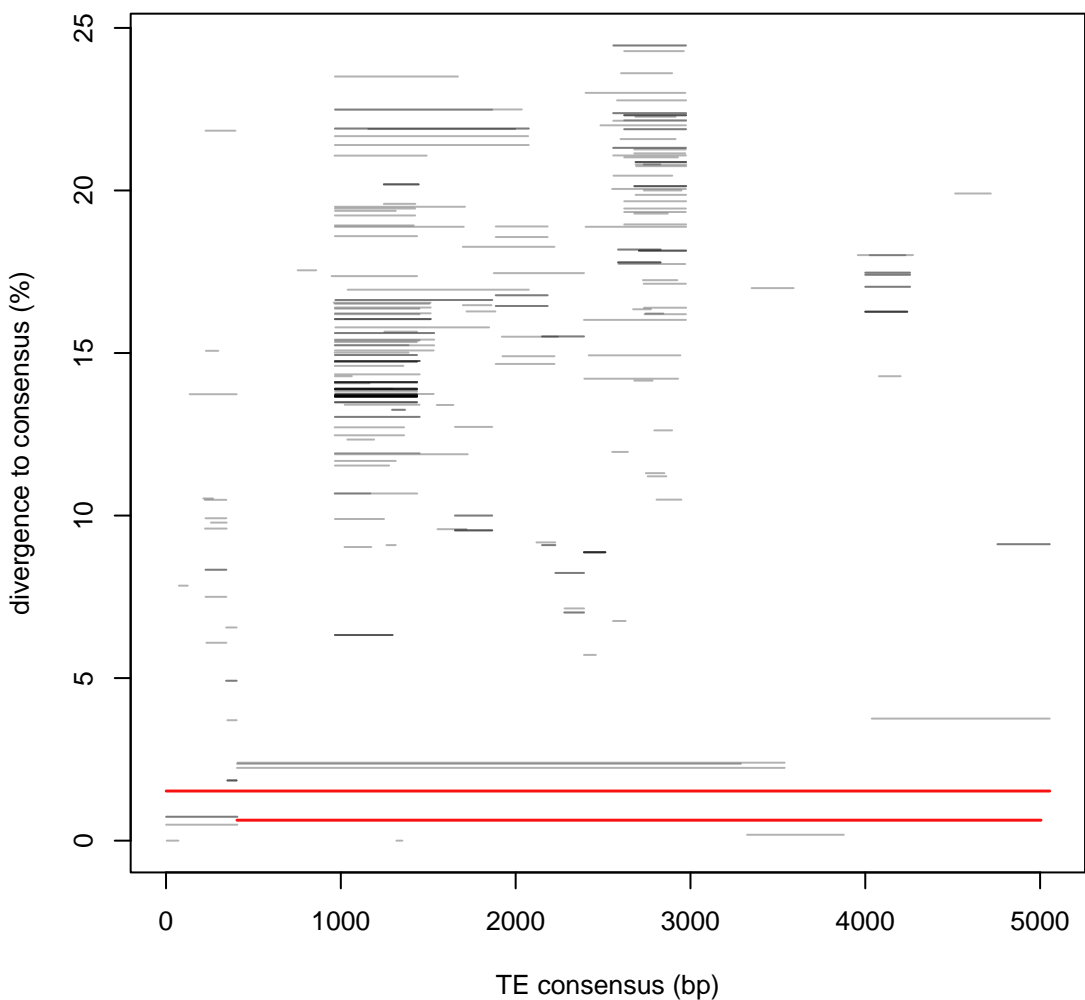
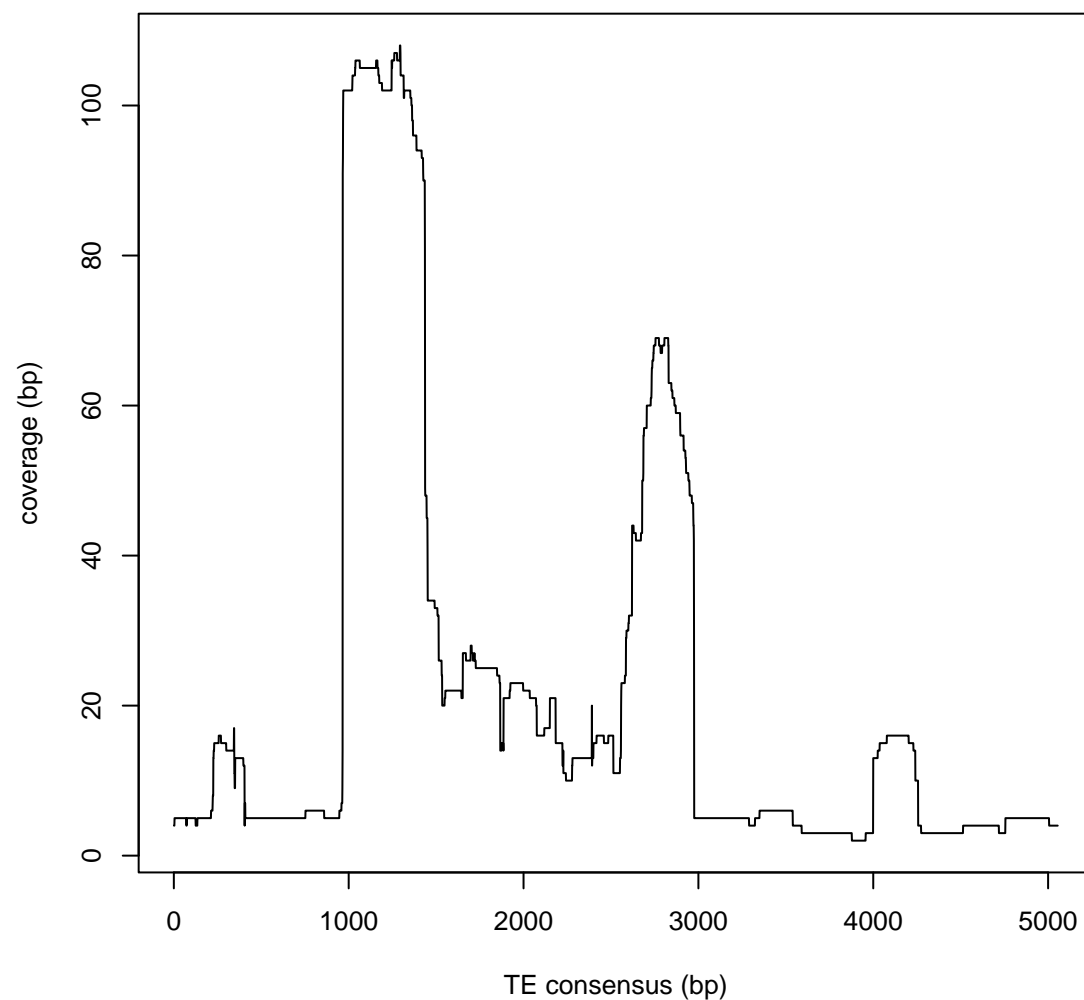


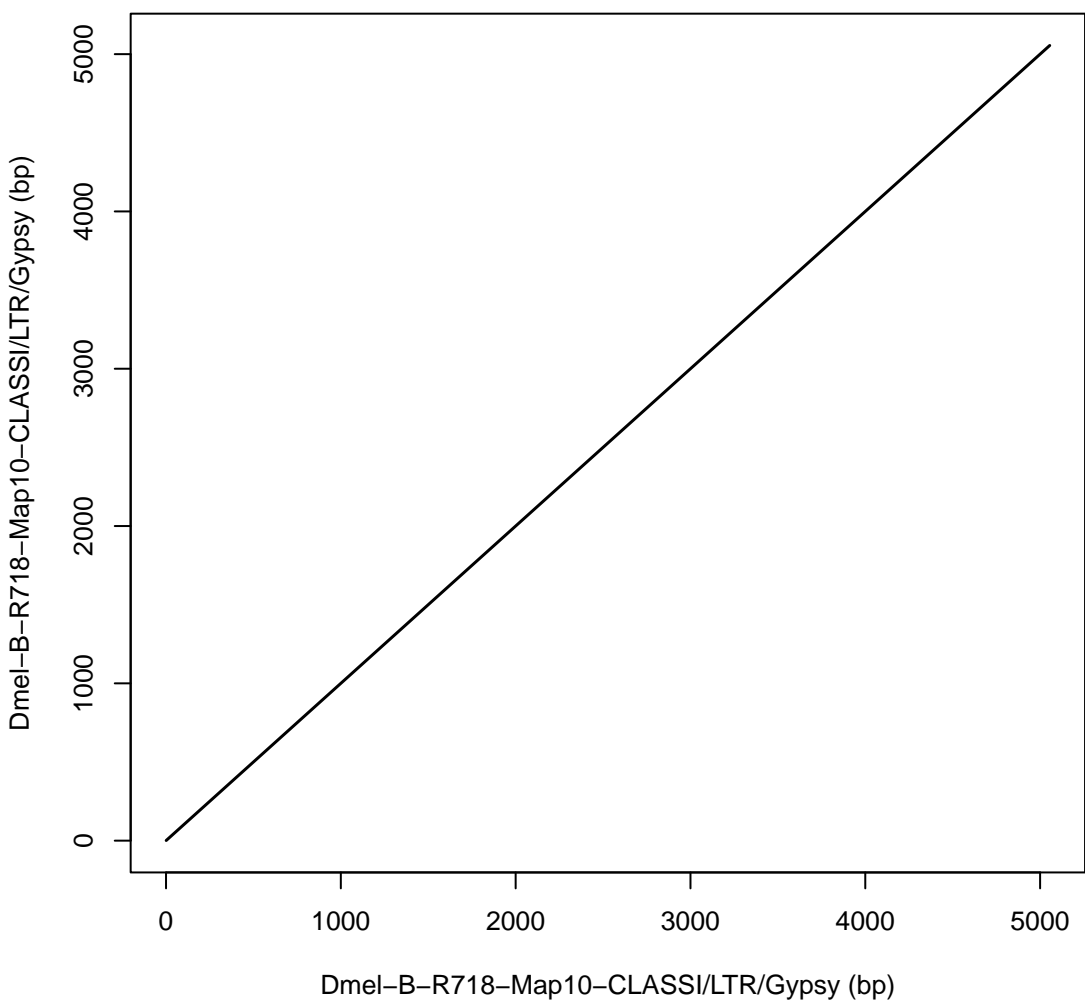
TE: Dmel-B-R718-Map10-CLASSI/LTR/Gypsy
consensus size: 5055bp; fragments: 265; full length: 2 (>=4549.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

