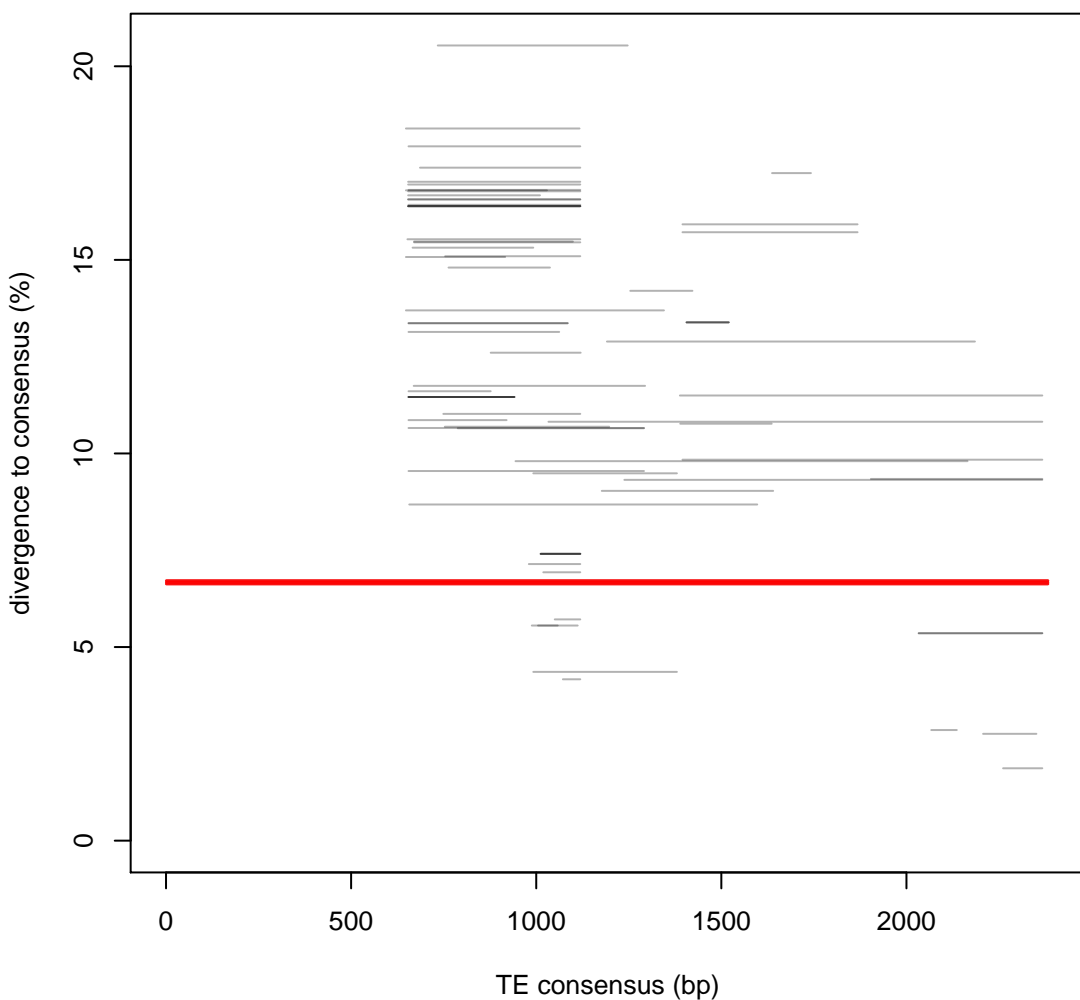
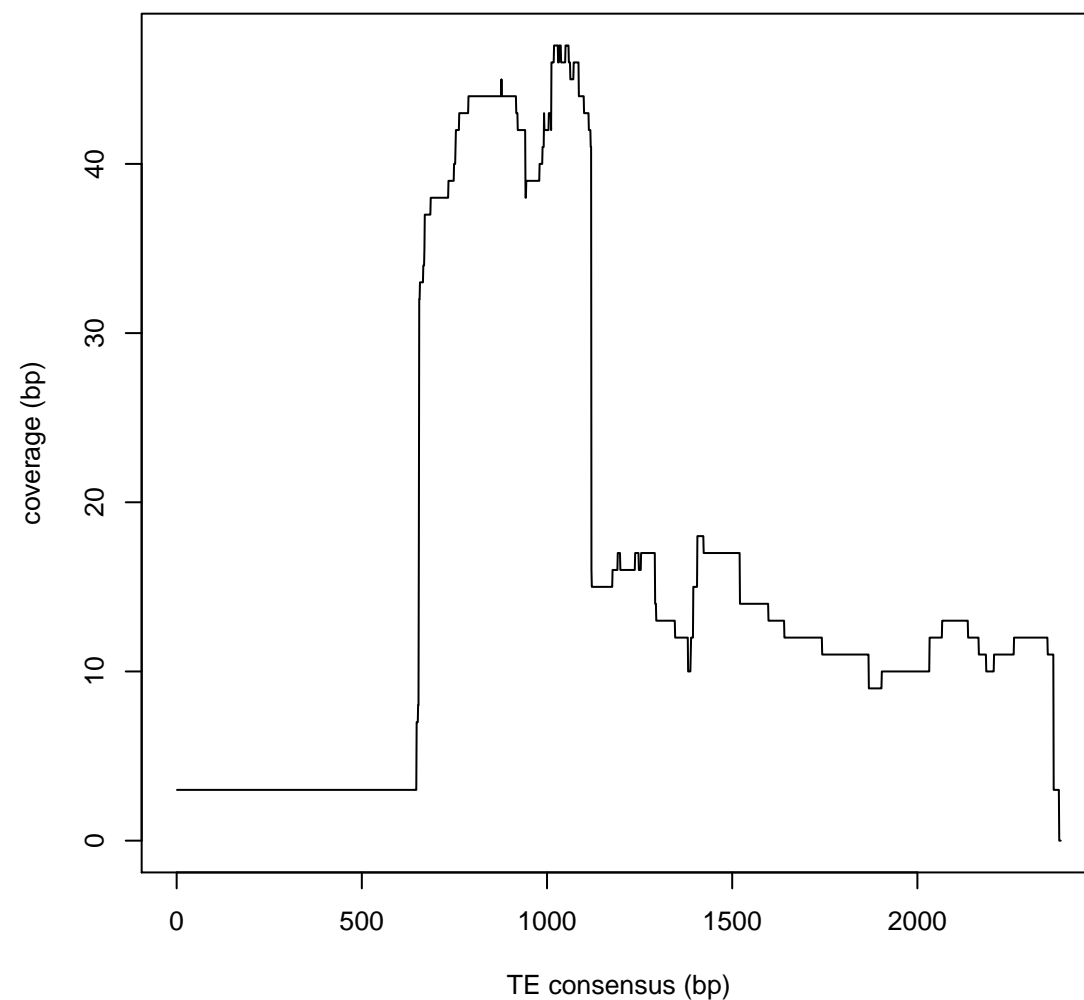


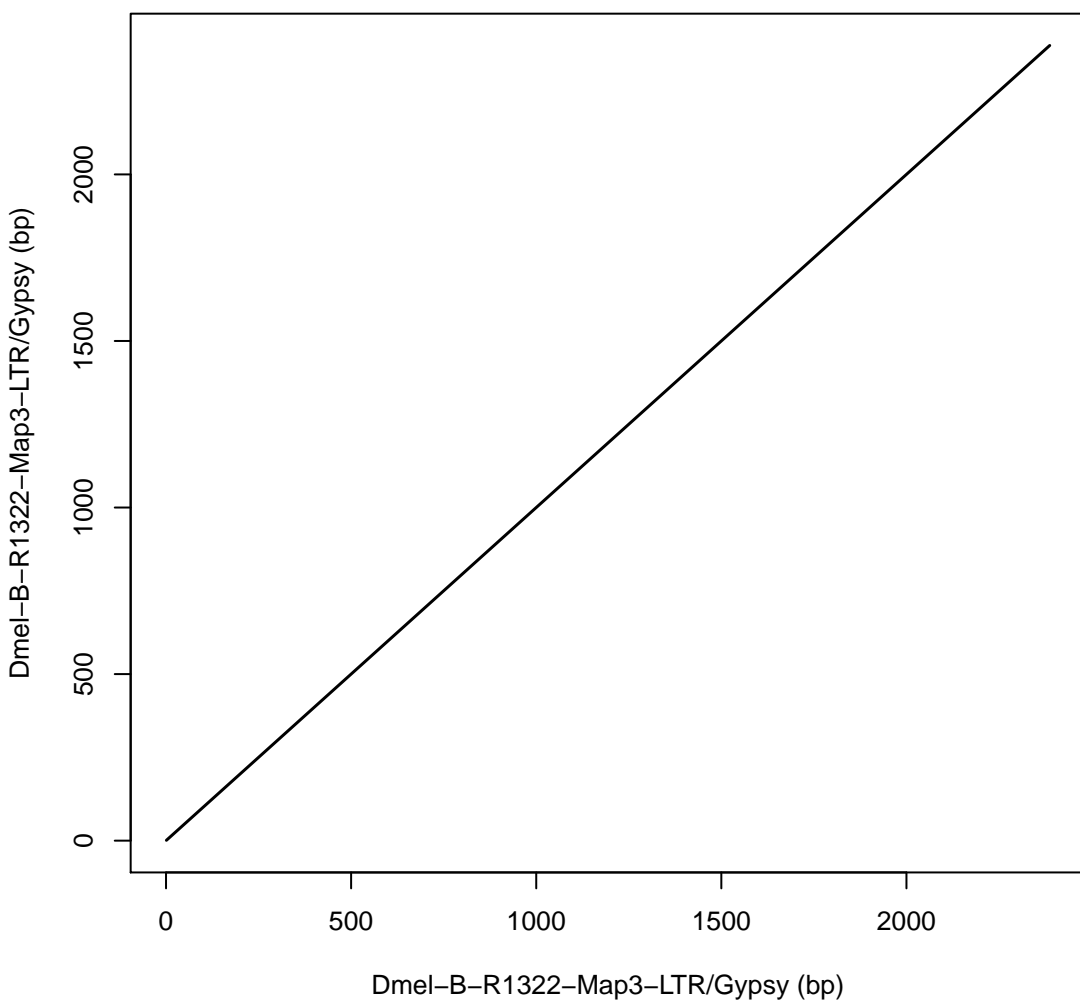
TE: Dmel-B-R1322-Map3-LTR/Gypsy  
consensus size: 2387bp; fragments: 78; full length: 3 (>=2148.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

