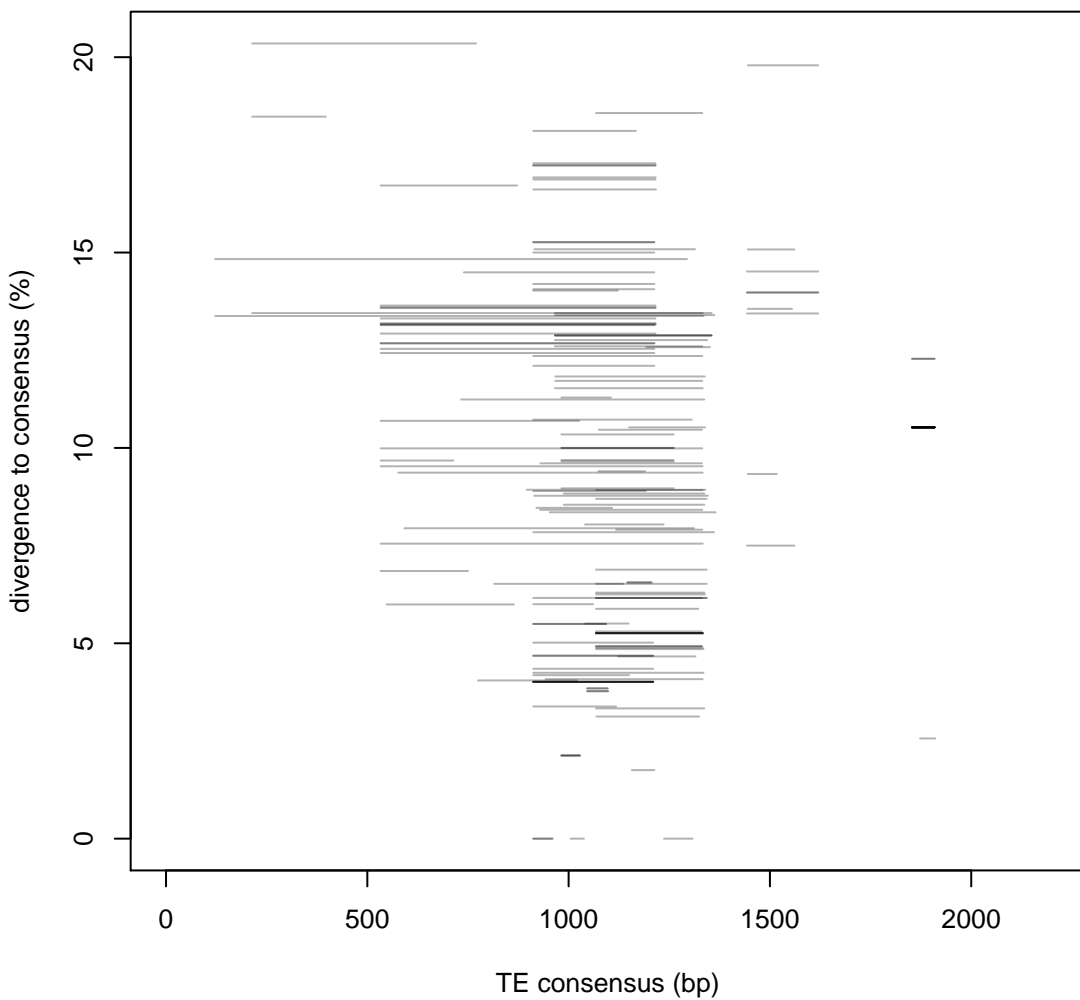
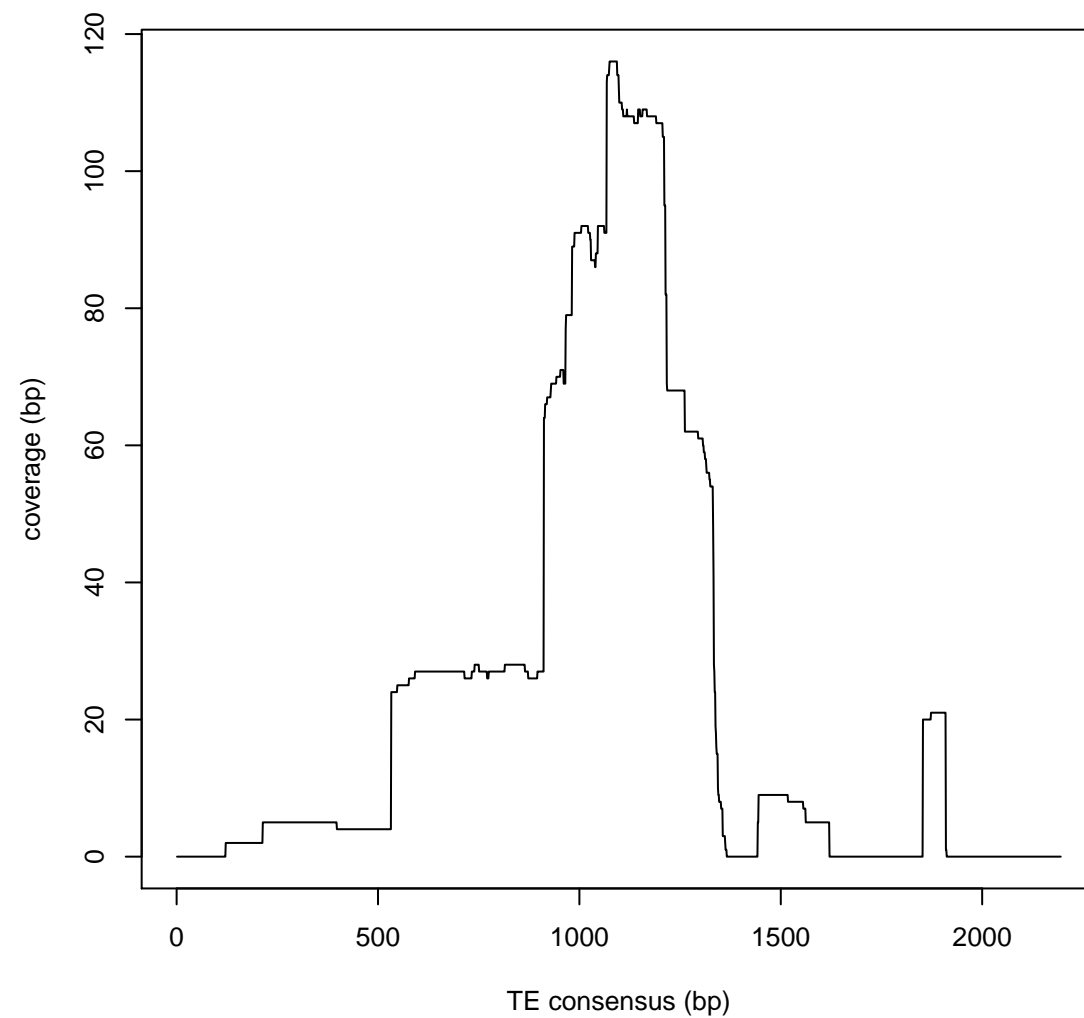


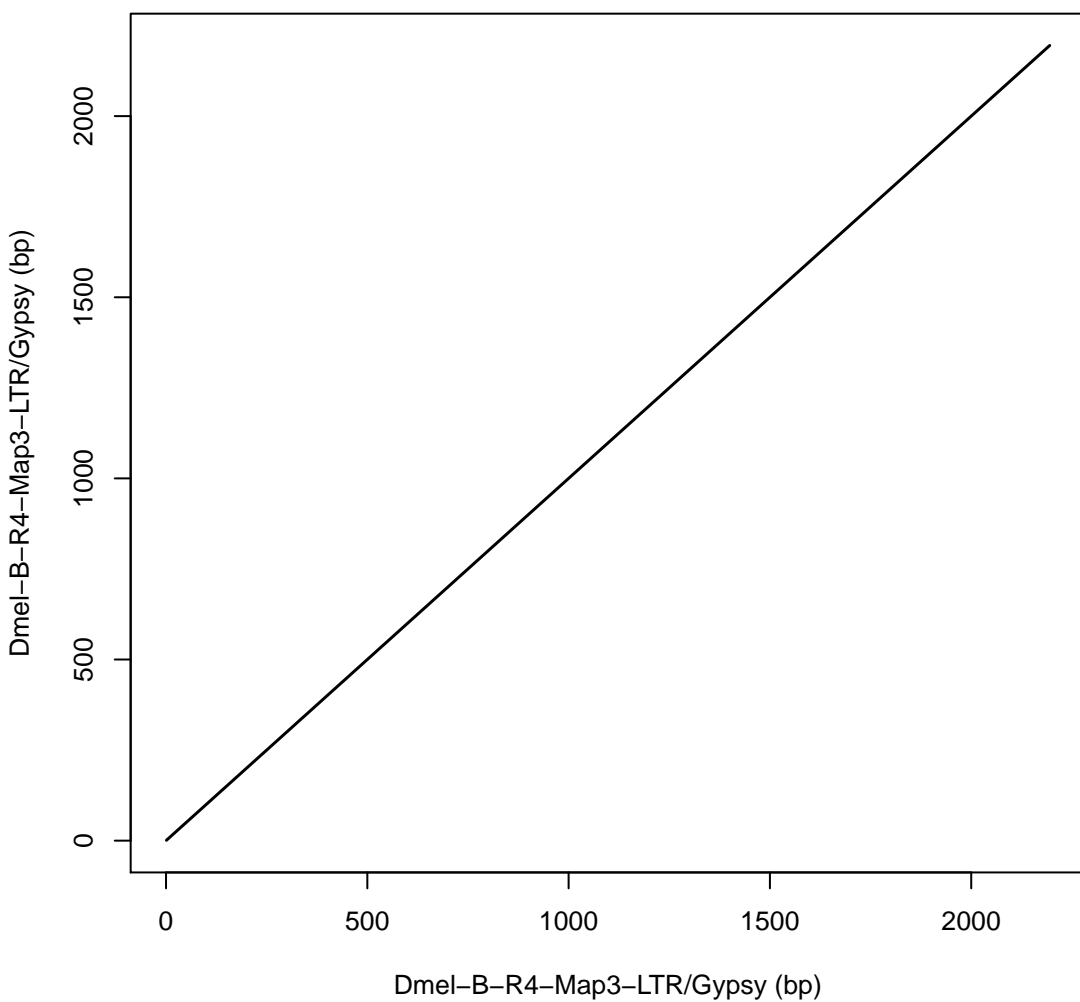
TE: Dmel-B-R4-Map3-LTR/Gypsy
consensus size: 2195bp; fragments: 169; full length: 0 (>=1975.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

