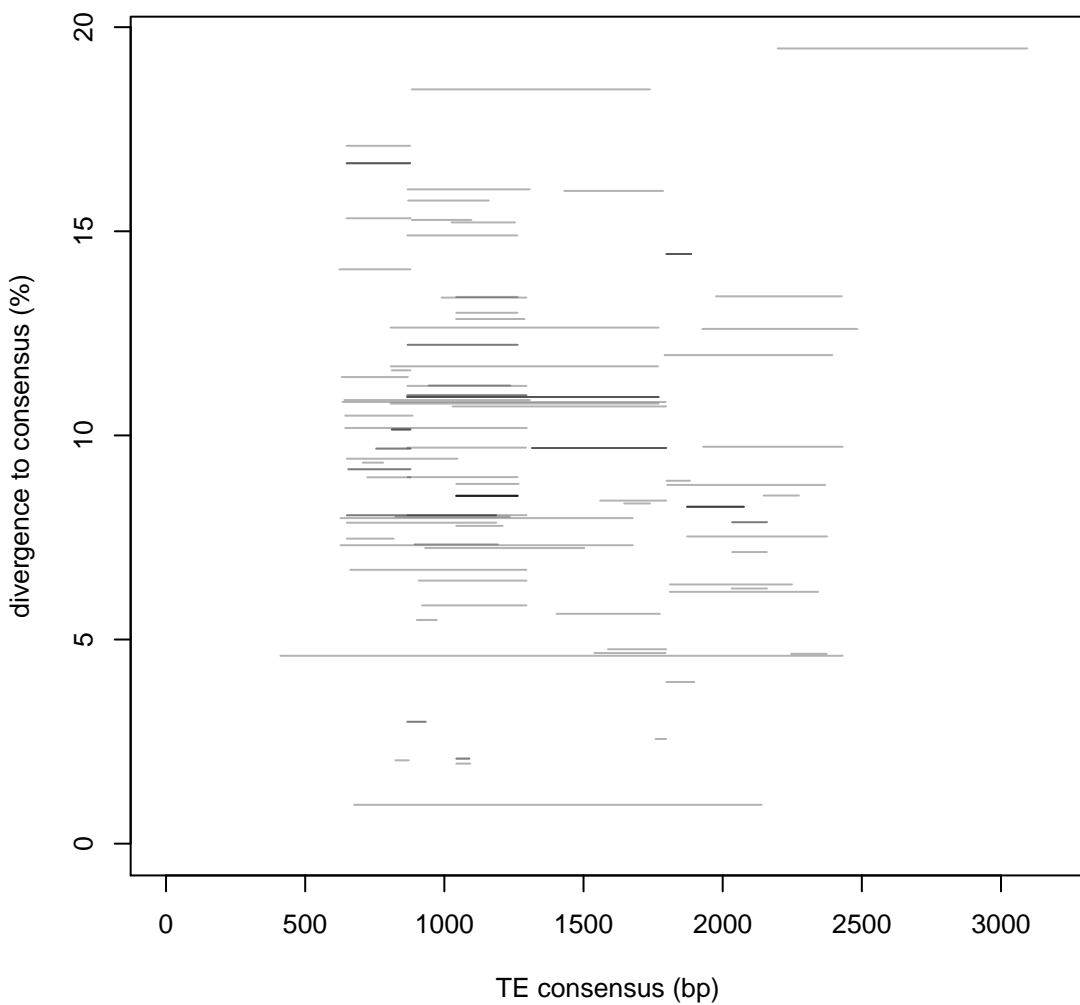
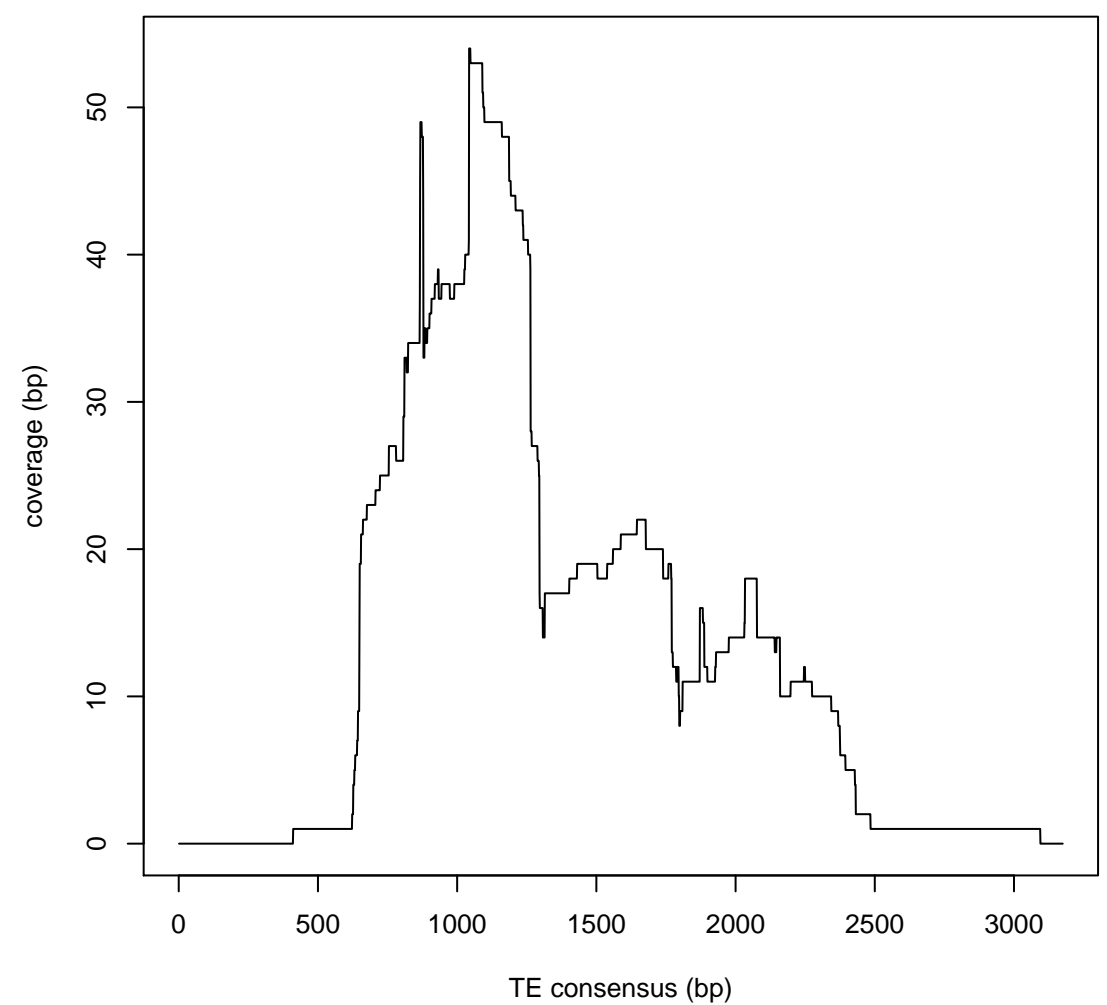


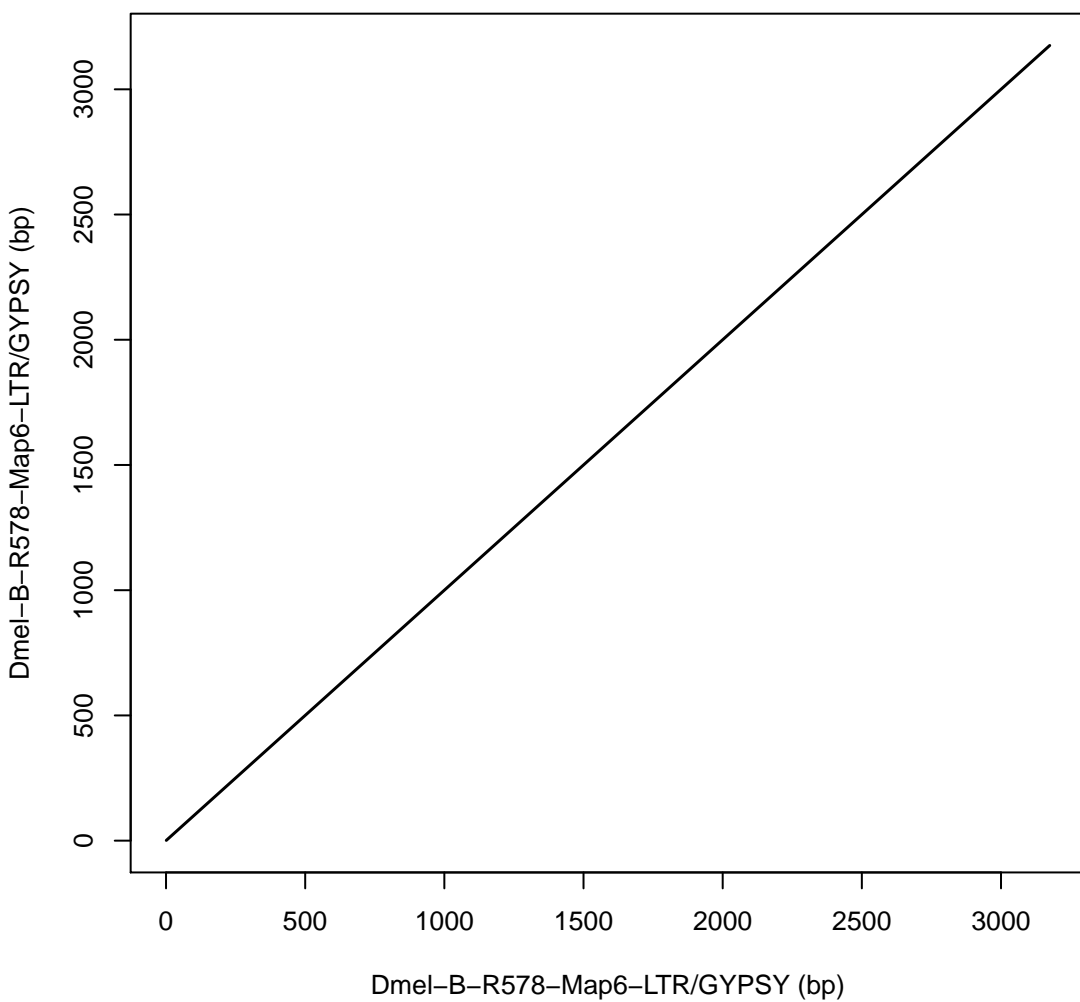
TE: Dmel-B-R578-Map6-LTR/GYPSY
consensus size: 3175bp; fragments: 111; full length: 0 (>=2857.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

