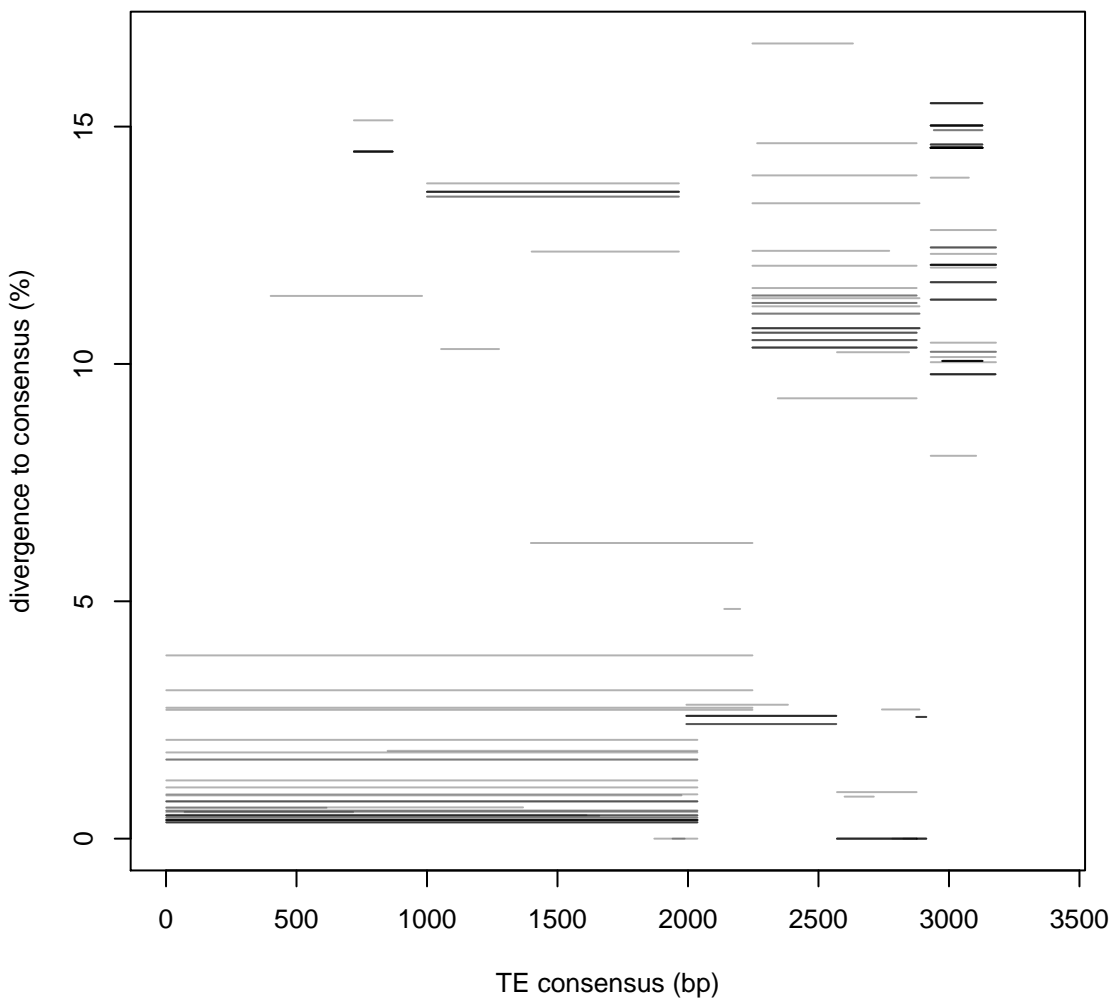
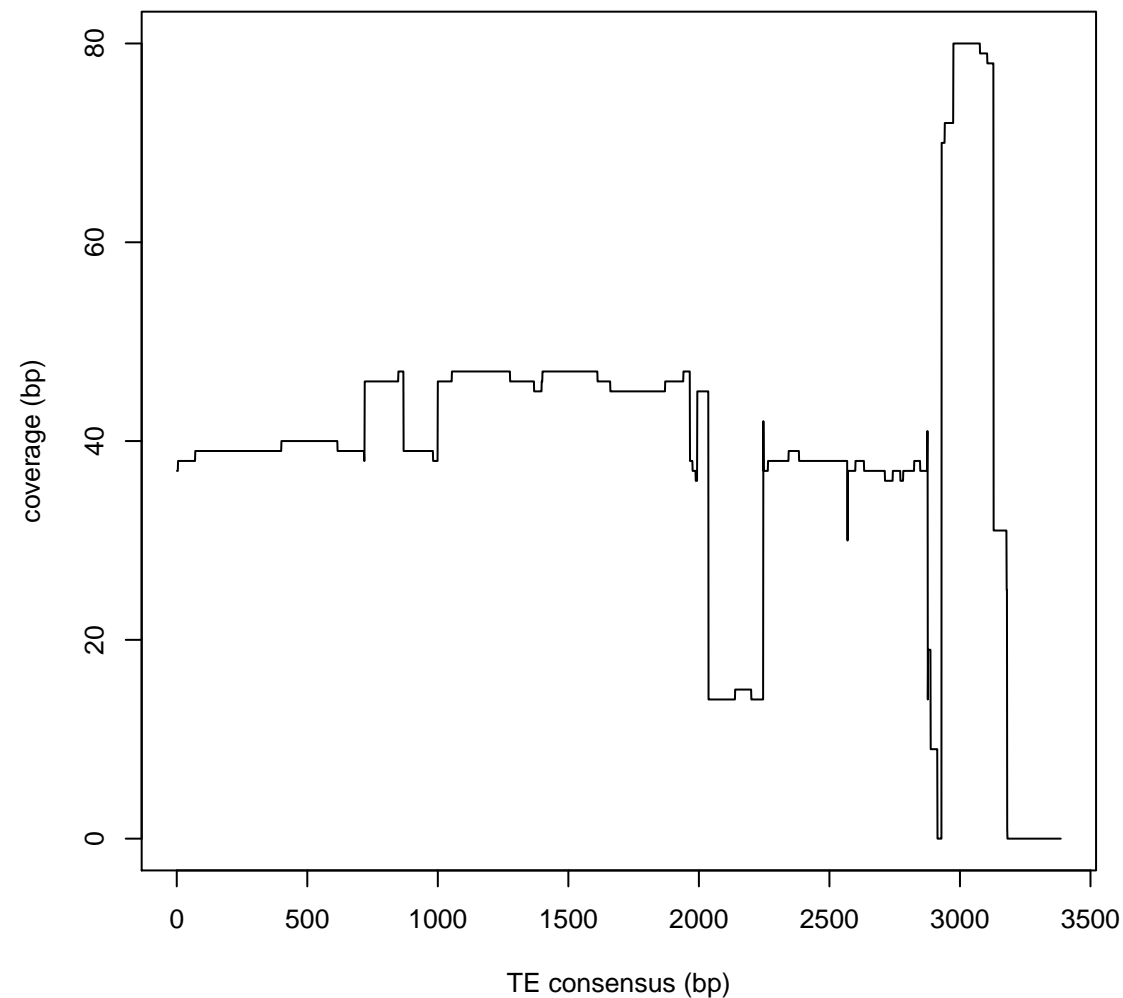


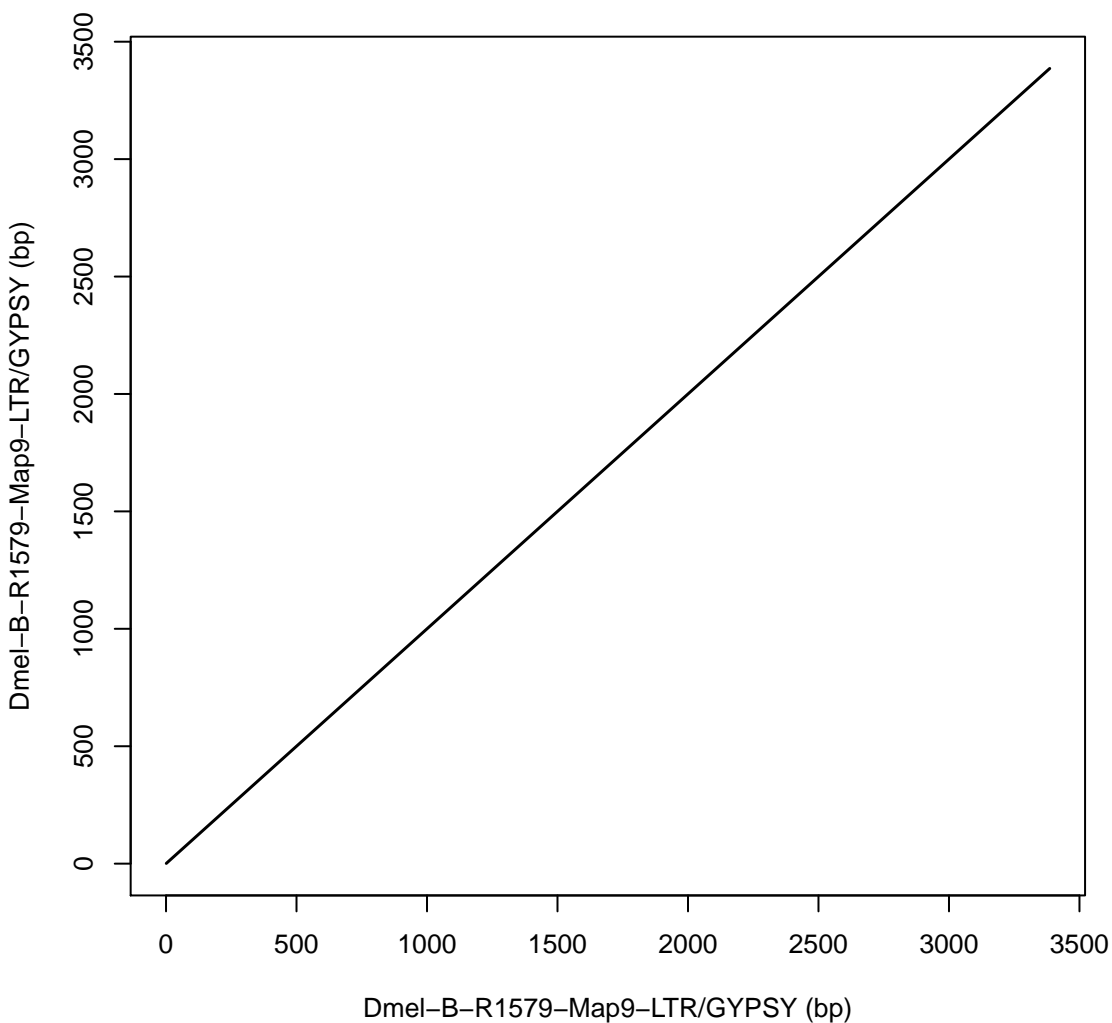
TE: Dmel-B-R1579-Map9-LTR/Gypsy  
consensus size: 3386bp; fragments: 202; full length: 0 (>=3047.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

