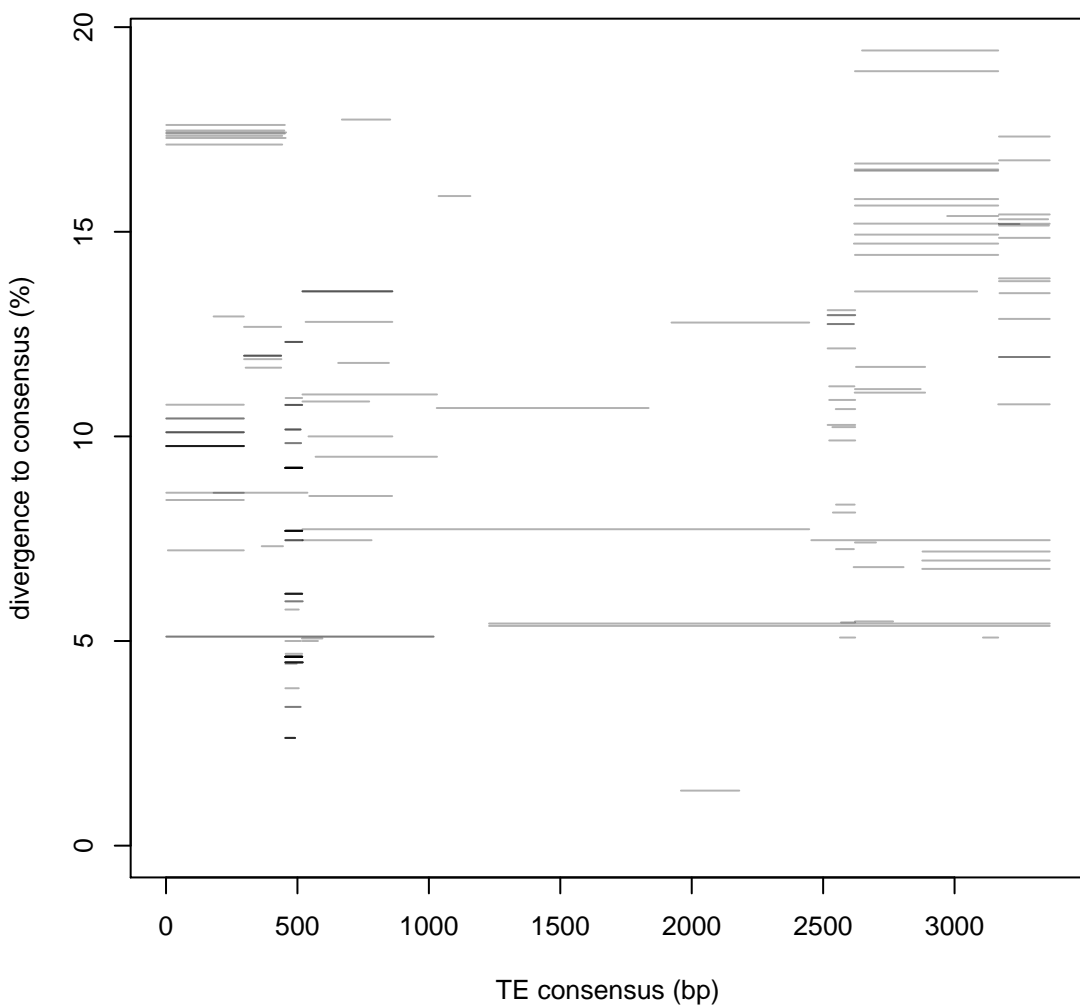
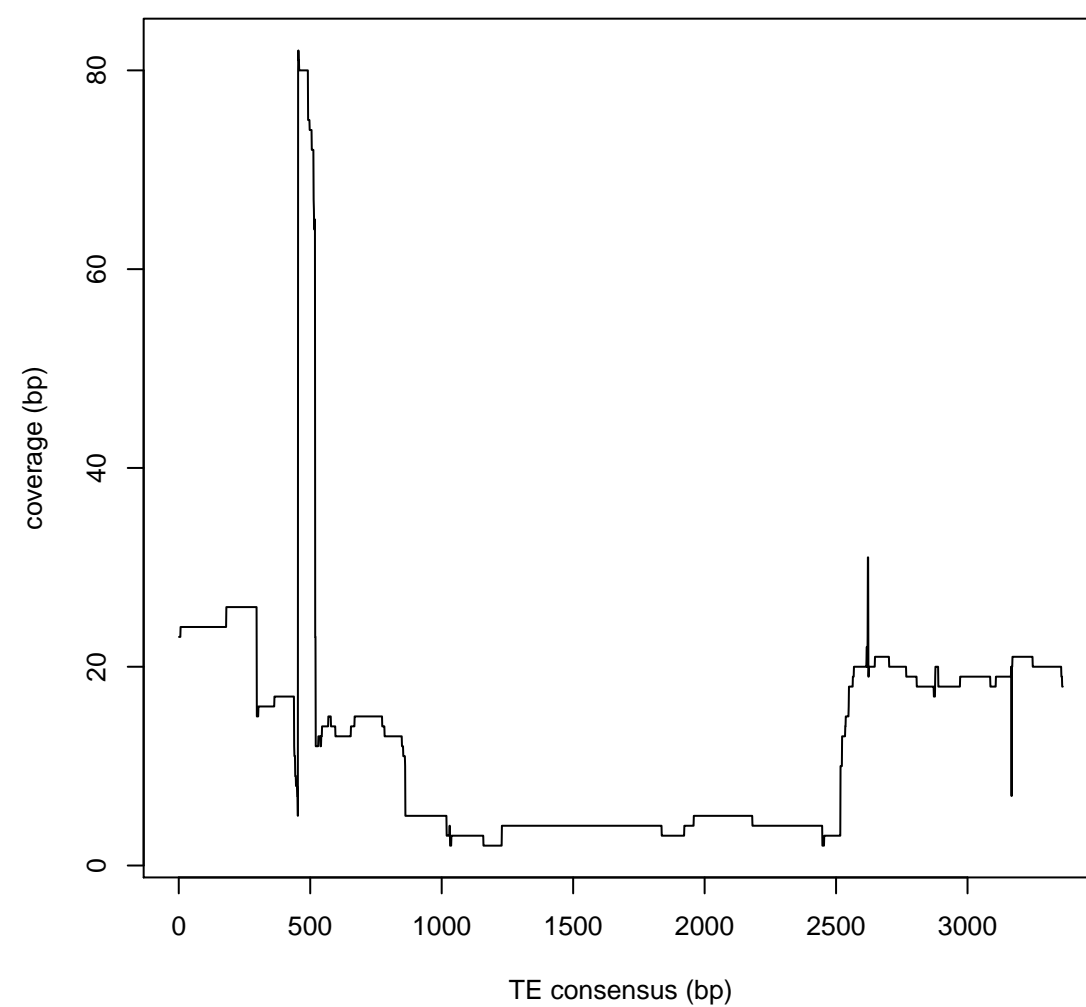


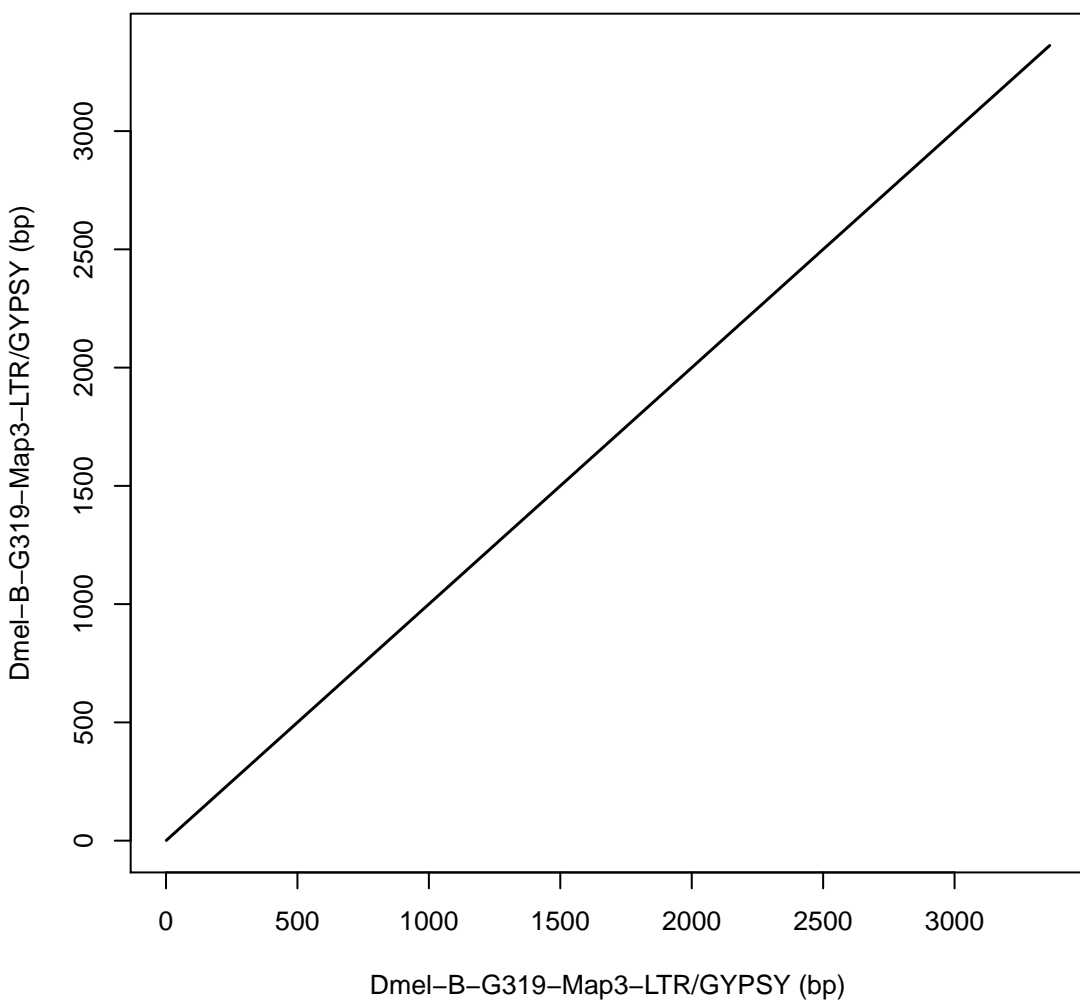
TE: Dmel-B-G319-Map3-LTR/GYPSY  
consensus size: 3362bp; fragments: 187; full length: 0 (>=3025.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

