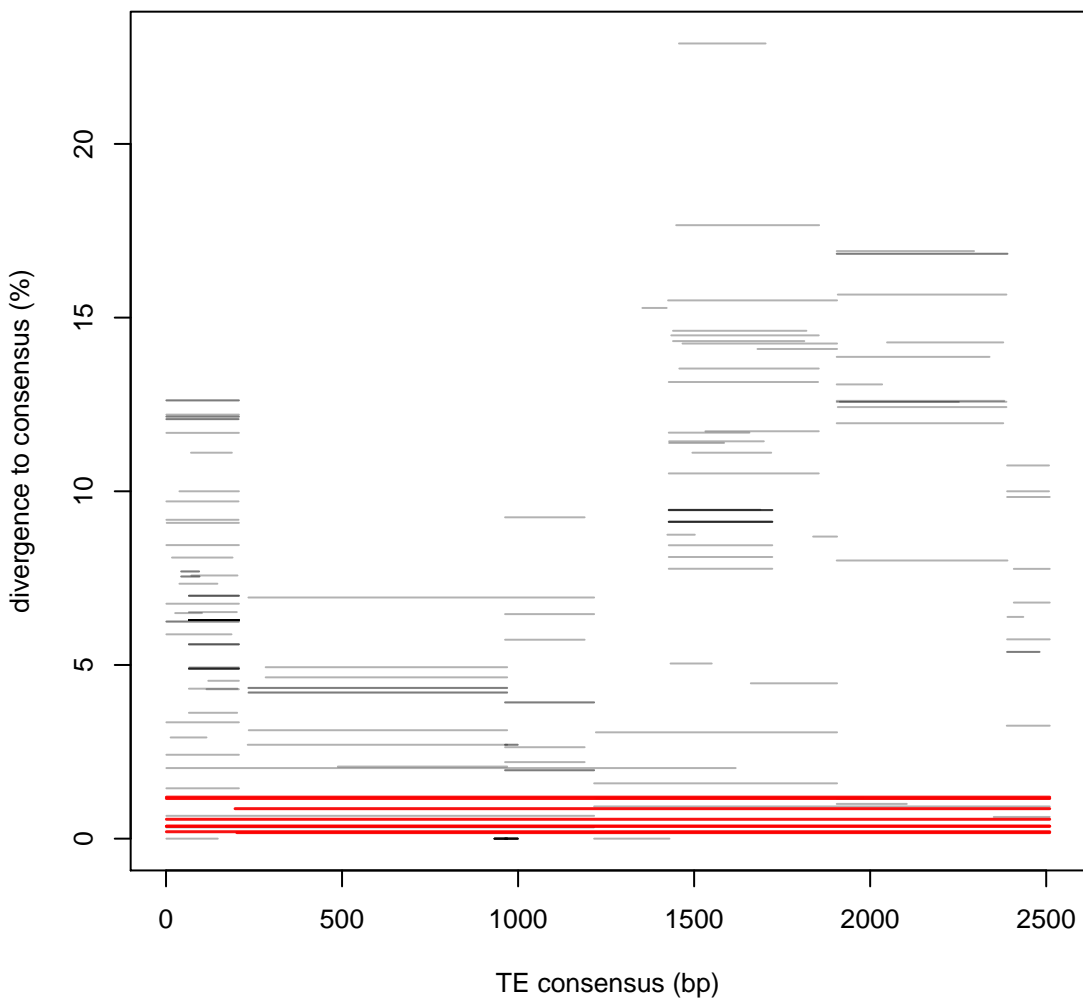
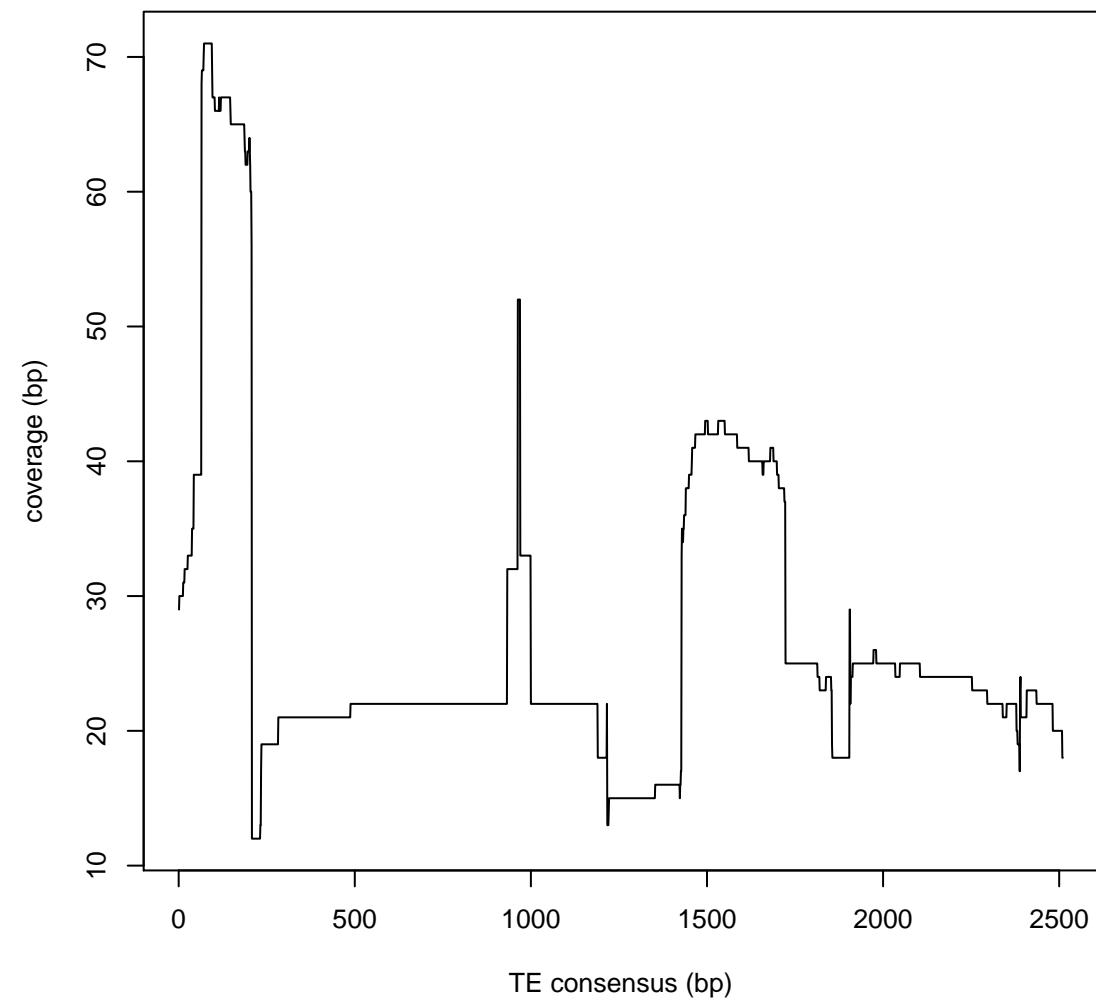


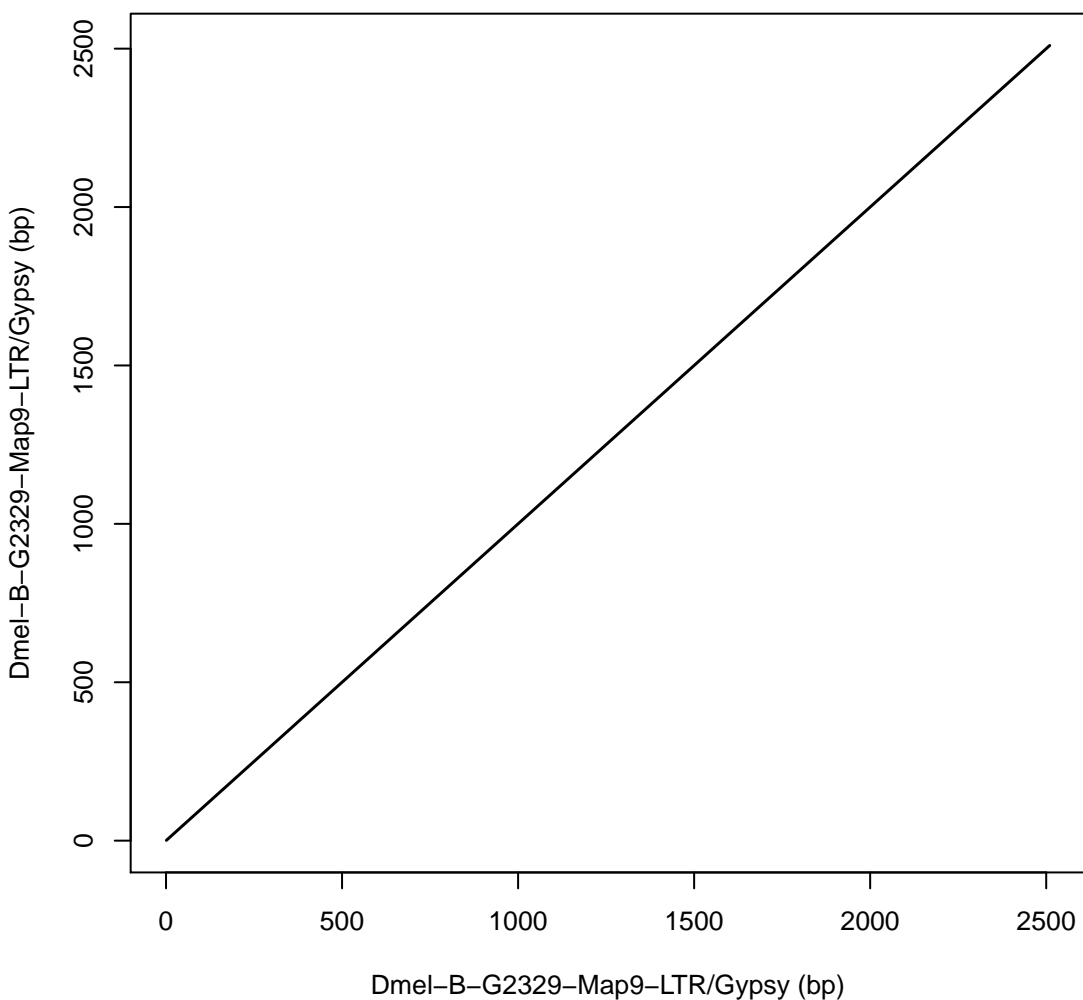
TE: Dmel-B-G2329-Map9-LTR/Gypsy  
consensus size: 2510bp; fragments: 182; full length: 8 (>=2259bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

