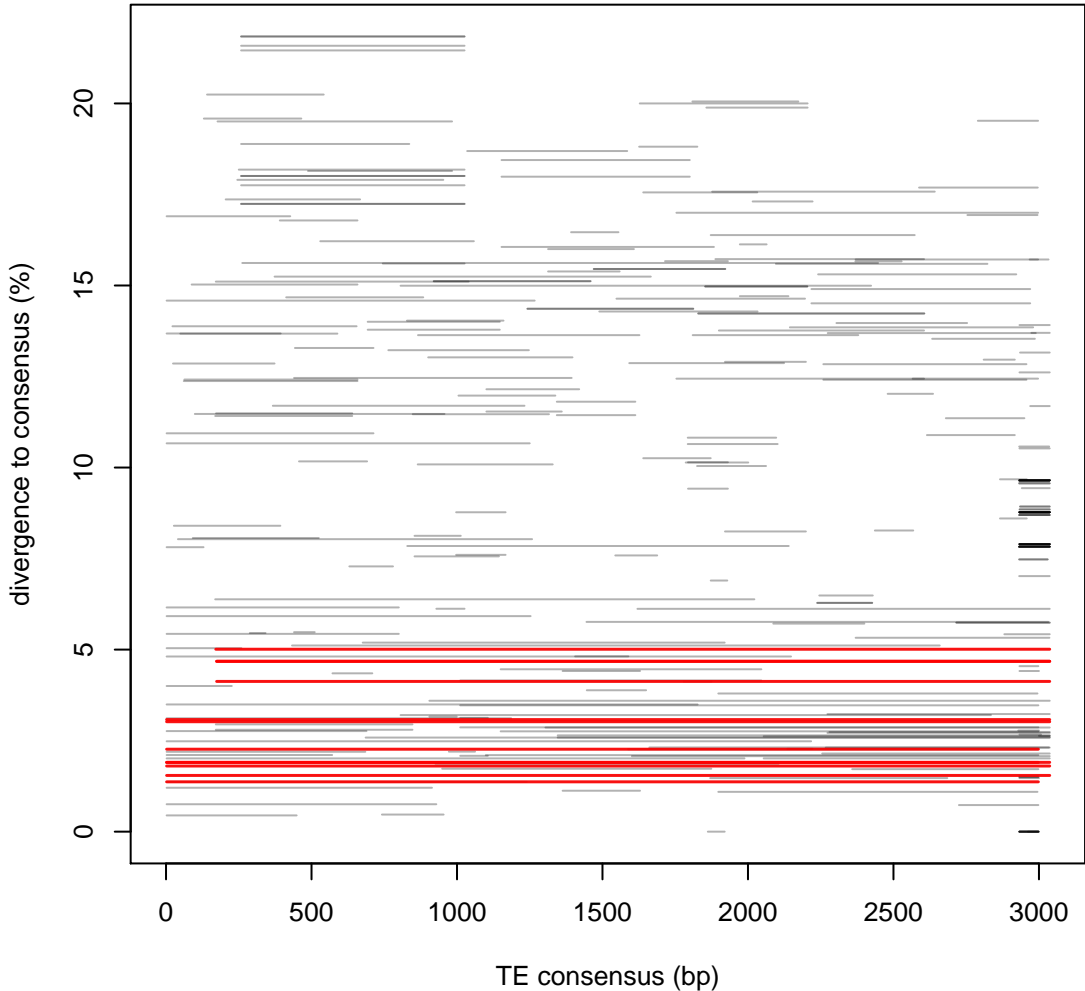
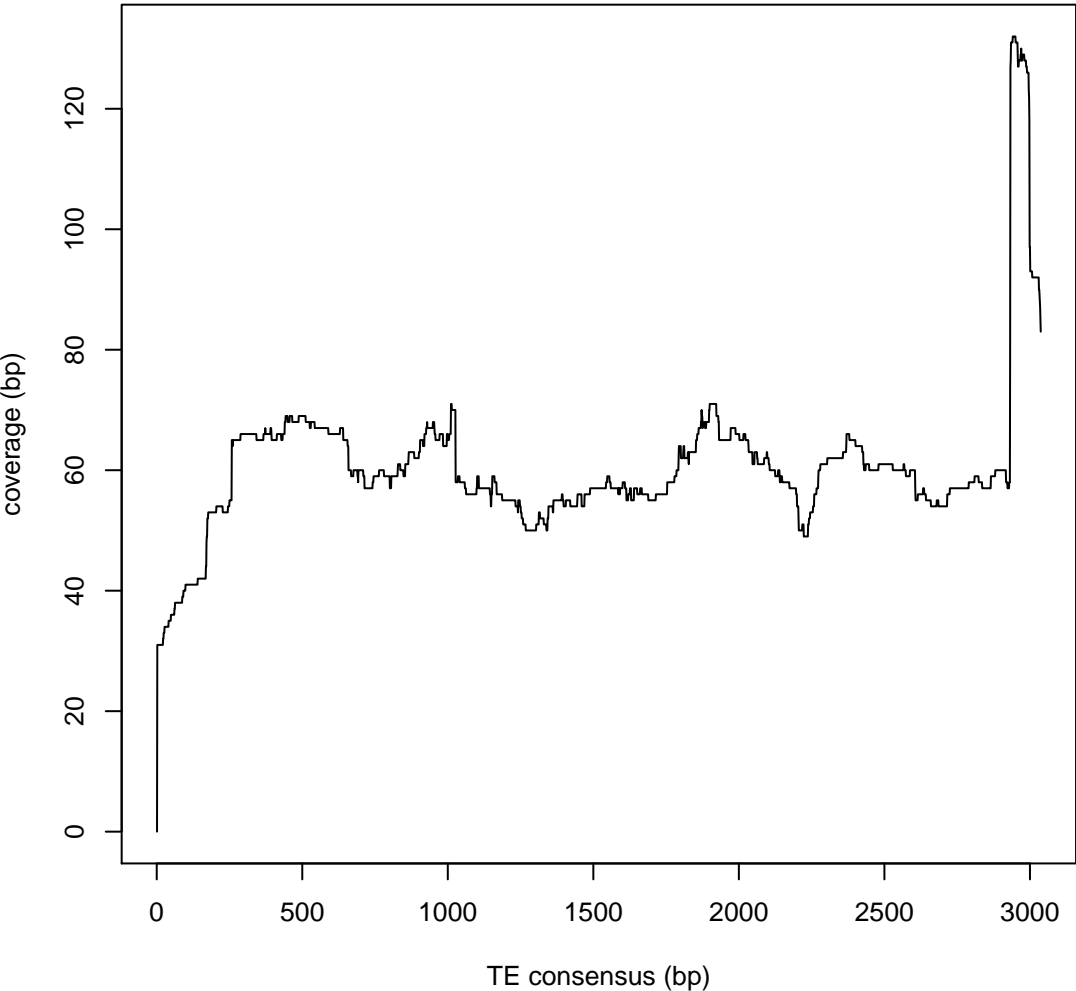


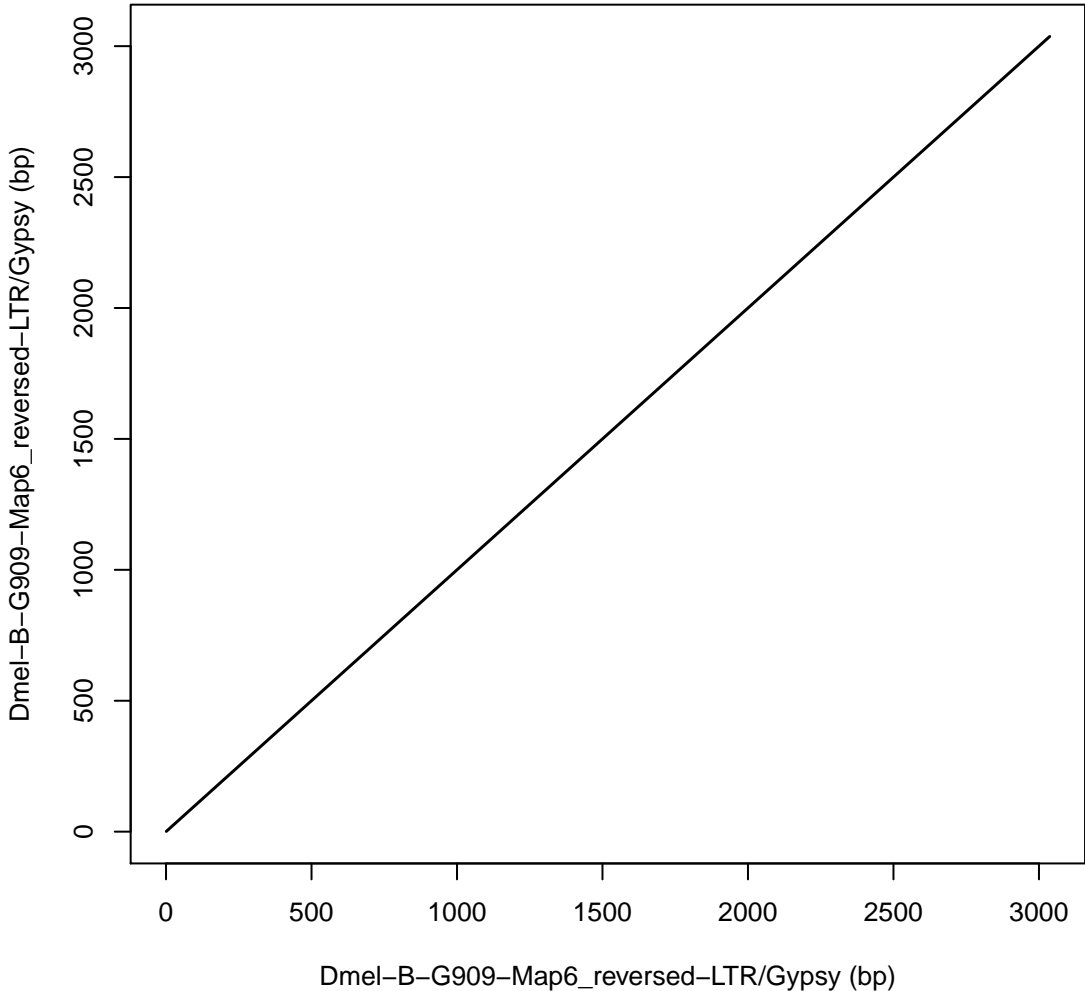
TE: Dmel-B-G909-Map6\_reversed-LTR/Gypsy  
consensus size: 3037bp; fragments: 305; full length: 12 (>=2733.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

