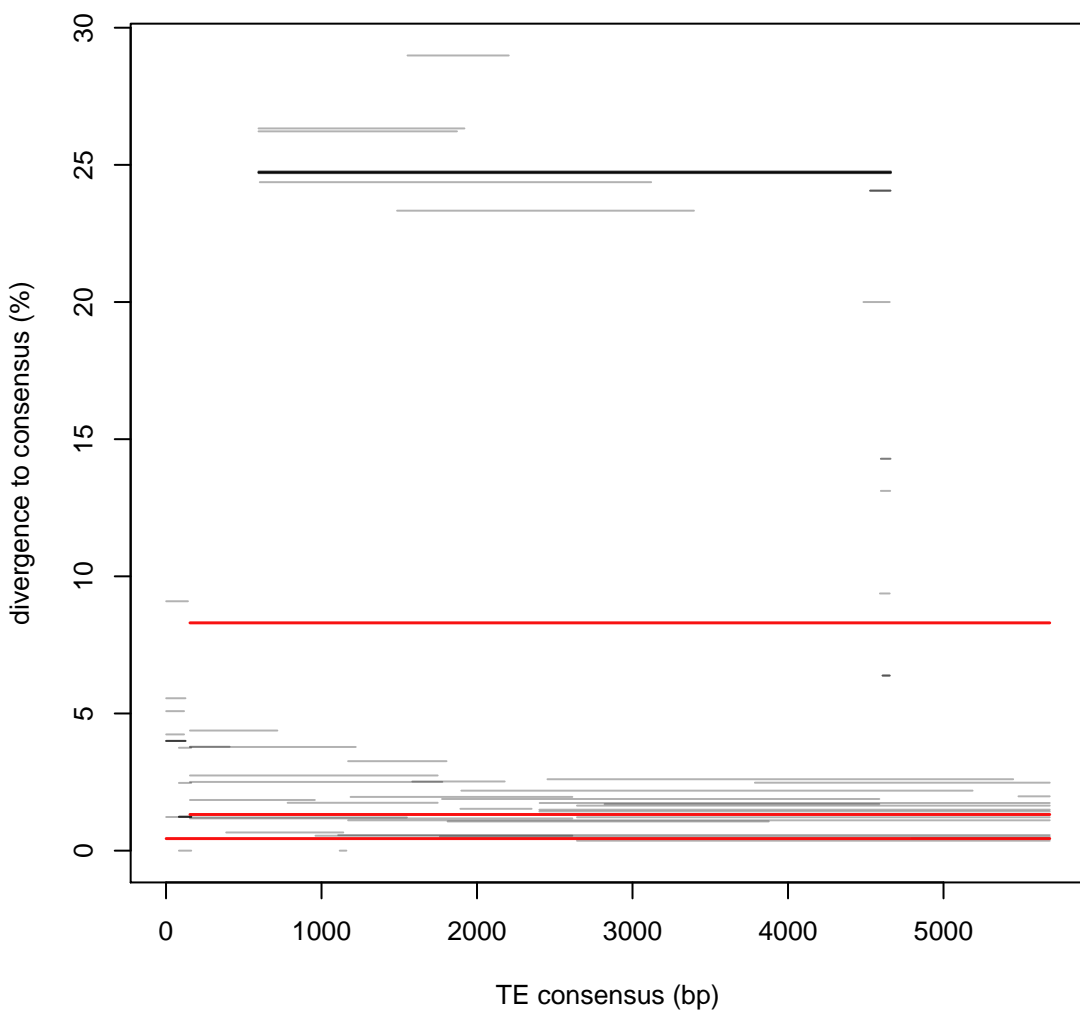
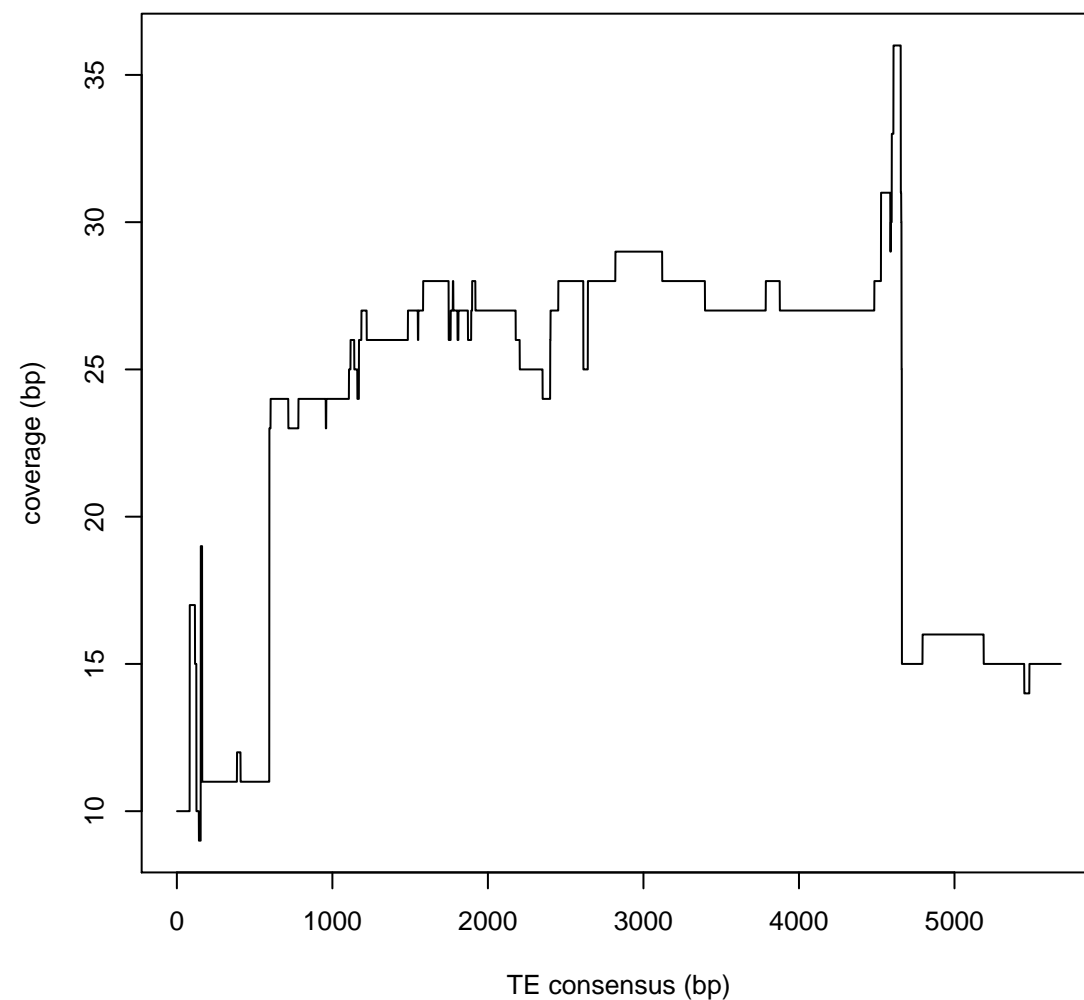


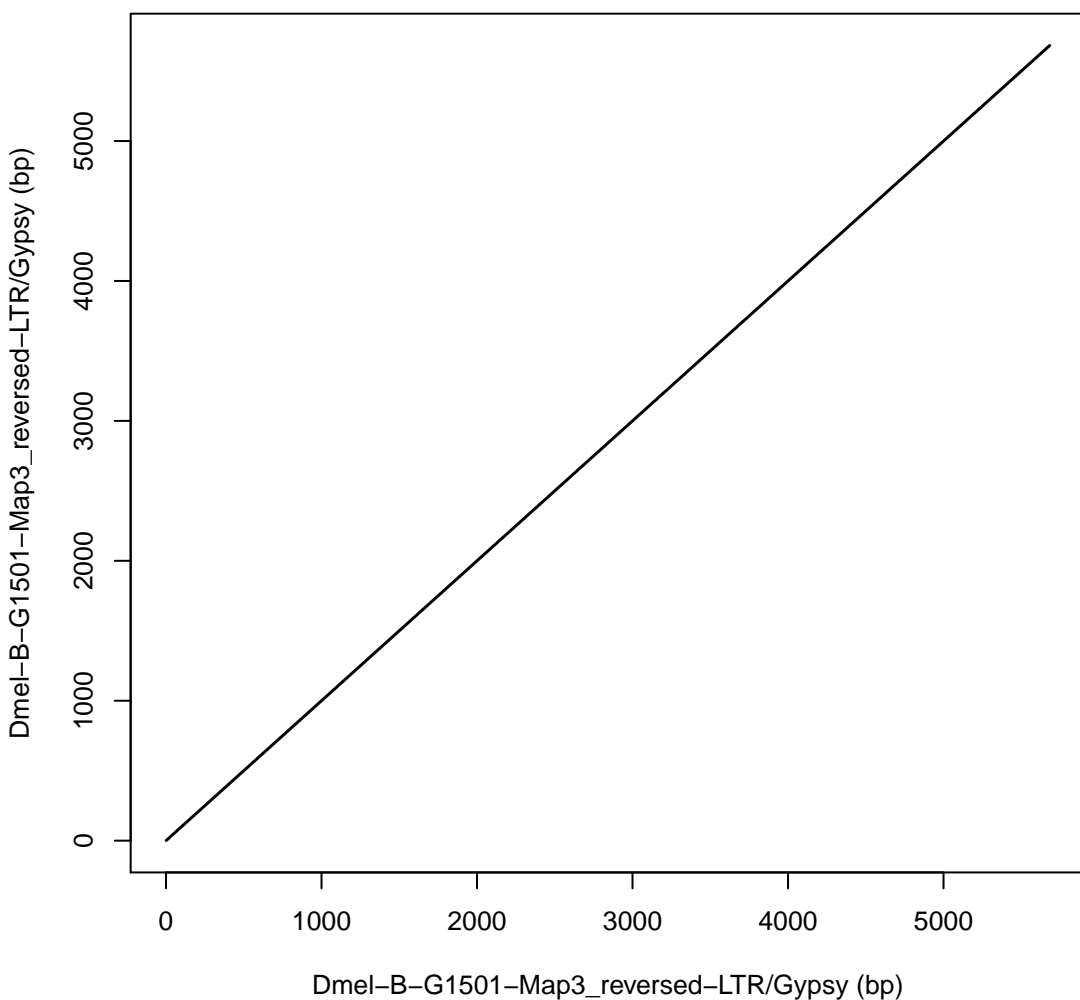
TE: Dmel-B-G1501-Map3\_reversed-LTR/Gypsy  
consensus size: 5683bp; fragments: 78; full length: 3 (>=5114.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

