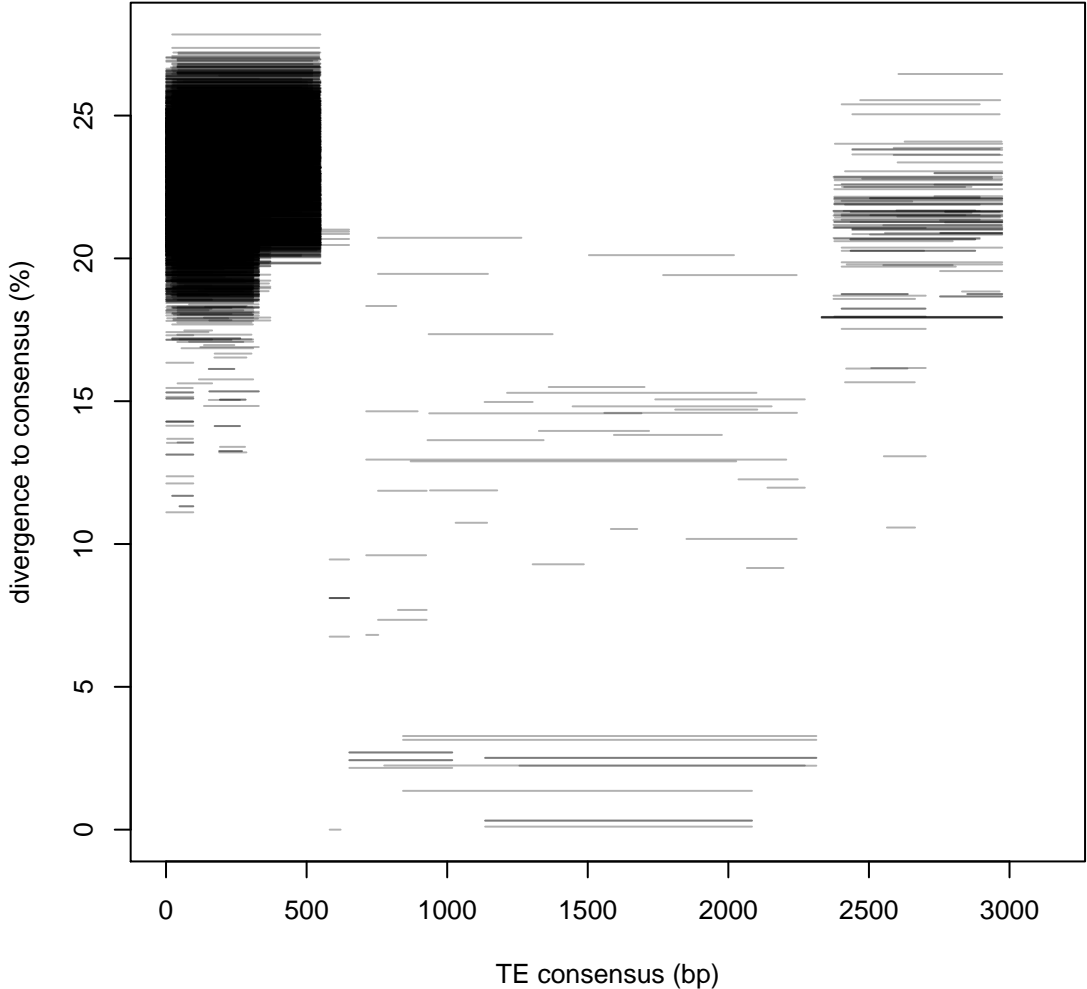
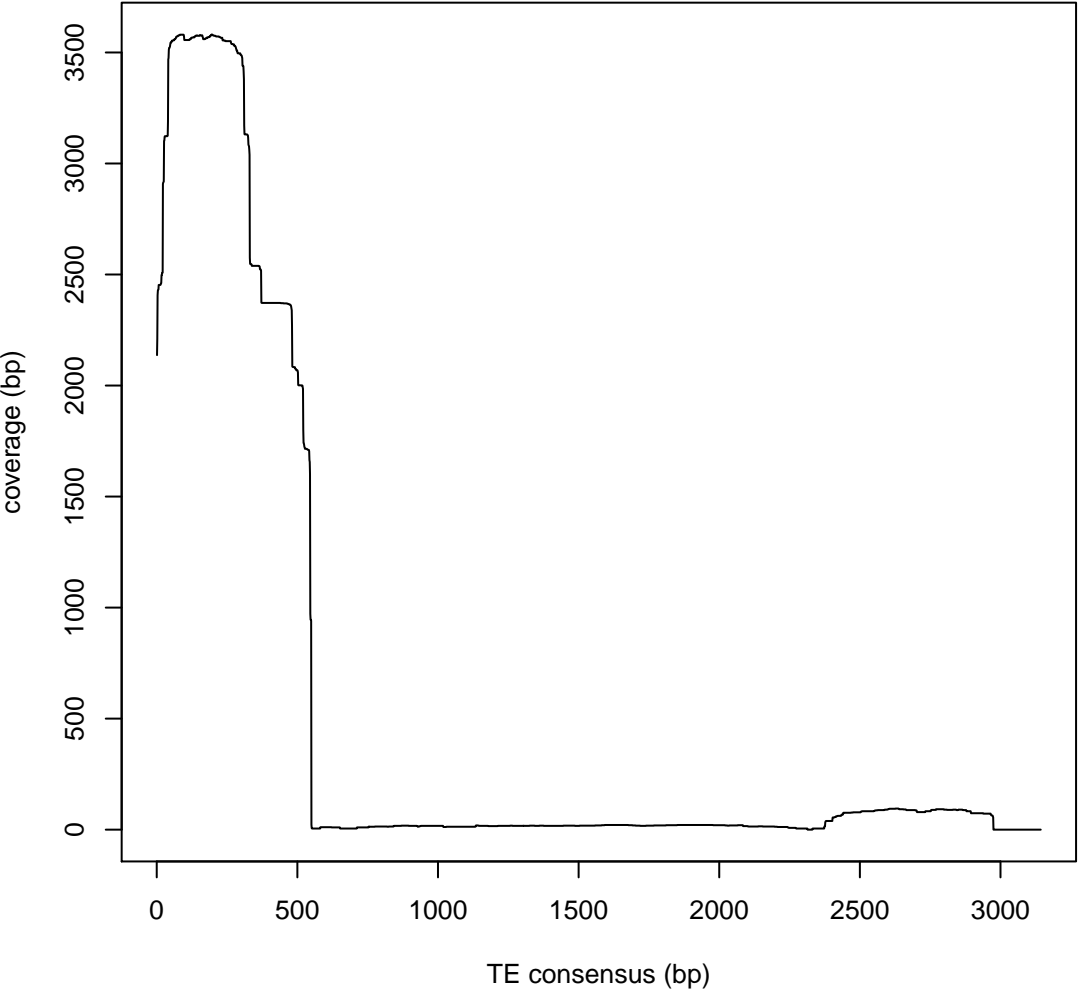


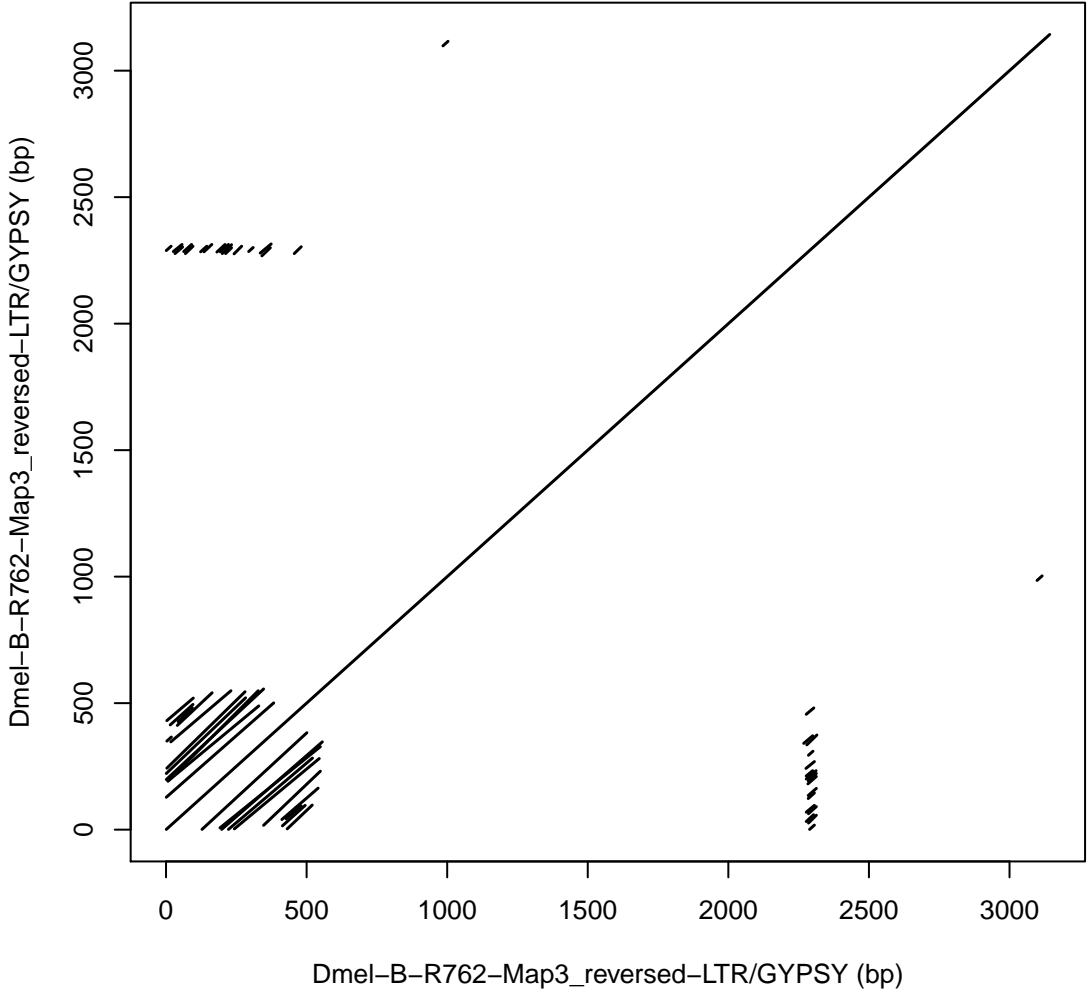
TE: Dmel-B-R762-Map3\_reversed-LTR/Gypsy  
consensus size: 3143bp; fragments: 3796; full length: 0 (>=2828.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

