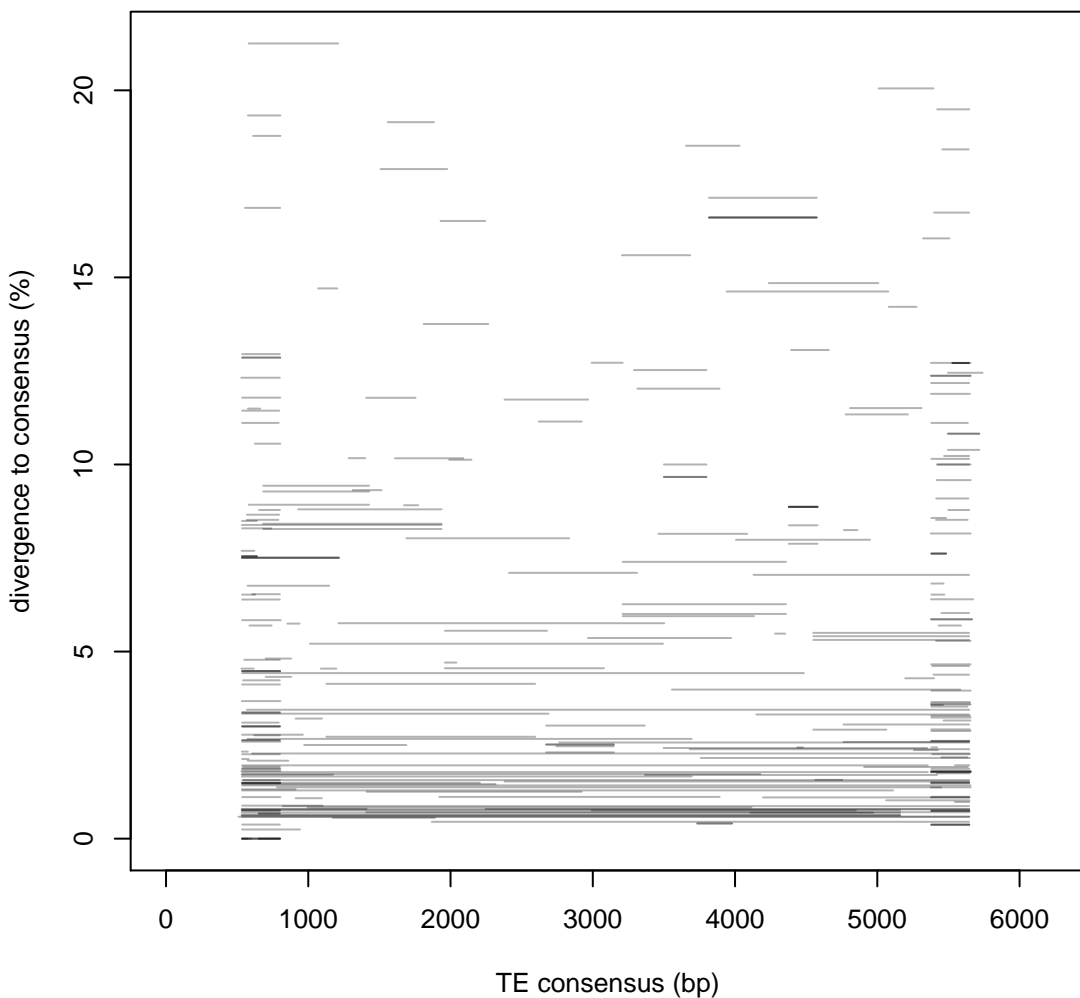
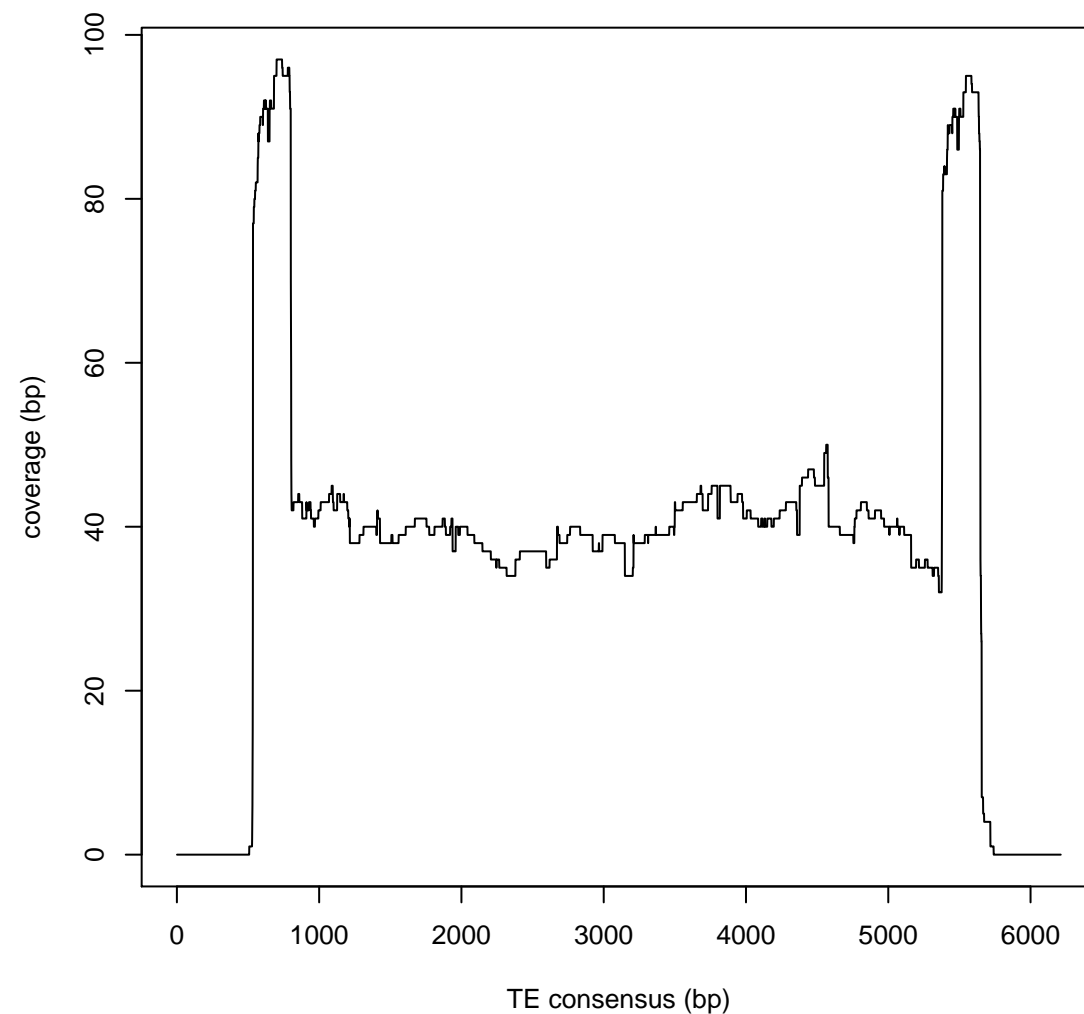


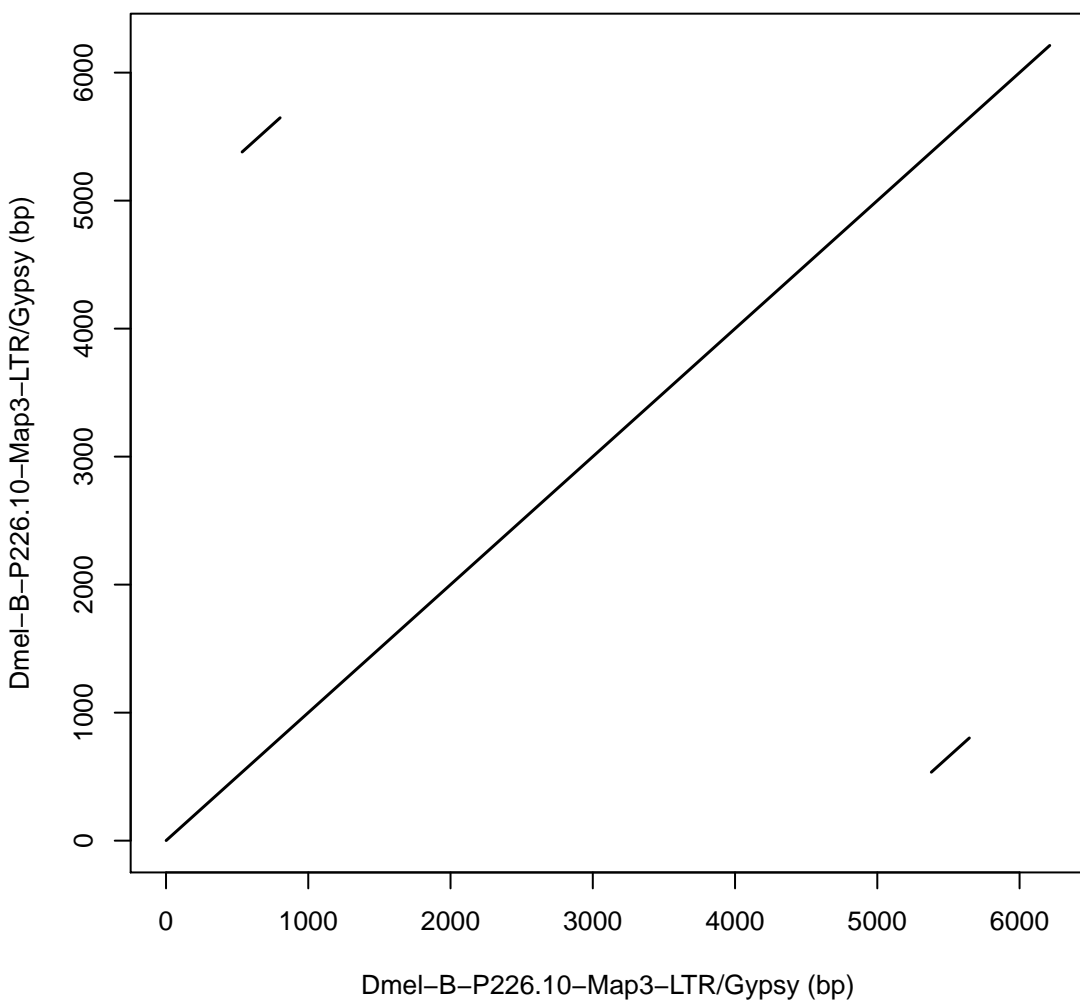
TE: Dmel-B-P226.10-Map3-LTR/Gypsy
consensus size: 6212bp; fragments: 289; full length: 0 (>=5590.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

