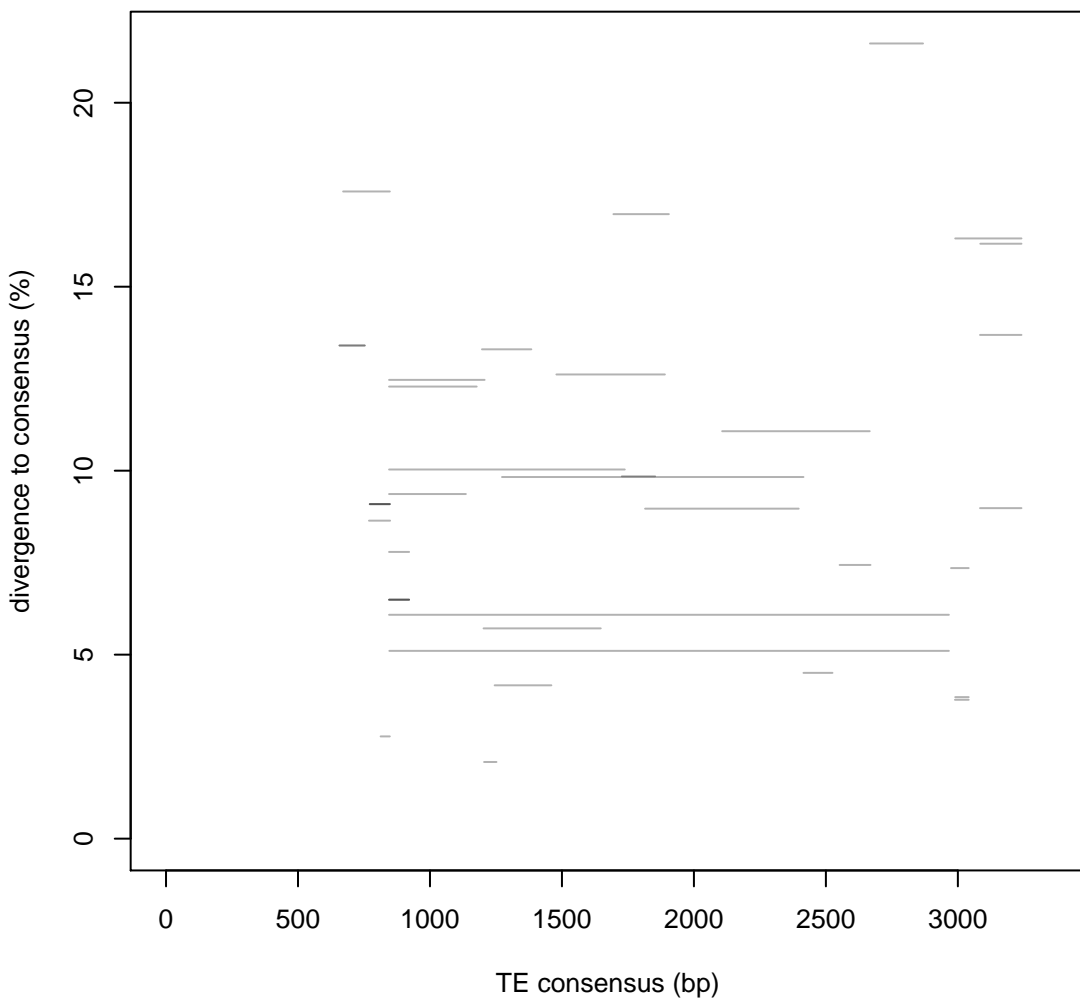
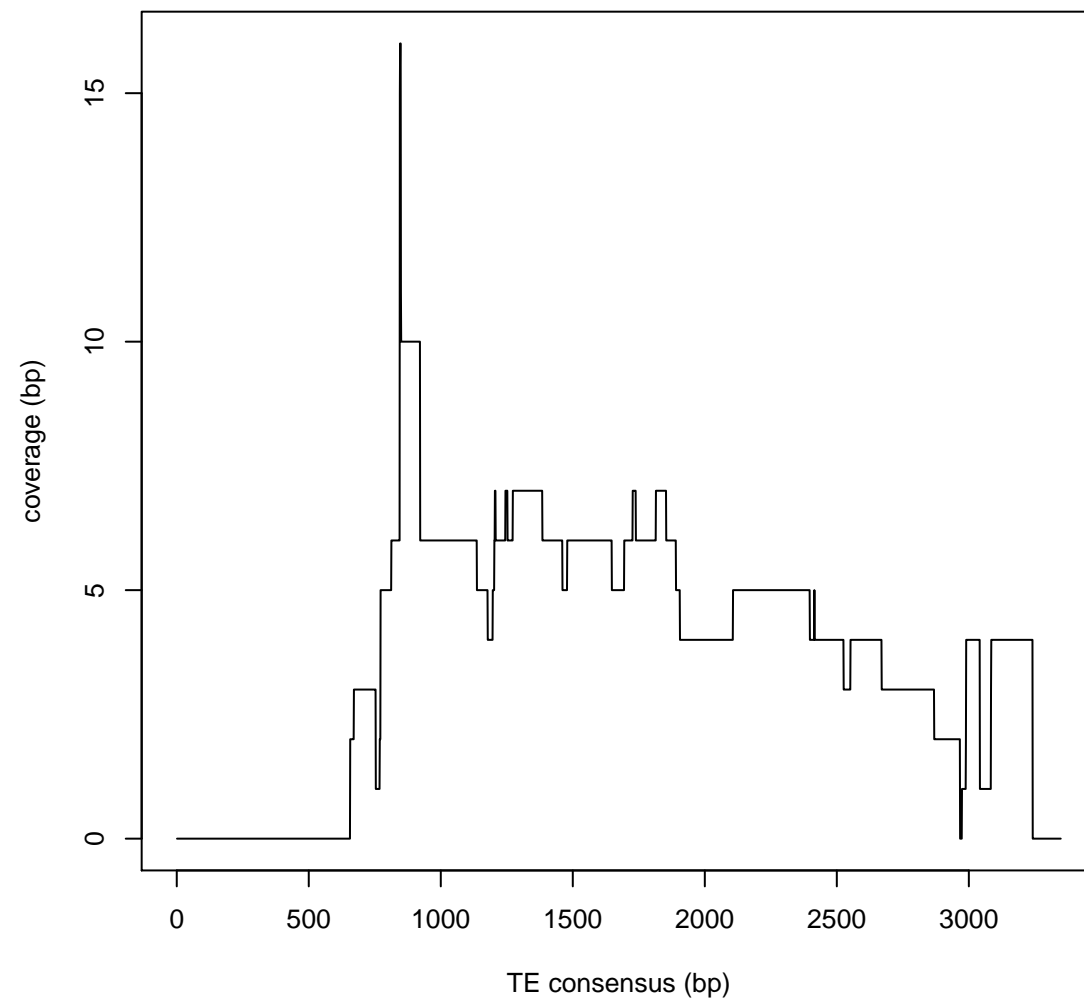


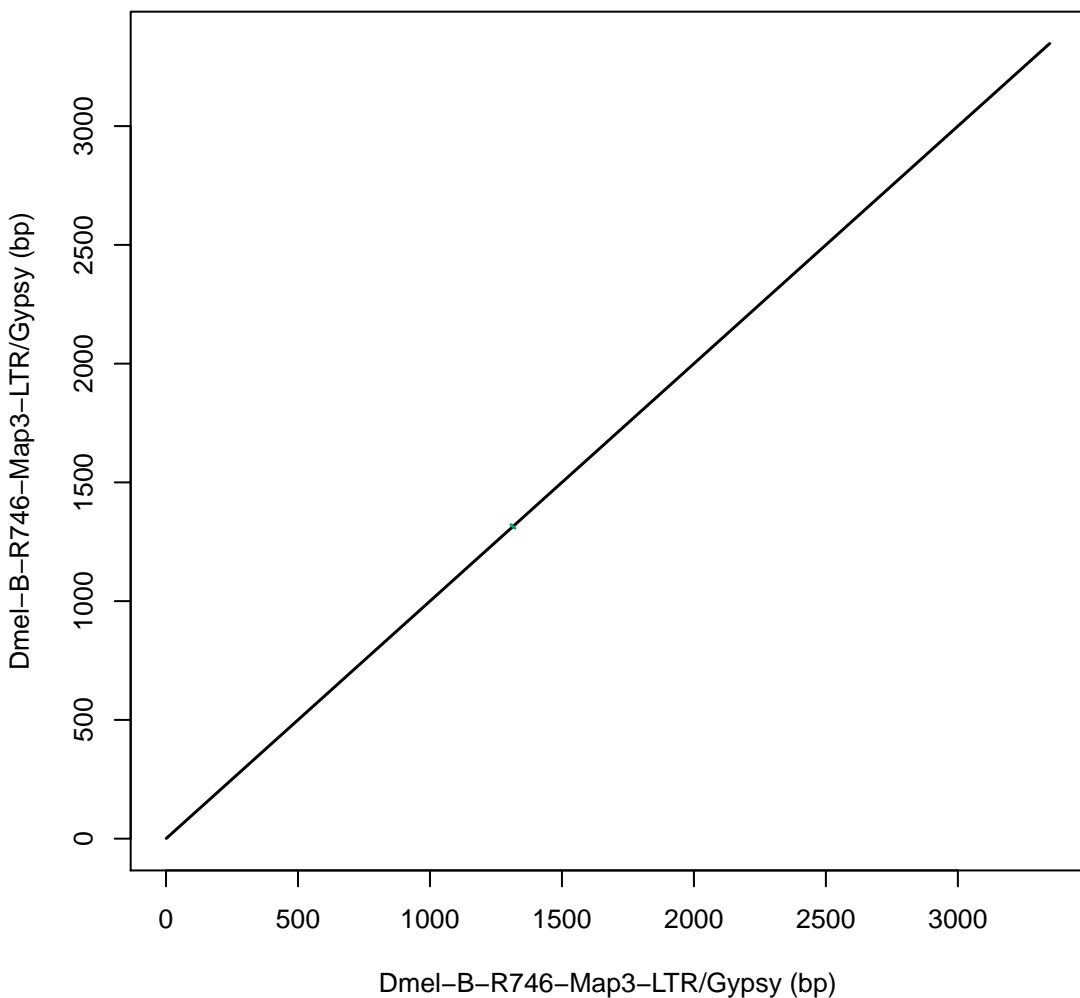
TE: Dmel-B-R746-Map3-LTR/Gypsy
consensus size: 3348bp; fragments: 38; full length: 0 (>=3013.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

