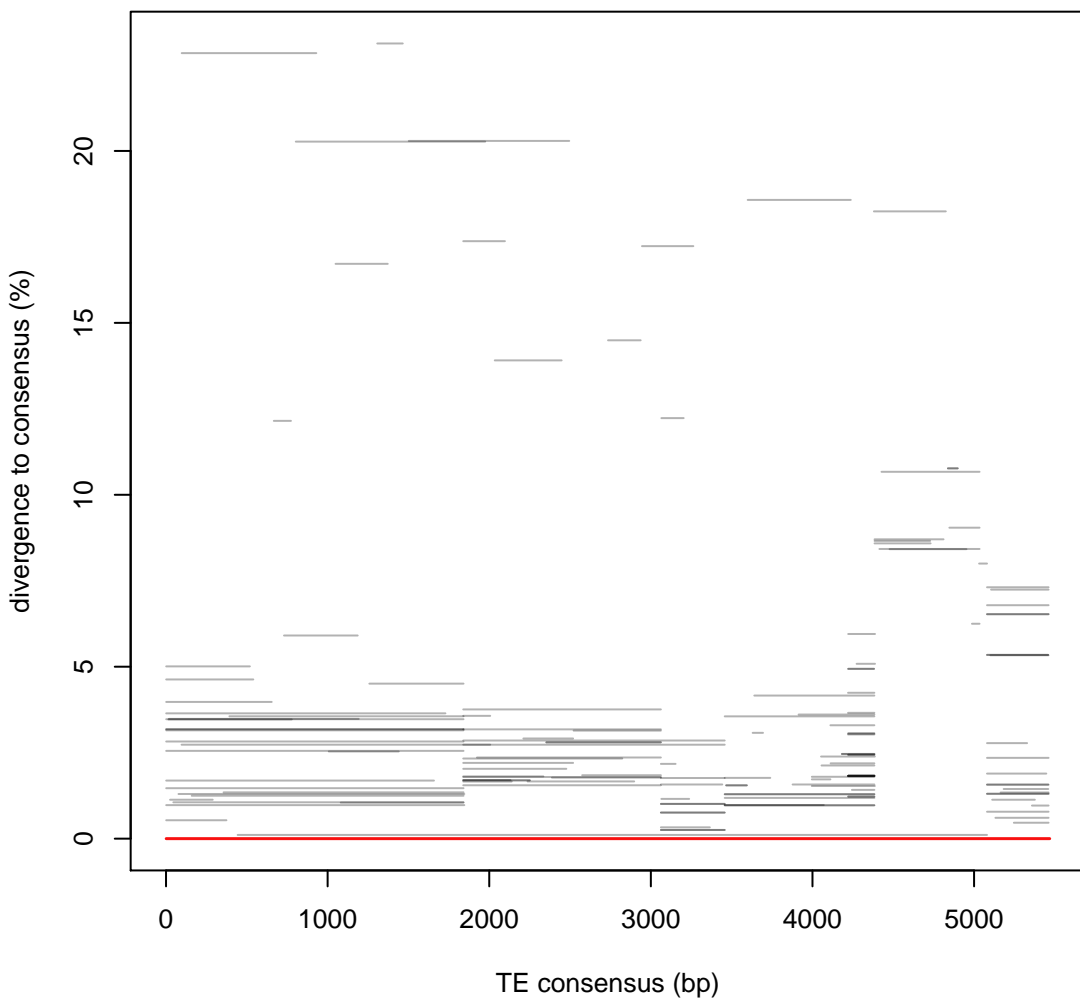
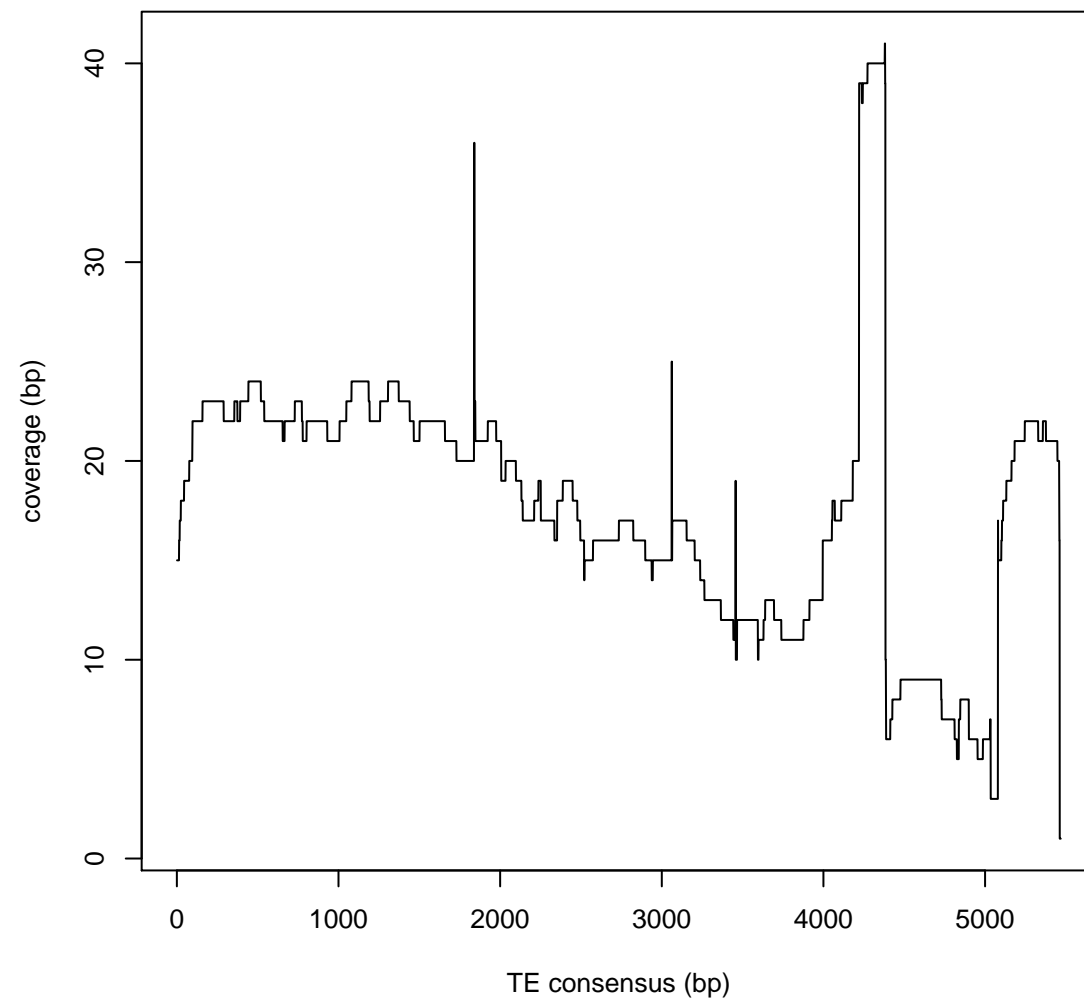


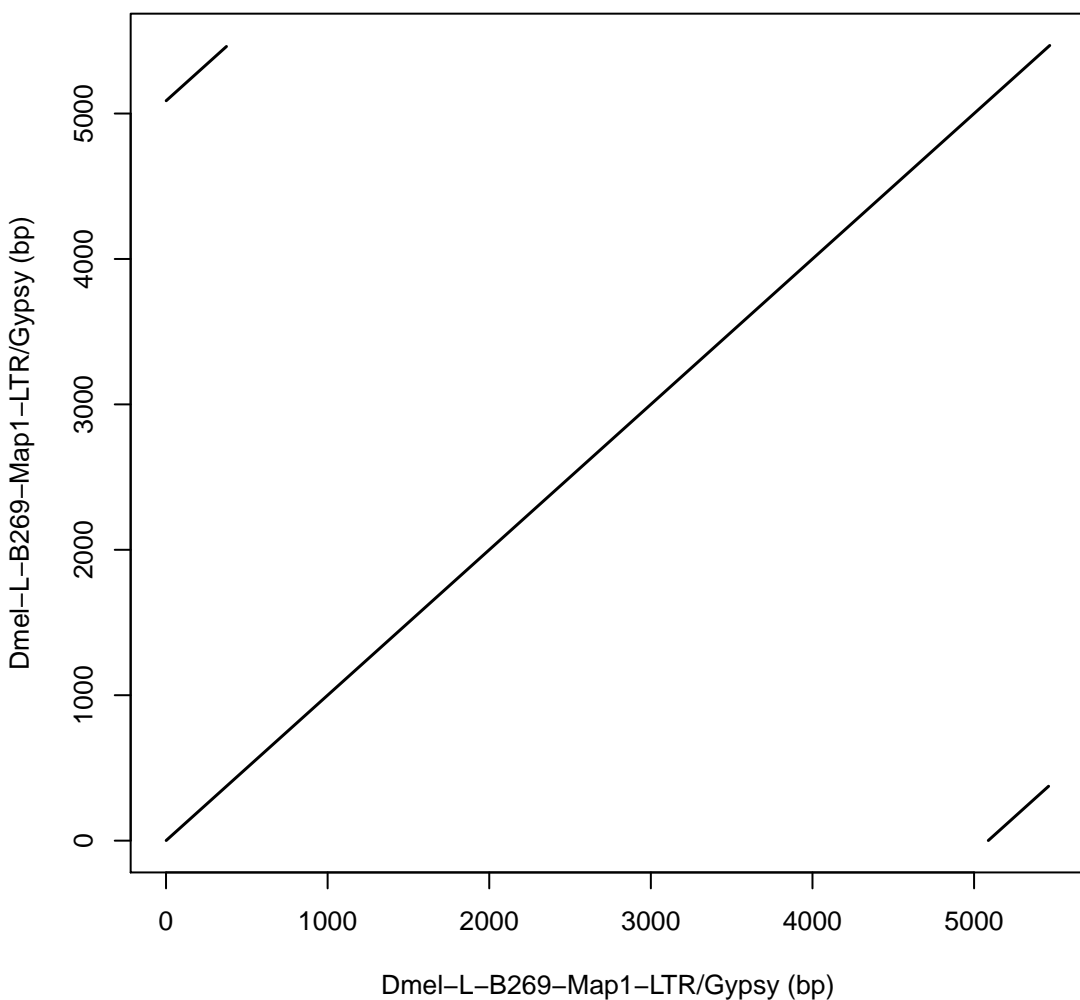
TE: Dmel-L-B269-Map1-LTR/Gypsy
consensus size: 5468bp; fragments: 153; full length: 1 (>=4921.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

