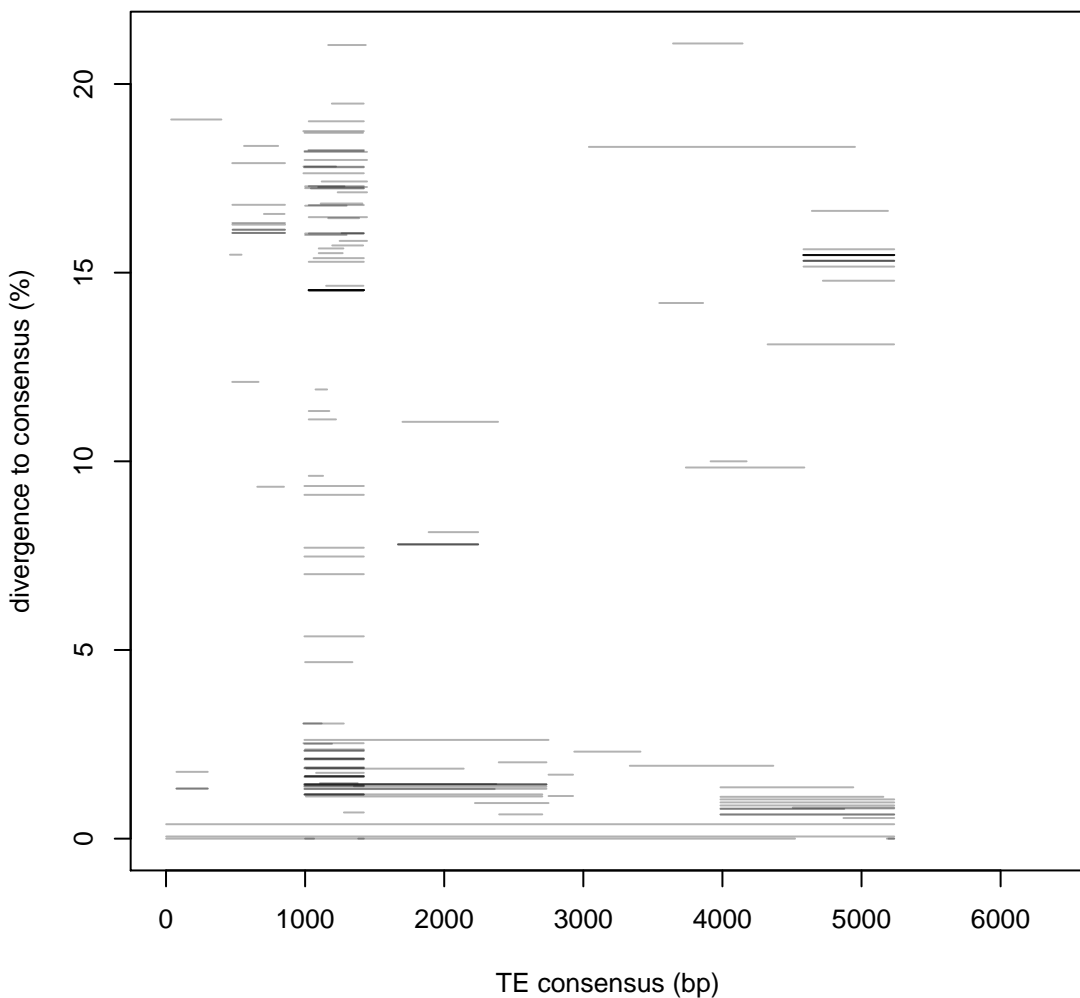
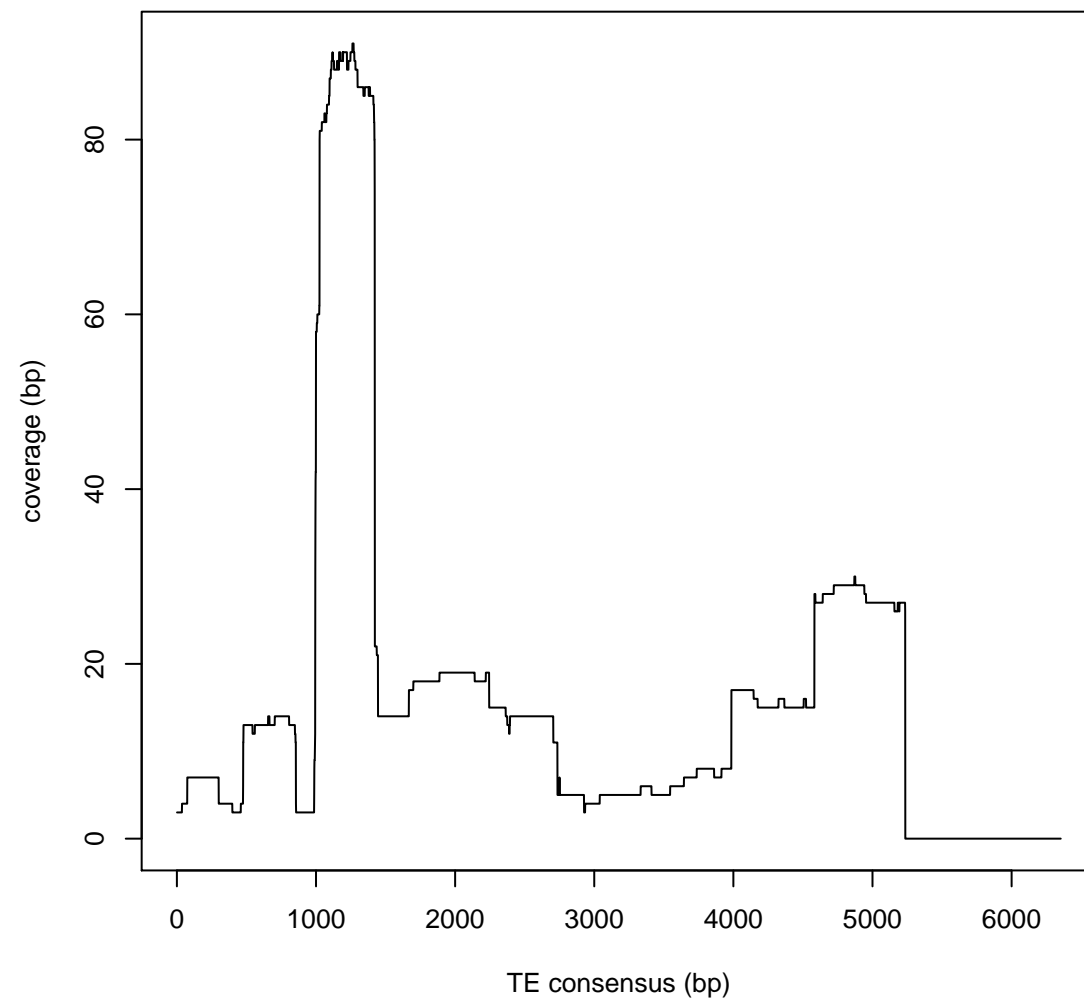


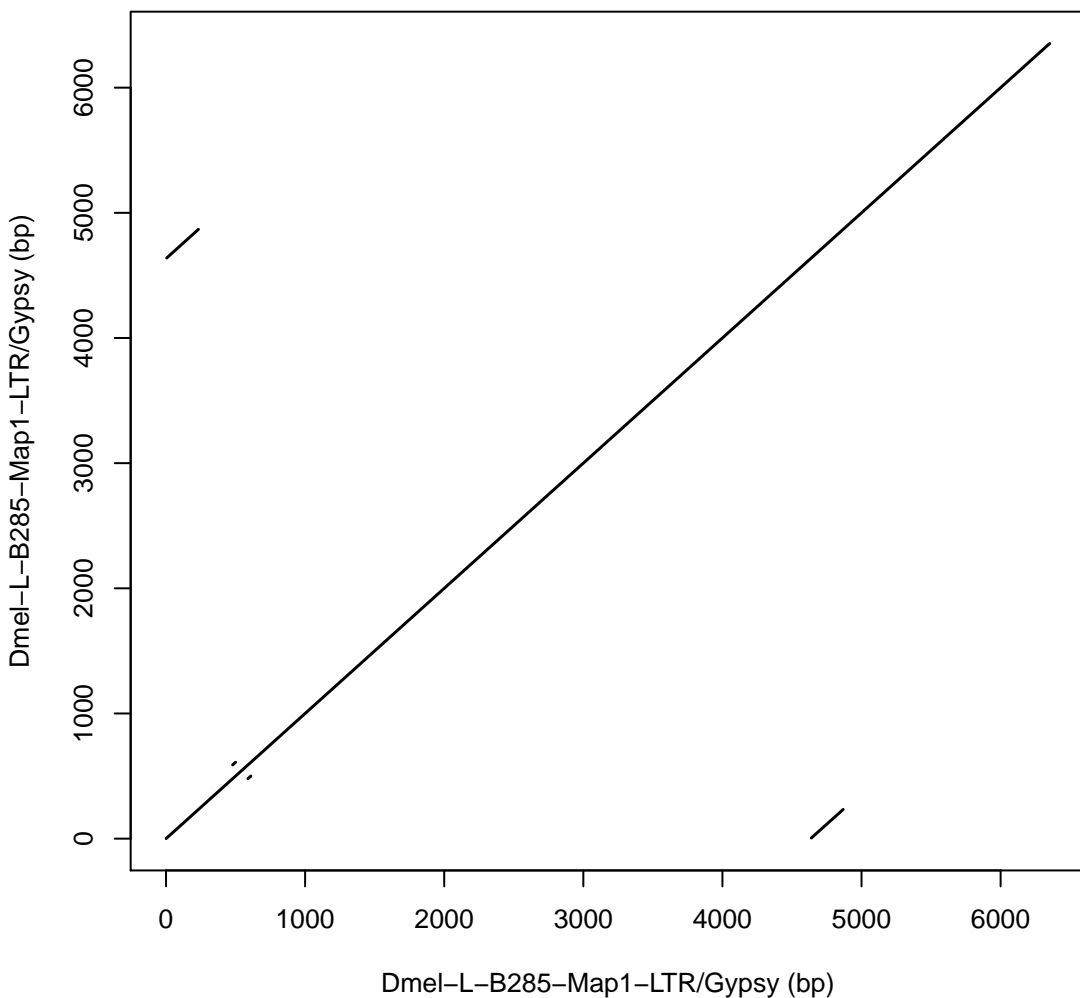
TE: Dmel-L-B285-Map1-LTR/Gypsy
consensus size: 6353bp; fragments: 165; full length: 0 (>=5717.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

