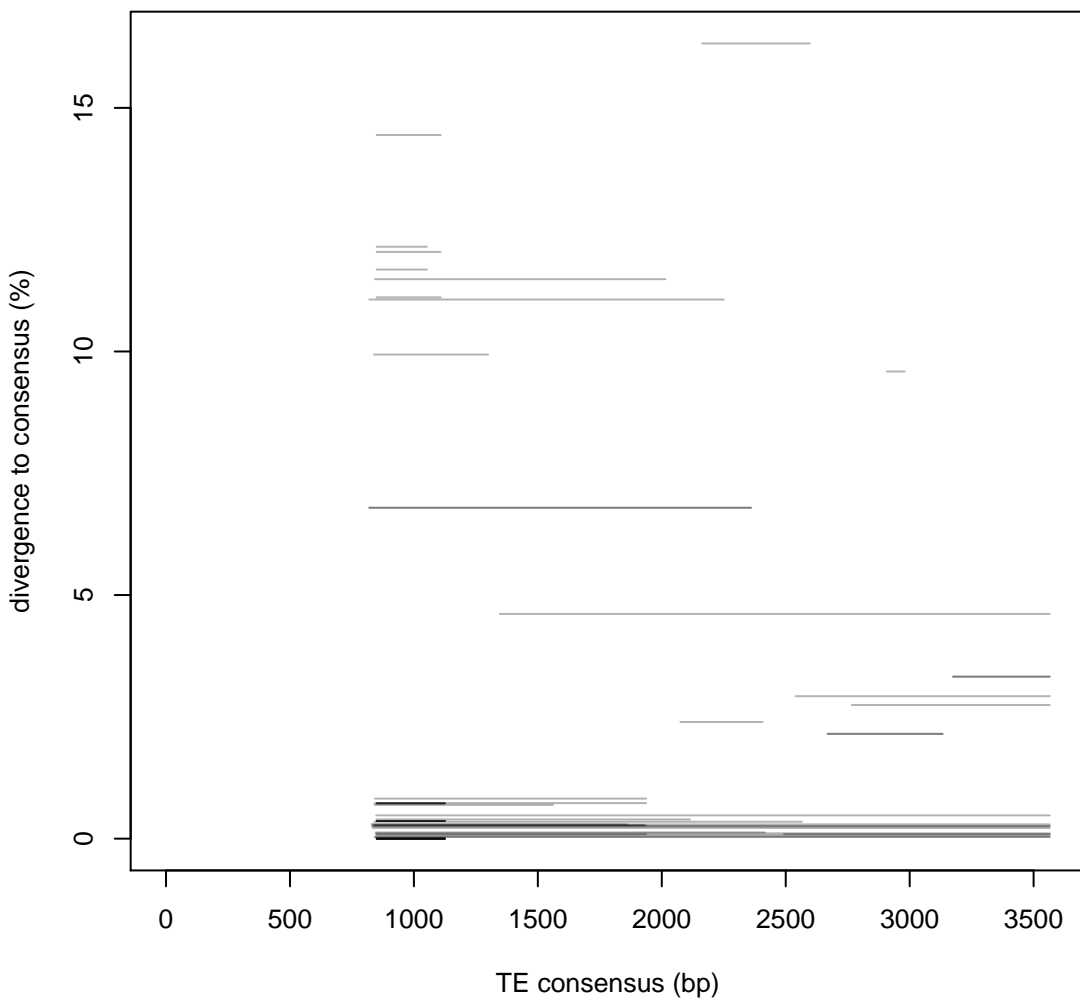
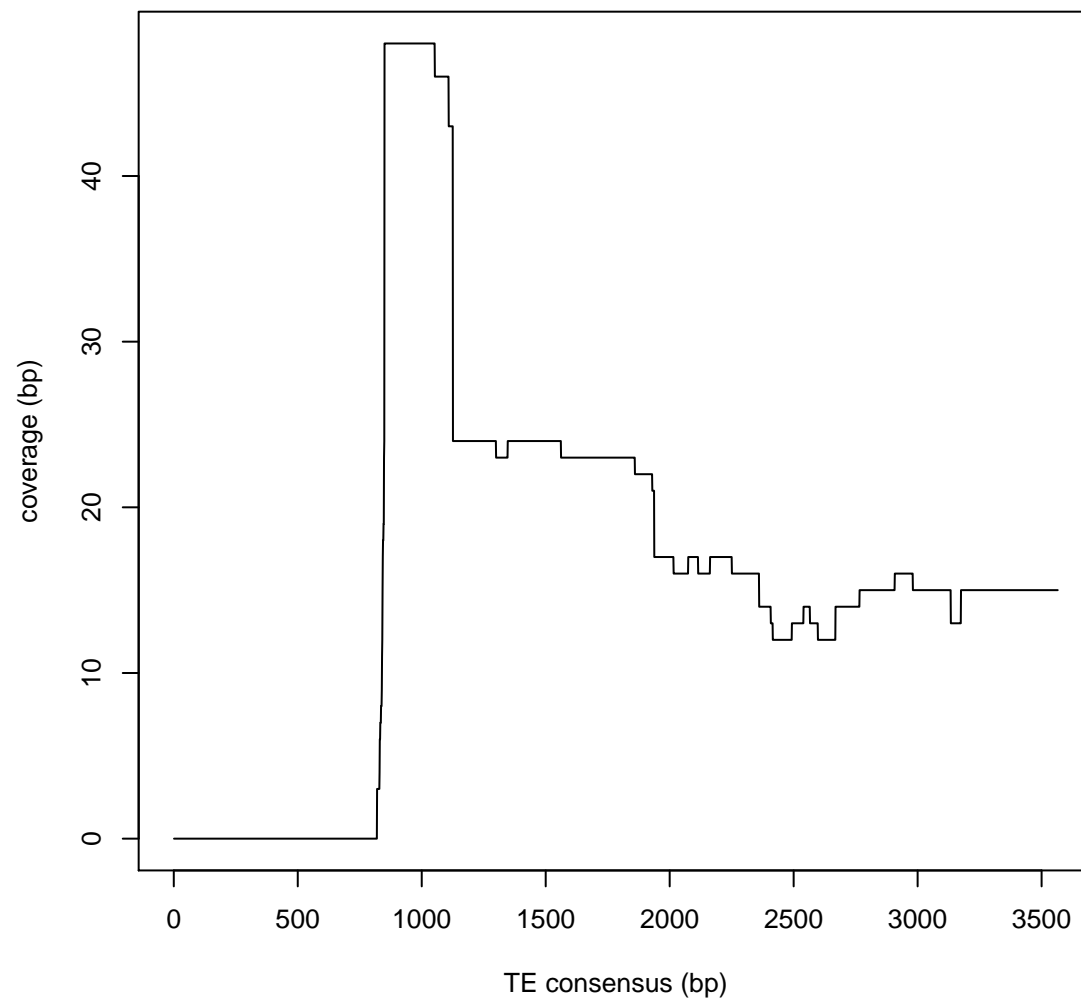


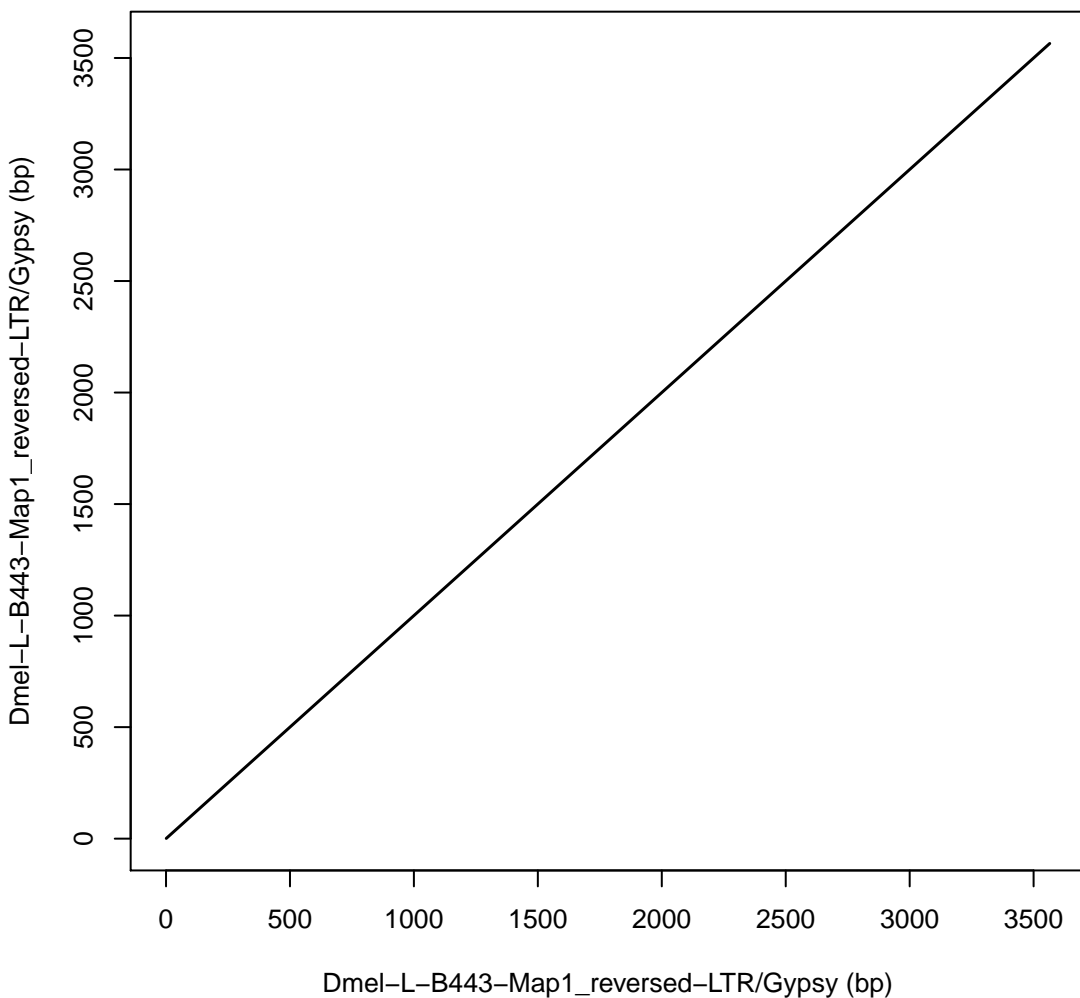
TE: Dmel-L-B443-Map1_reversed-LTR/Gypsy
consensus size: 3565bp; fragments: 59; full length: 0 (>=3208.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

