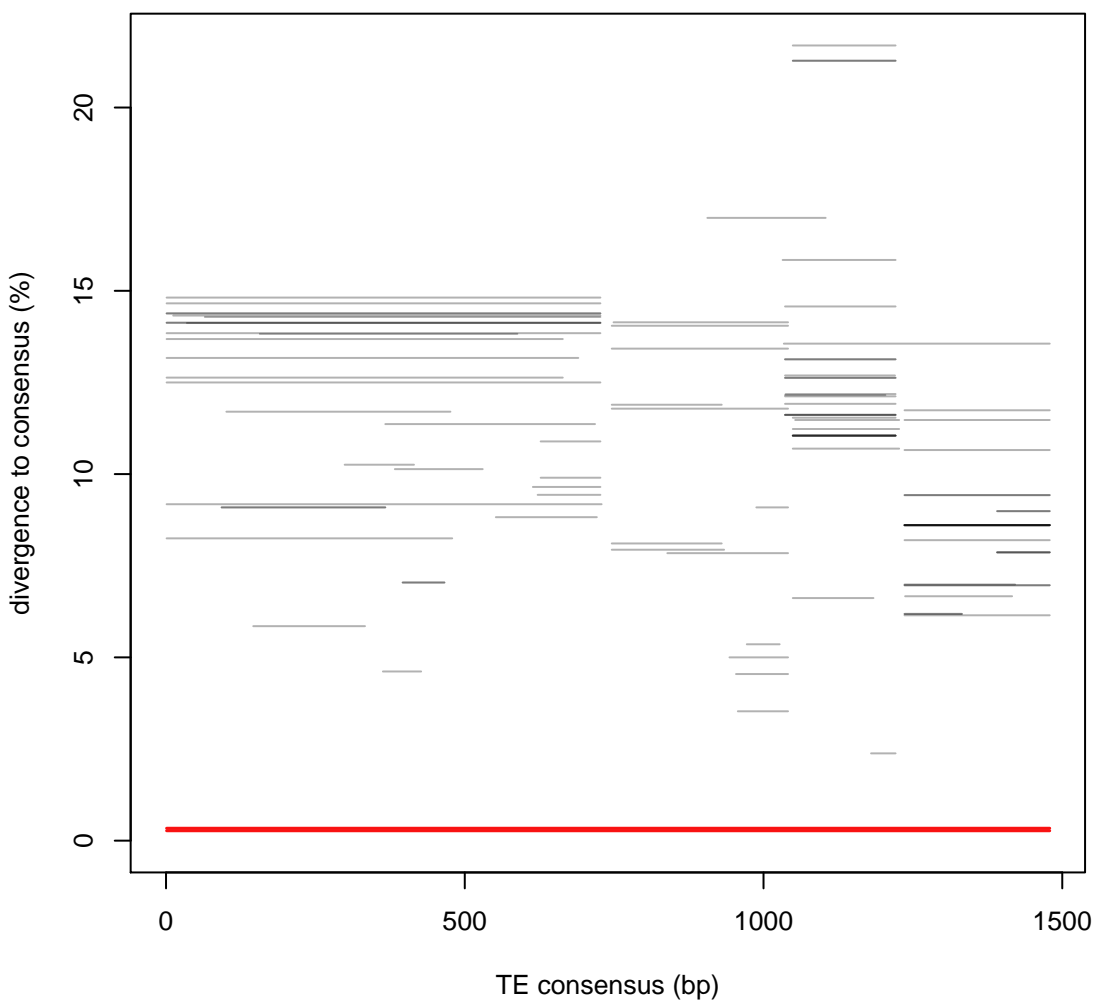
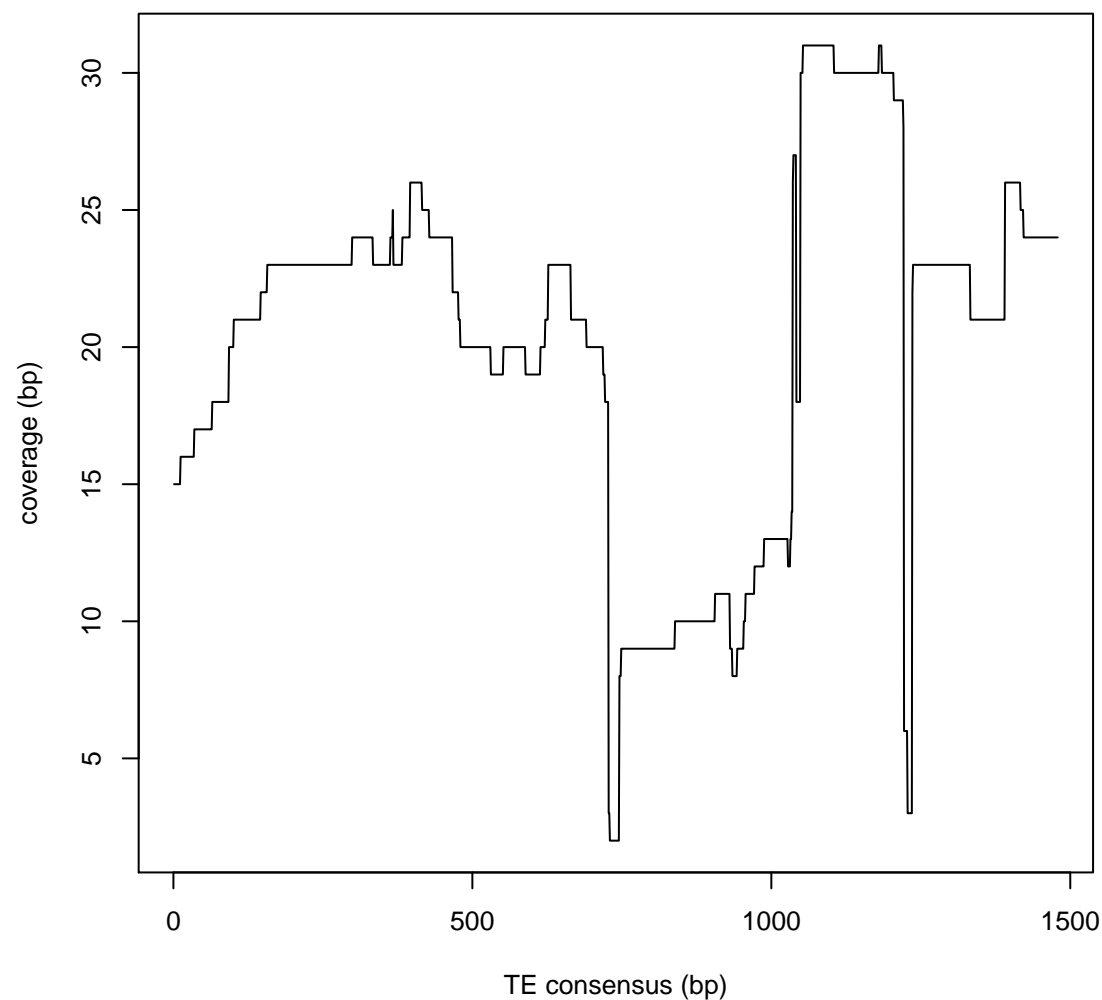


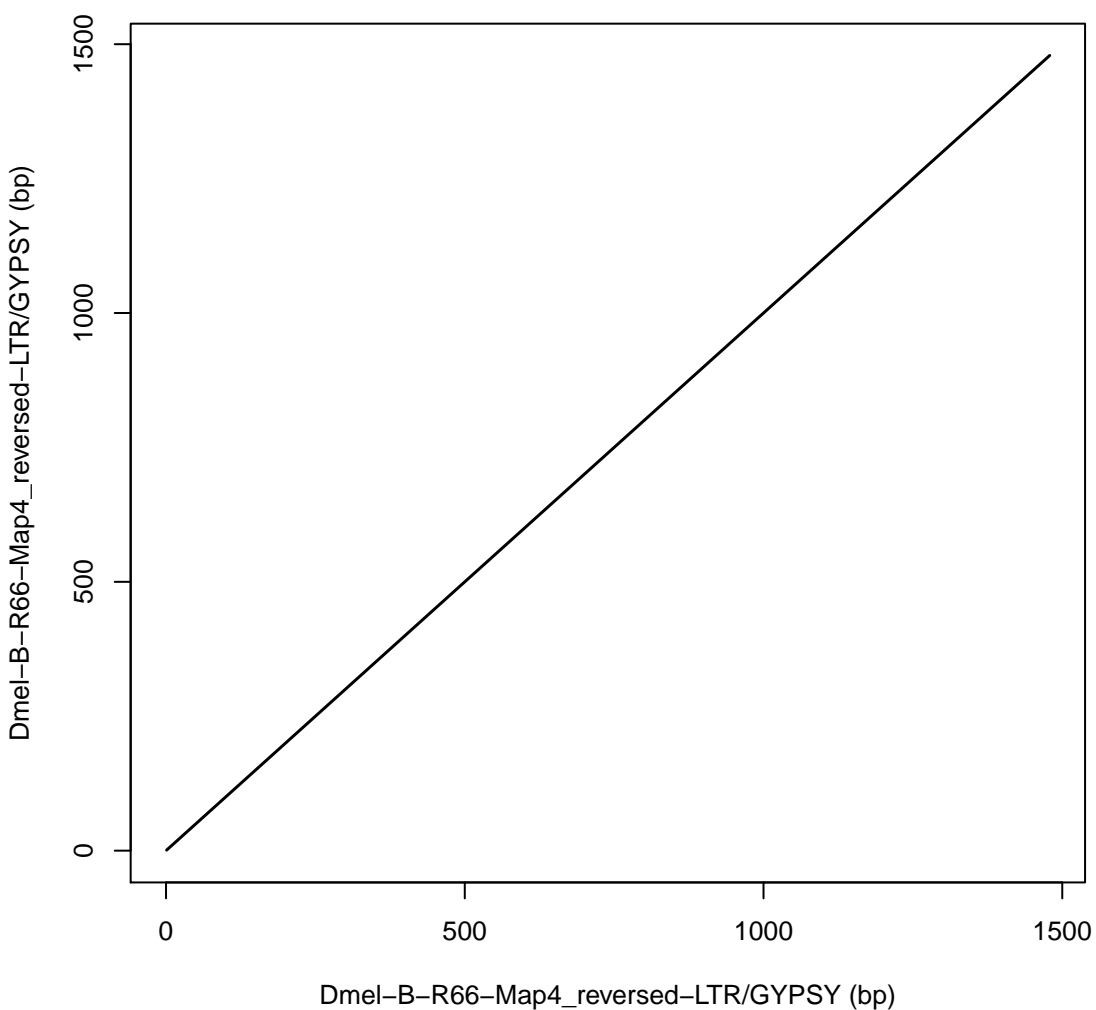
TE: Dmel-B-R66-Map4_reversed-LTR/GYPSY
consensus size: 1479bp; fragments: 102; full length: 2 (>=1331.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

