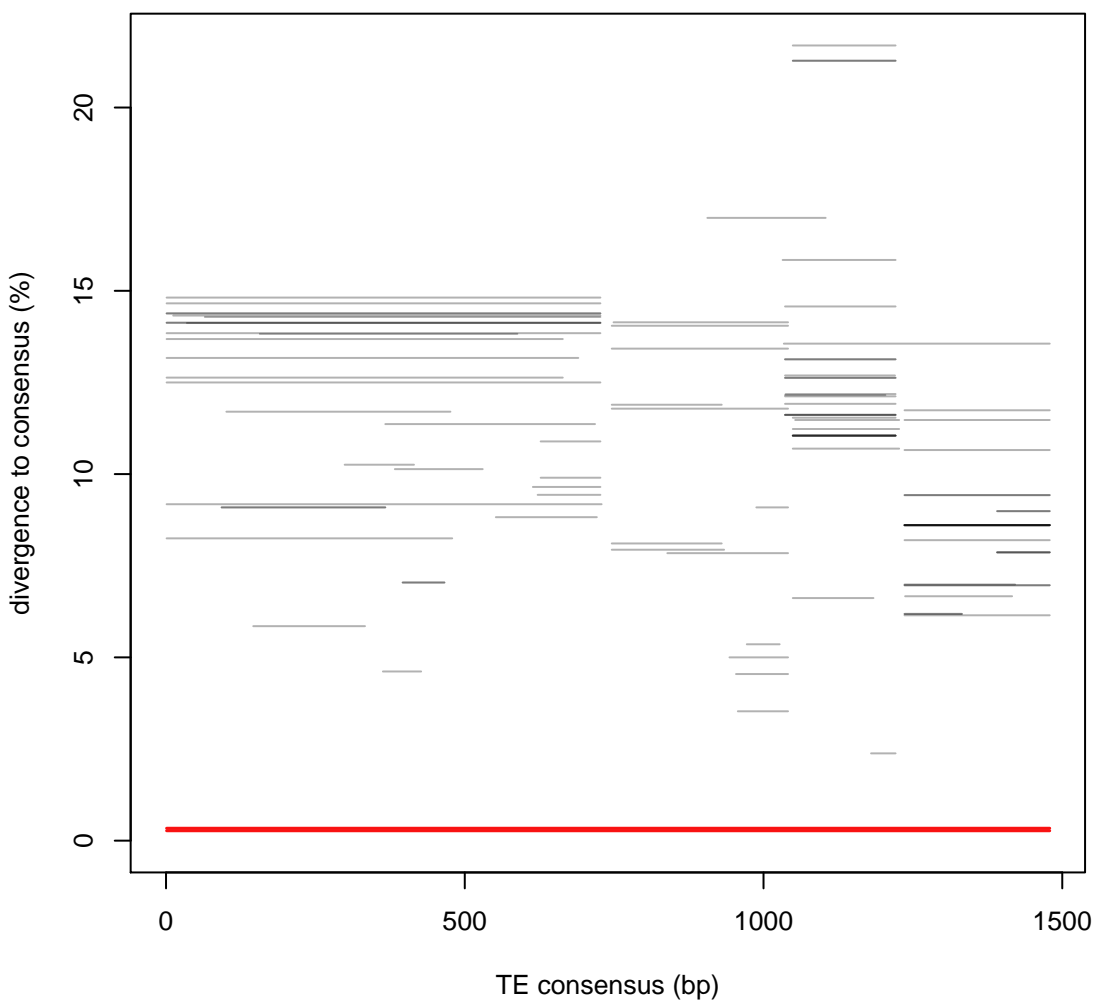
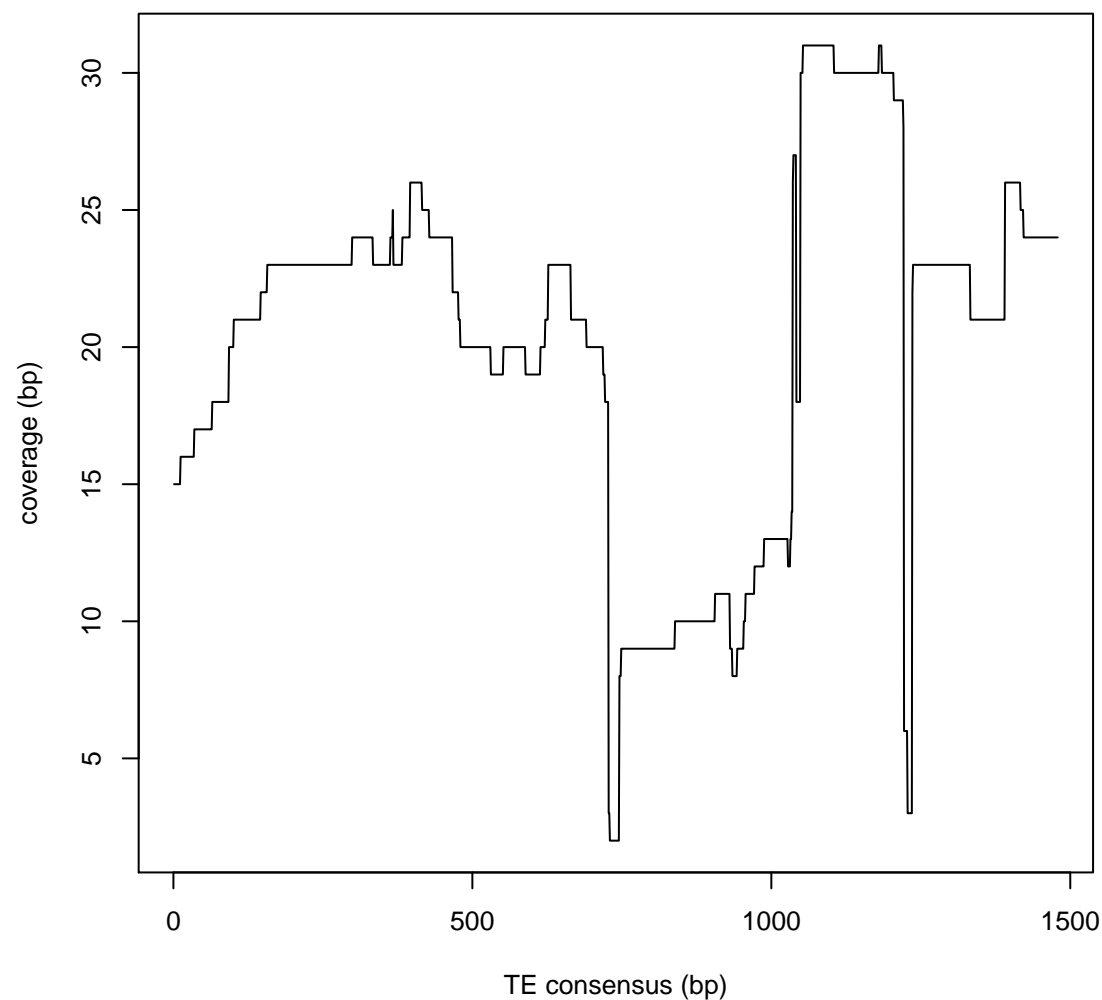


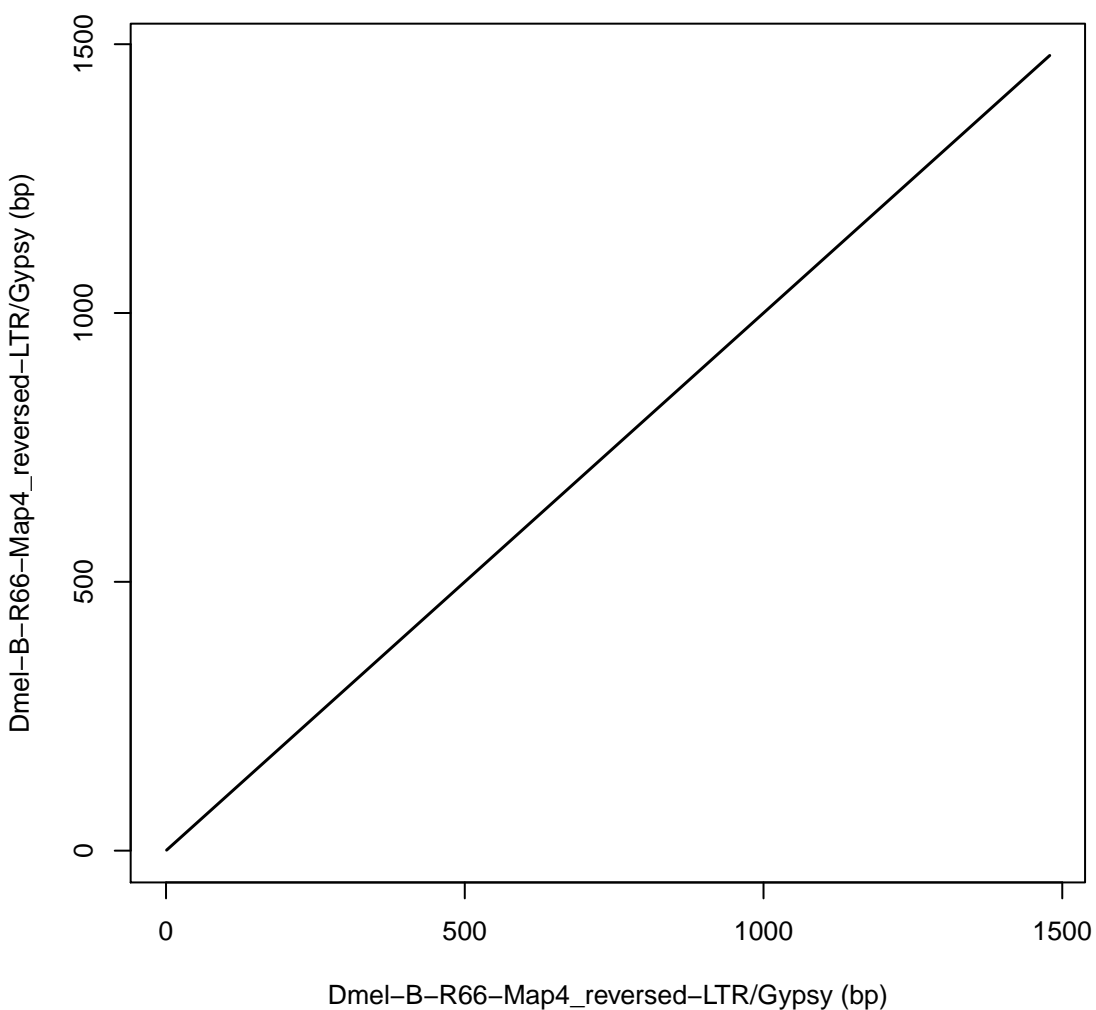
TE: Dmel-B-R66-Map4_reversed-LTR/Gypsy
consensus size: 1479bp; fragments: 102; full length: 2 (≥ 1331.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

