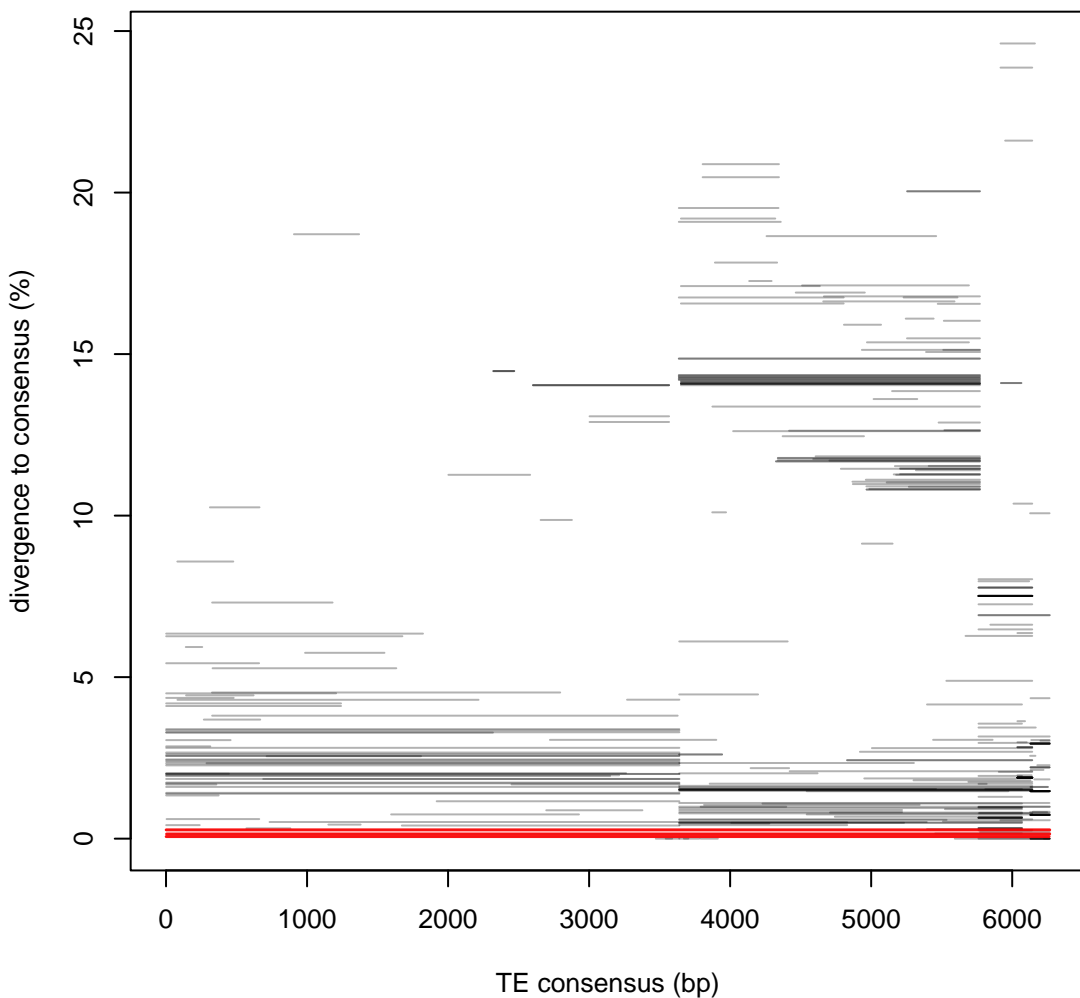
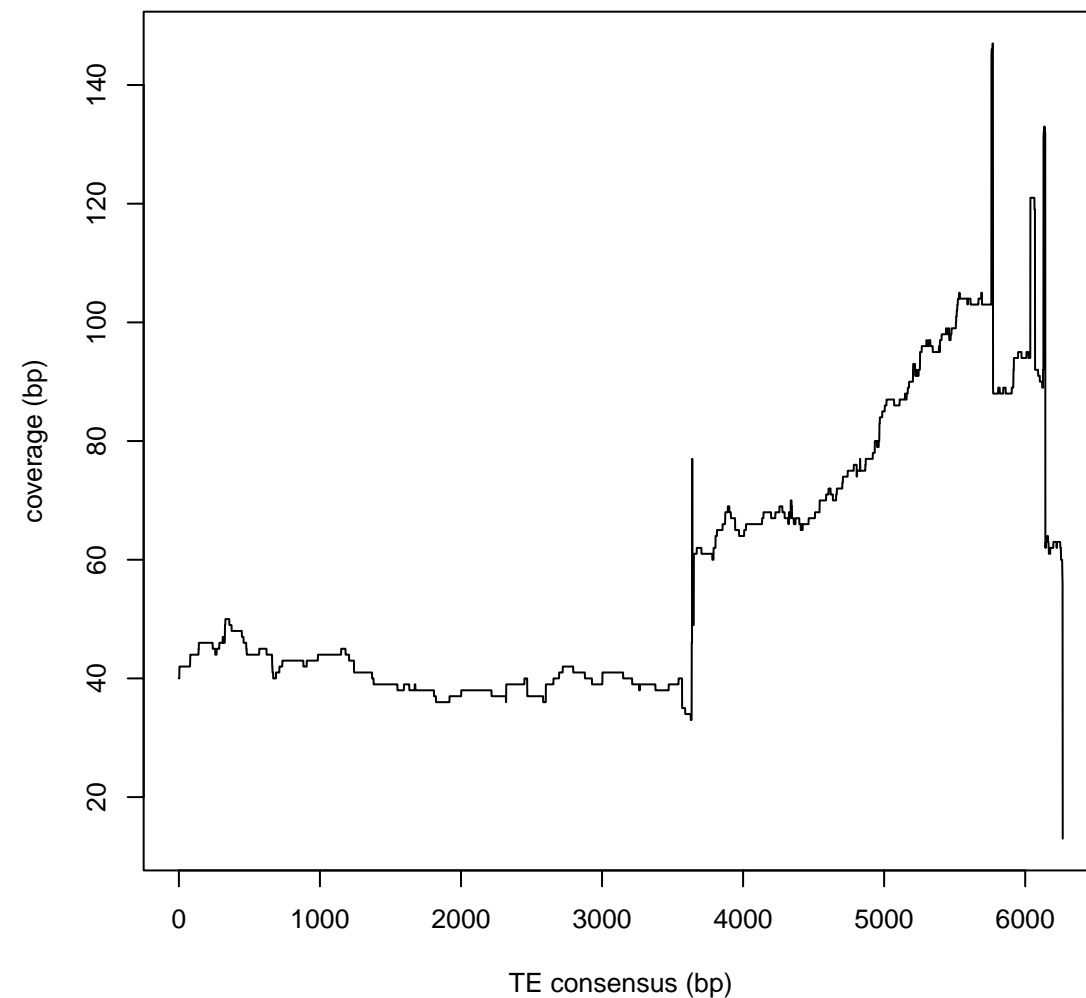


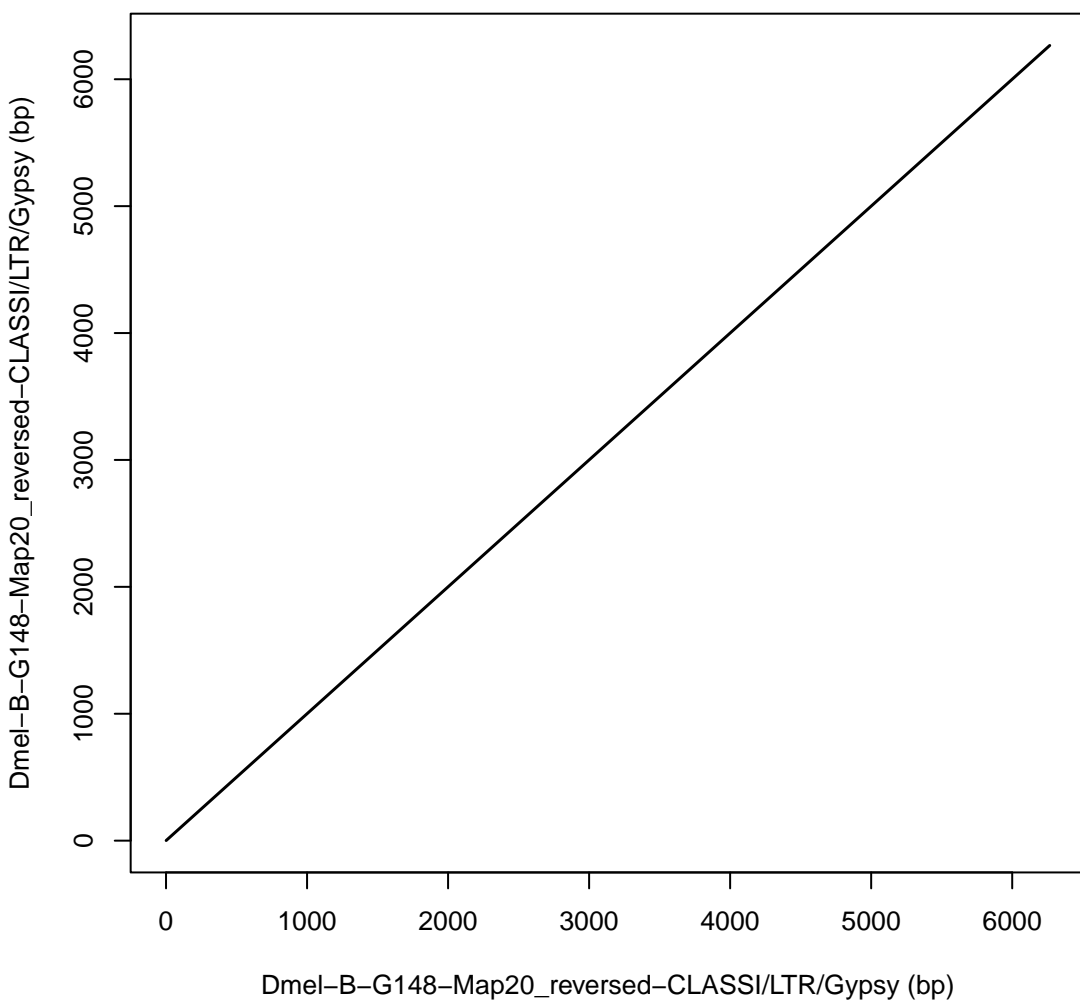
TE: Dmel-B-G148-Map20_reversed-CLASSI/LTR/Gypsy
consensus size: 6266bp; fragments: 353; full length: 3 (>=5639.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

