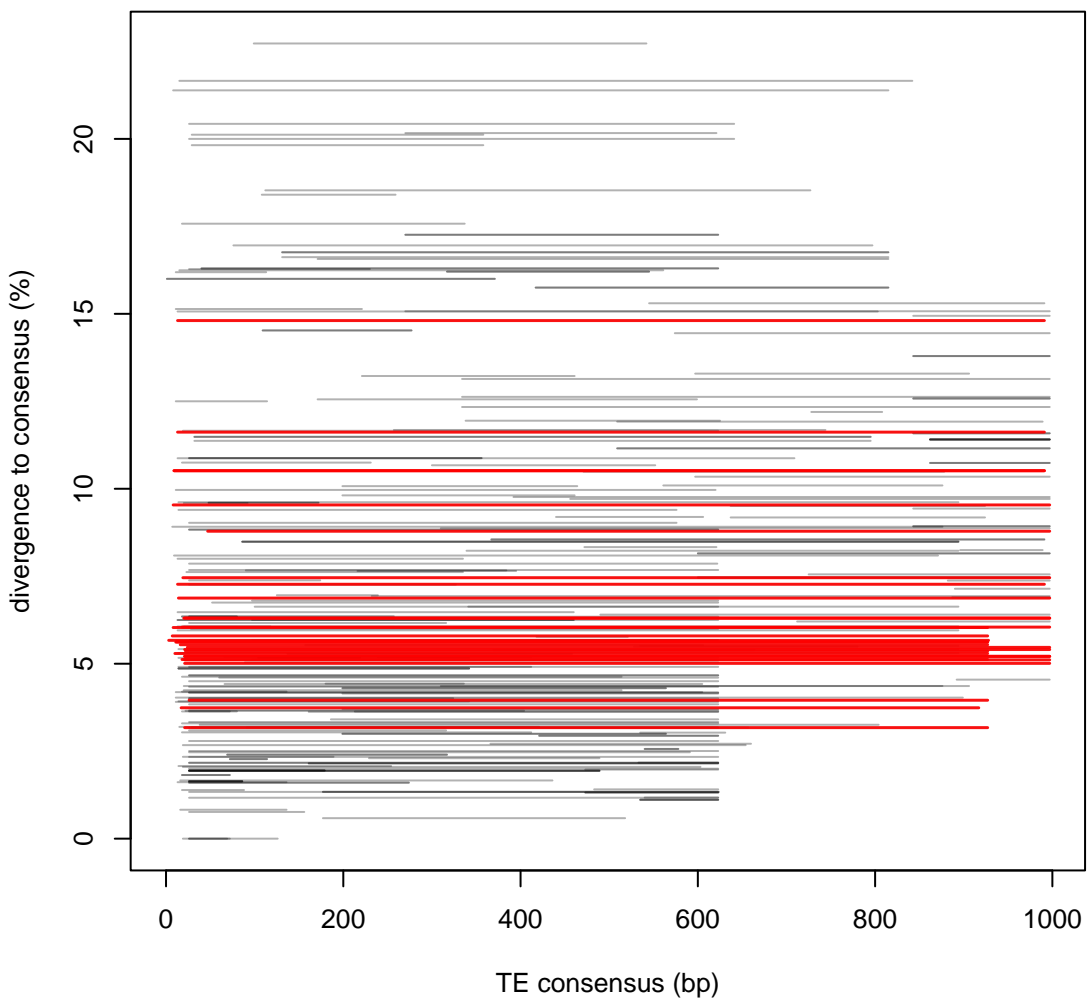
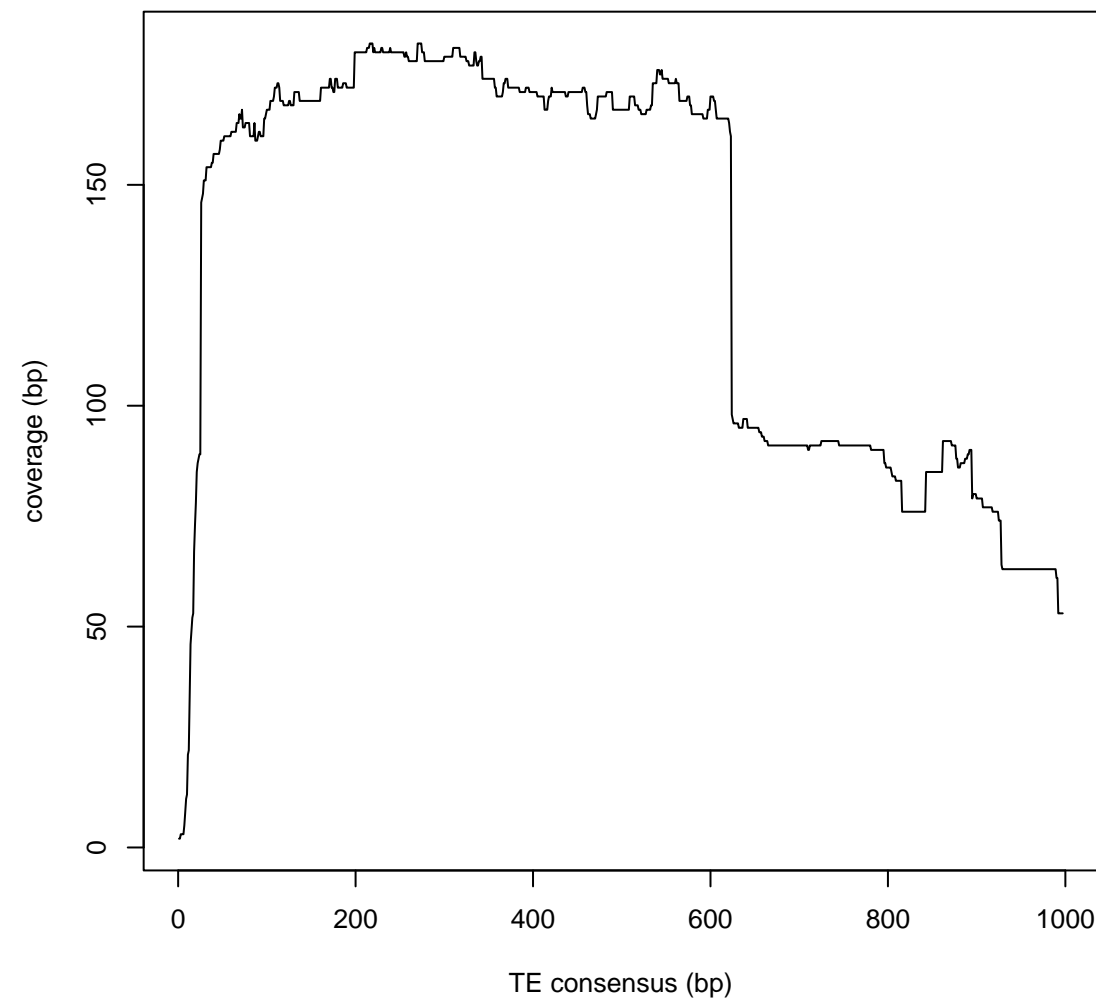


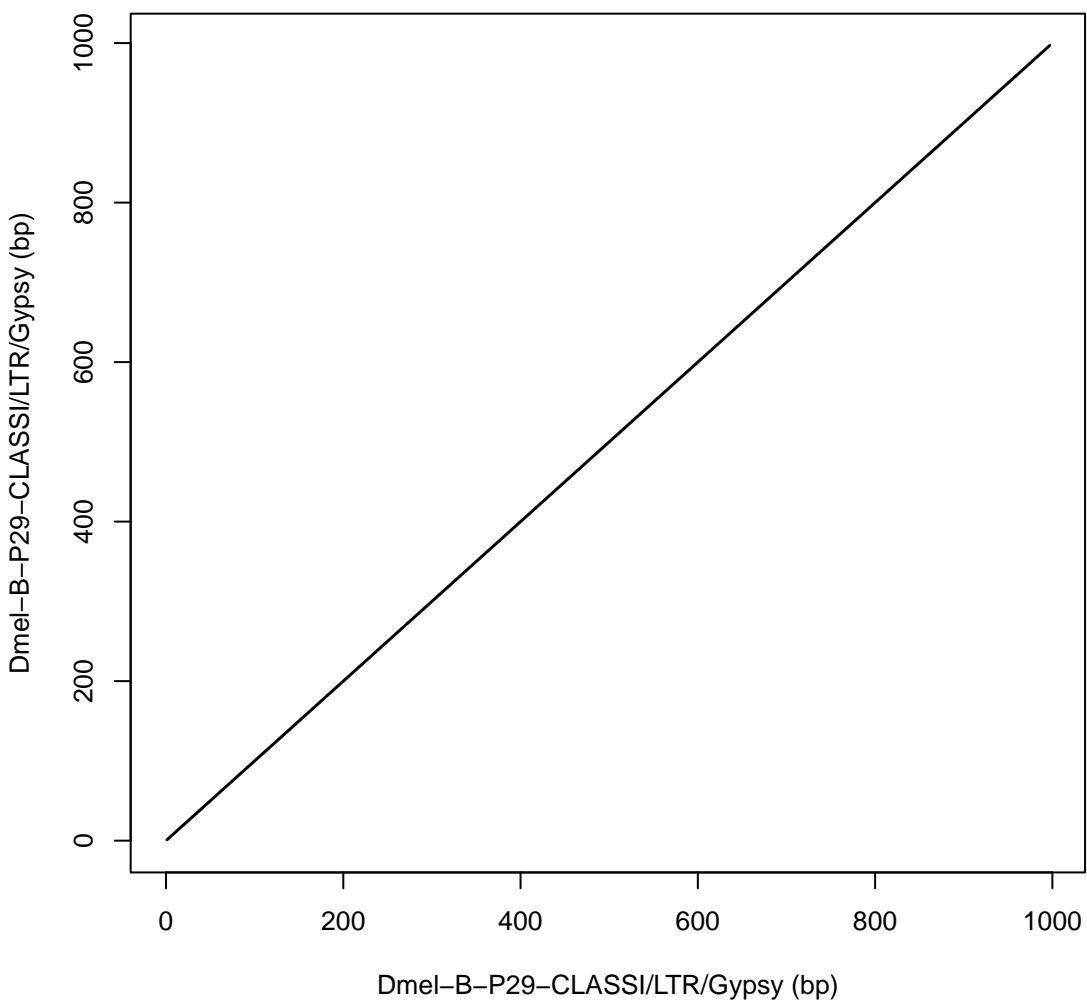
TE: Dmel-B-P29-CLASSI/LTR/Gypsy
consensus size: 997bp; fragments: 301; full length: 30 (>=897.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

