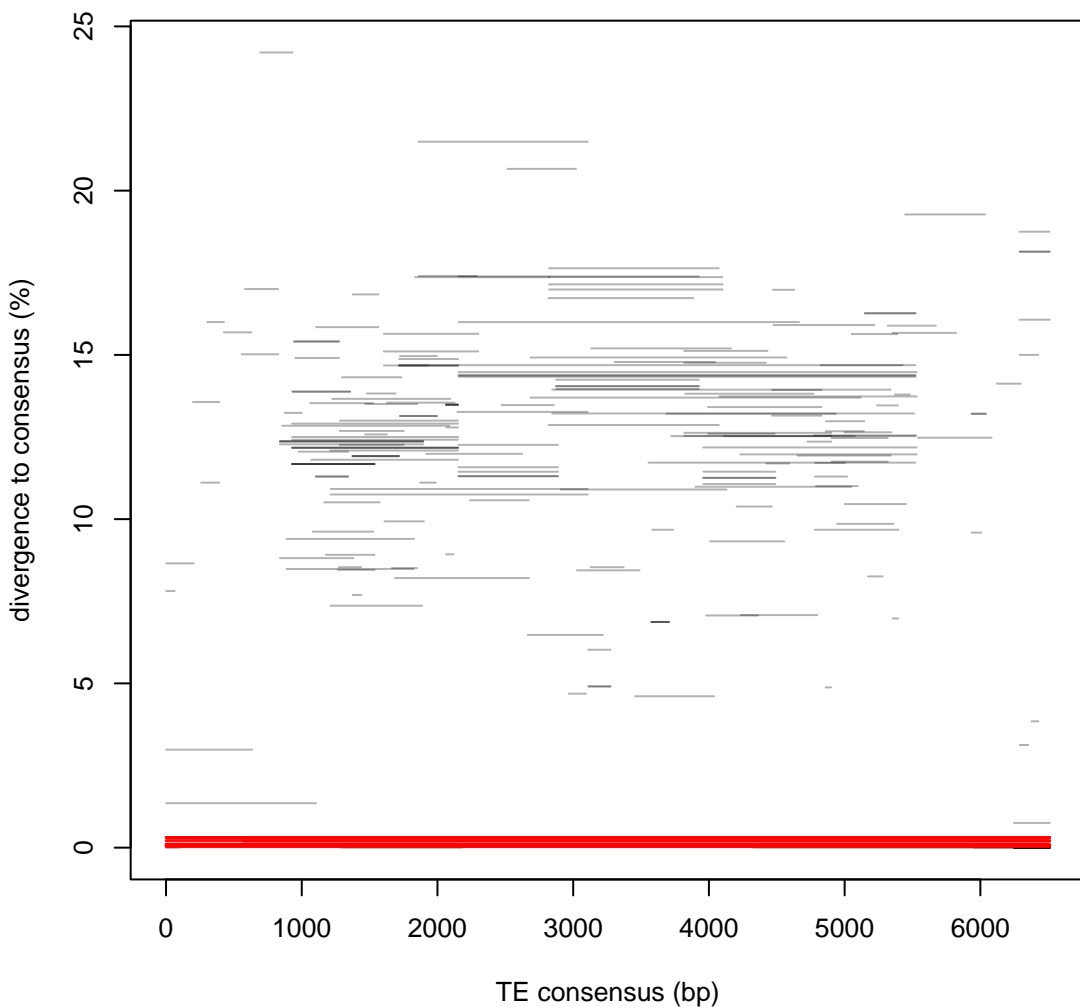
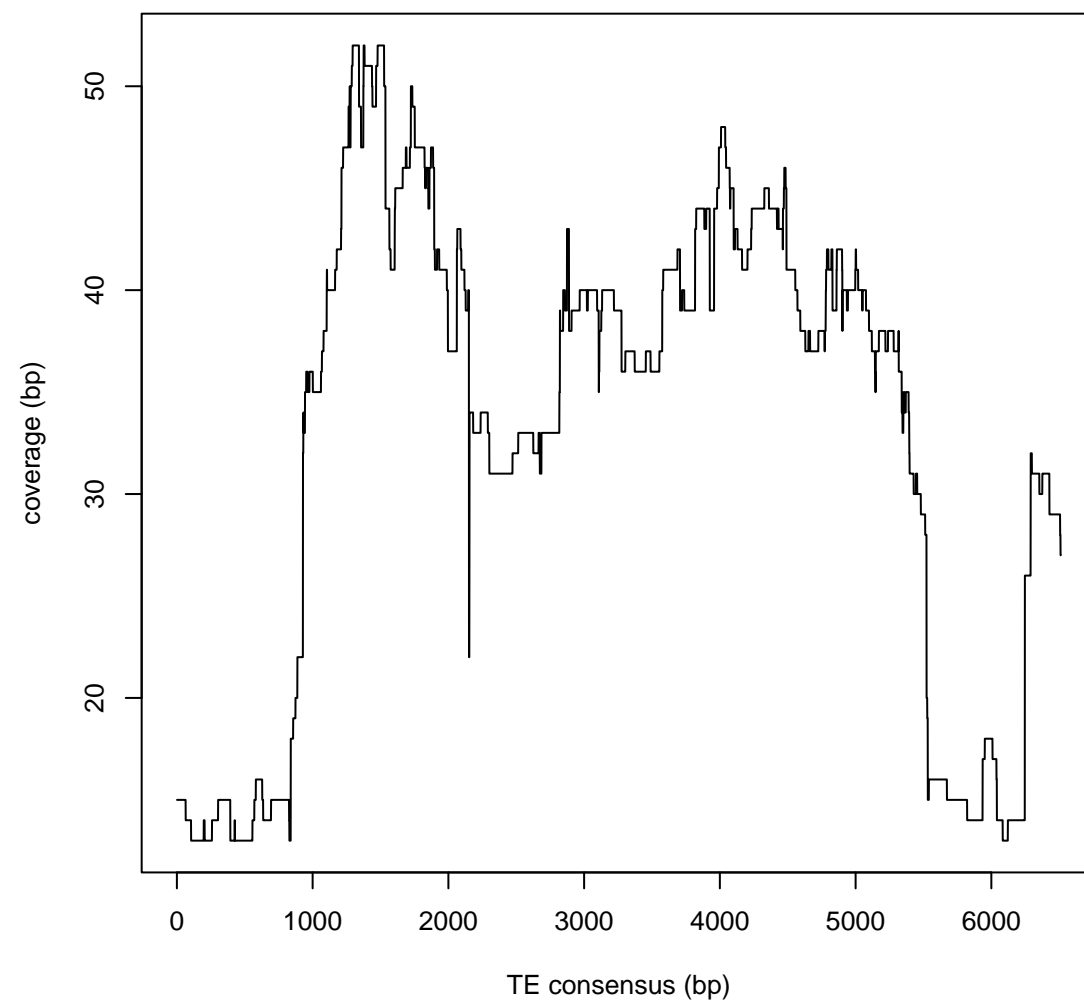


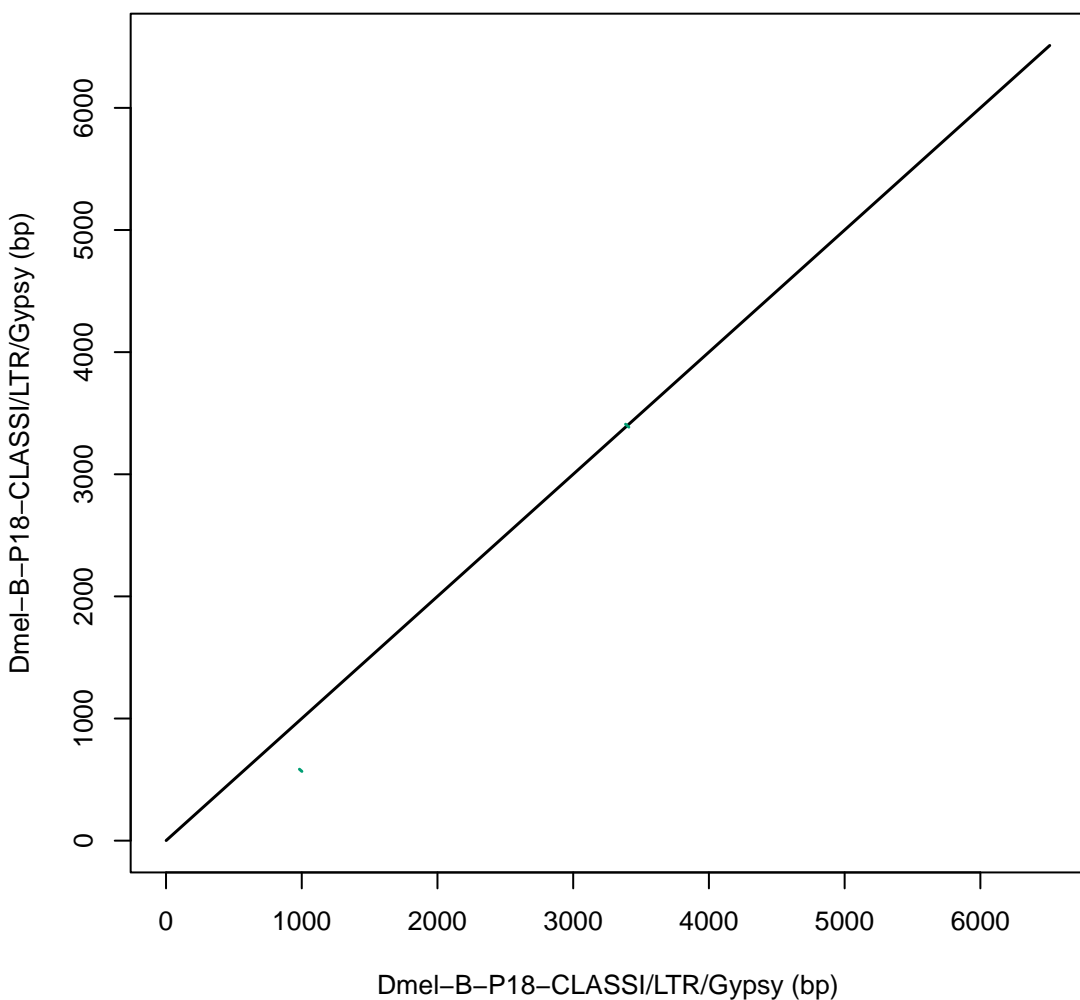
TE: Dmel-B-P18-CLASSI/LTR/Gypsy
consensus size: 6511bp; fragments: 227; full length: 10 (>=5859.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

