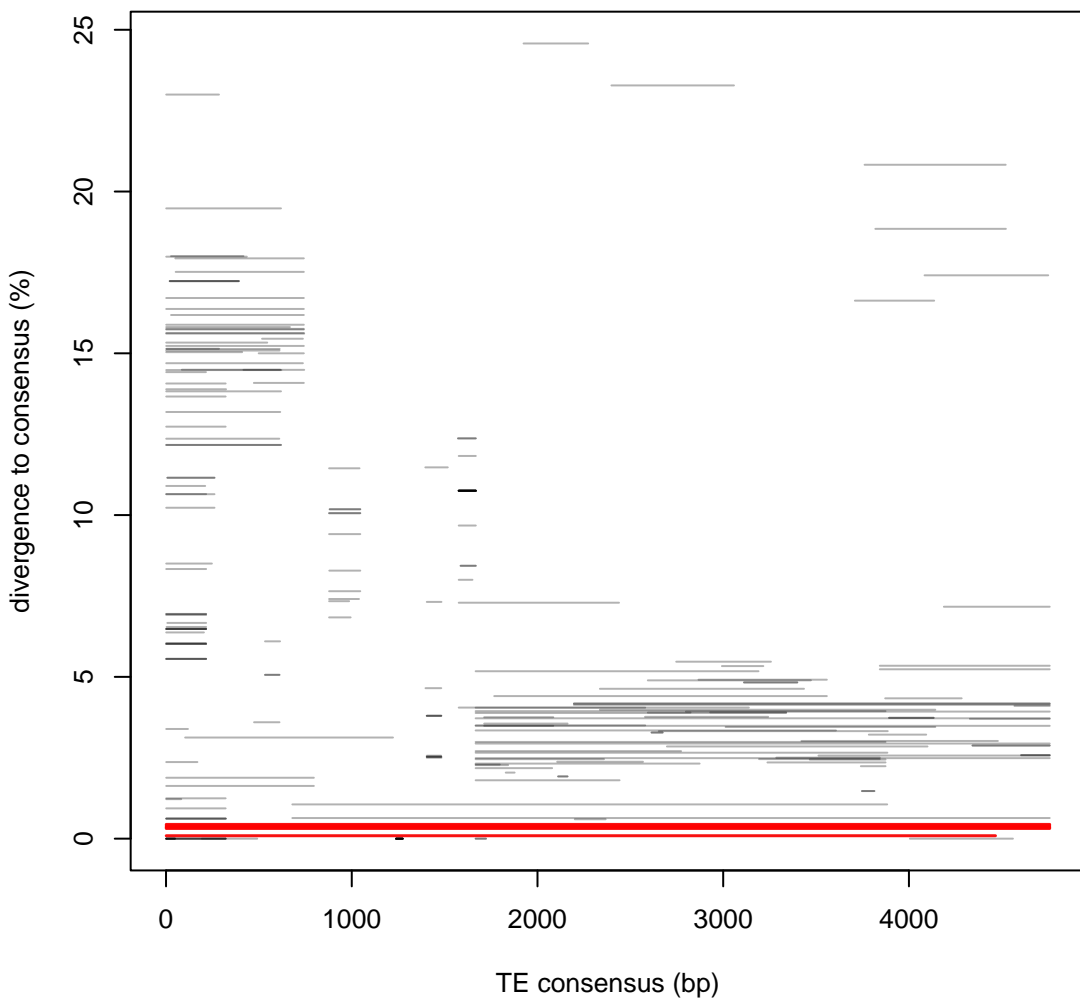
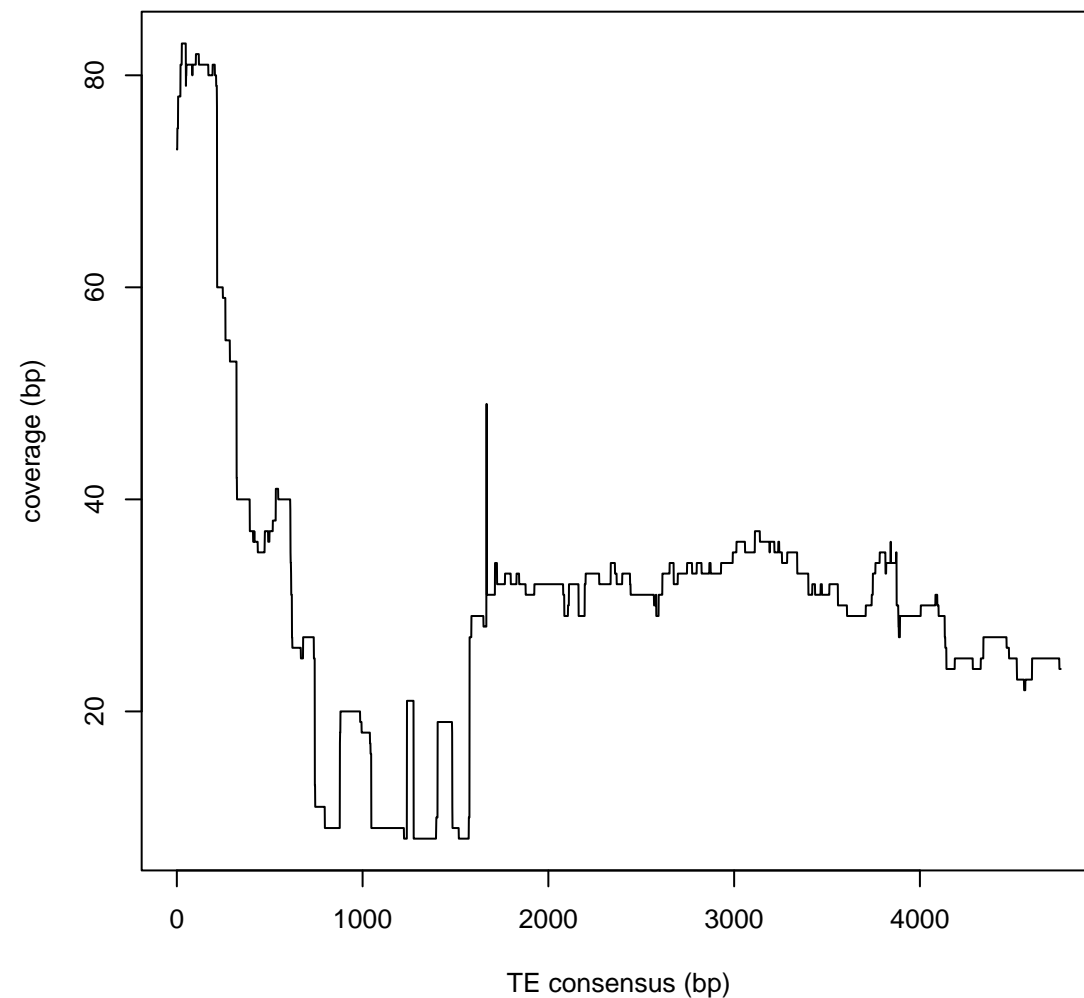


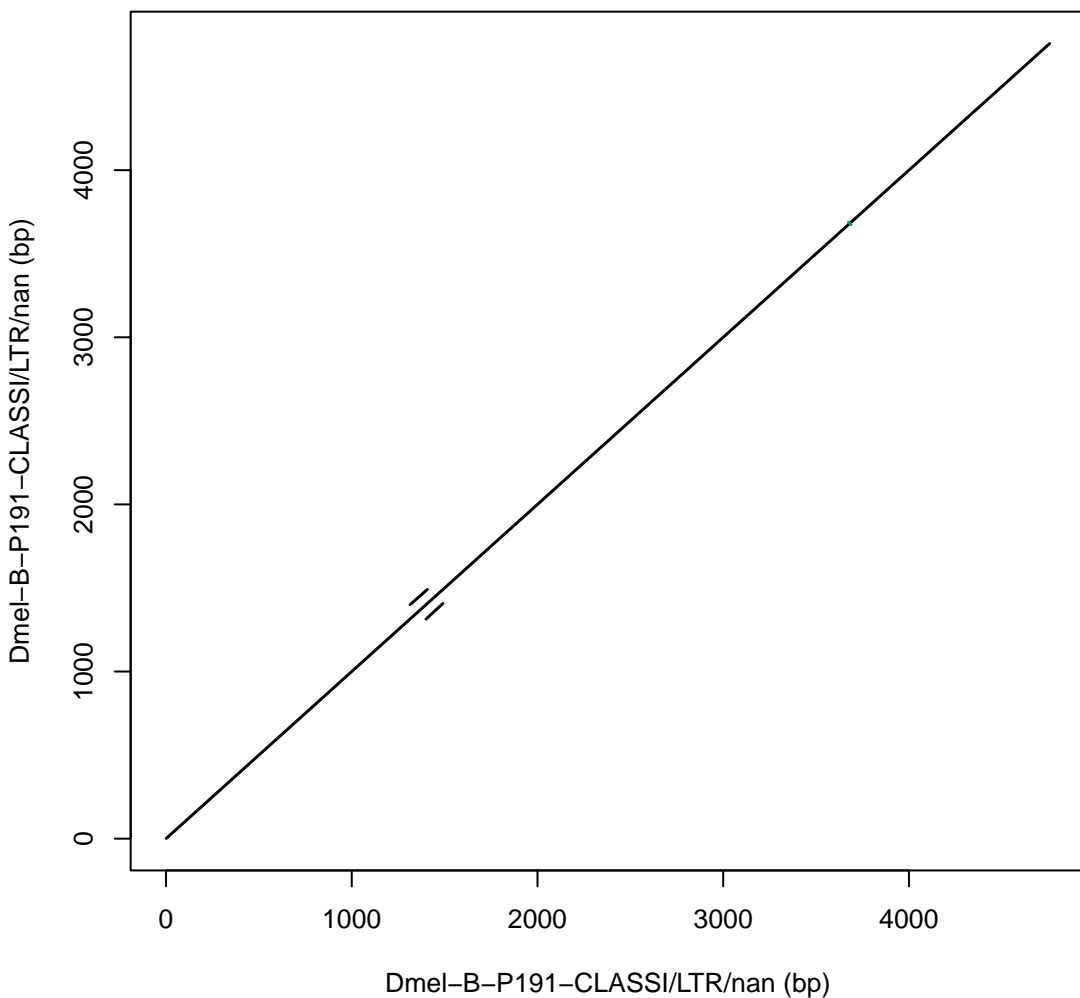
TE: Dmel-B-P191-CLASSI/LTR/nan
consensus size: 4758bp; fragments: 233; full length: 6 (>=4282.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

