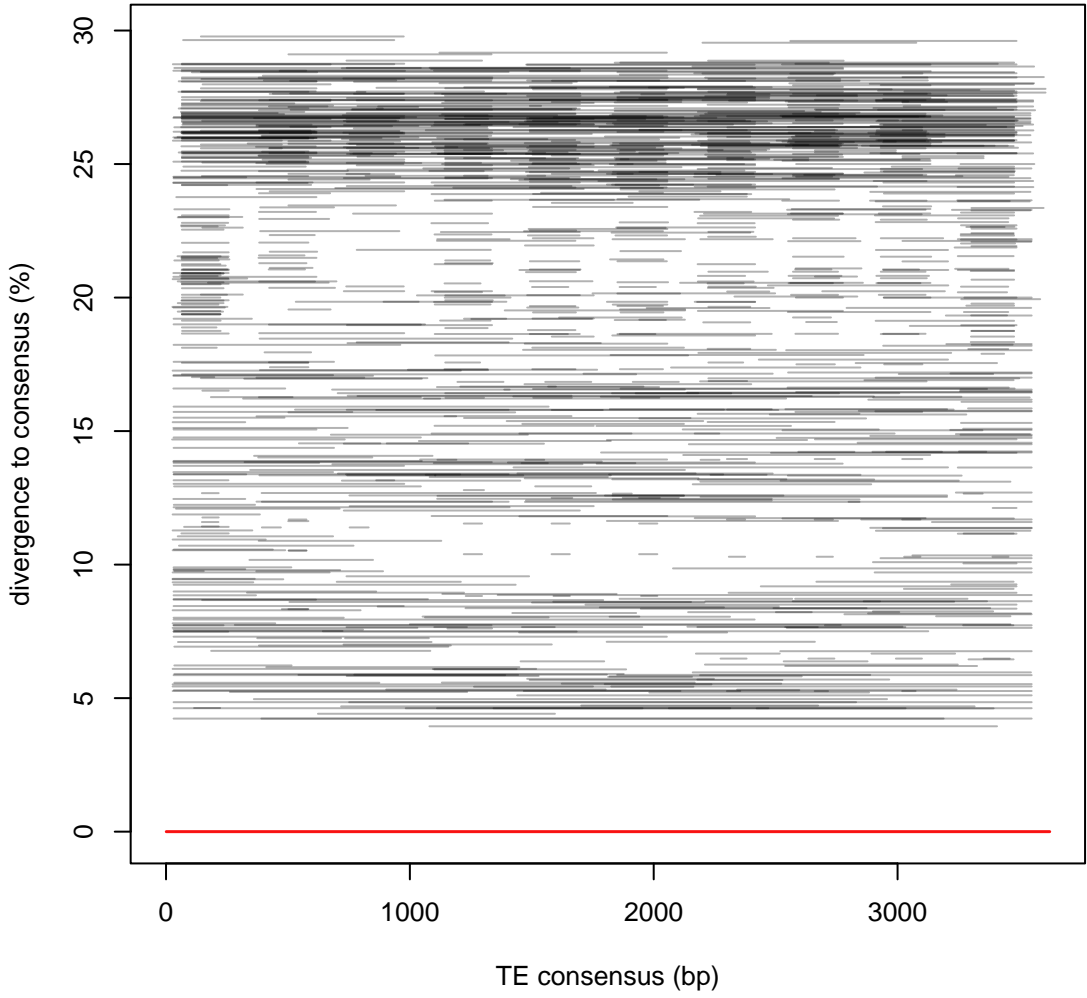
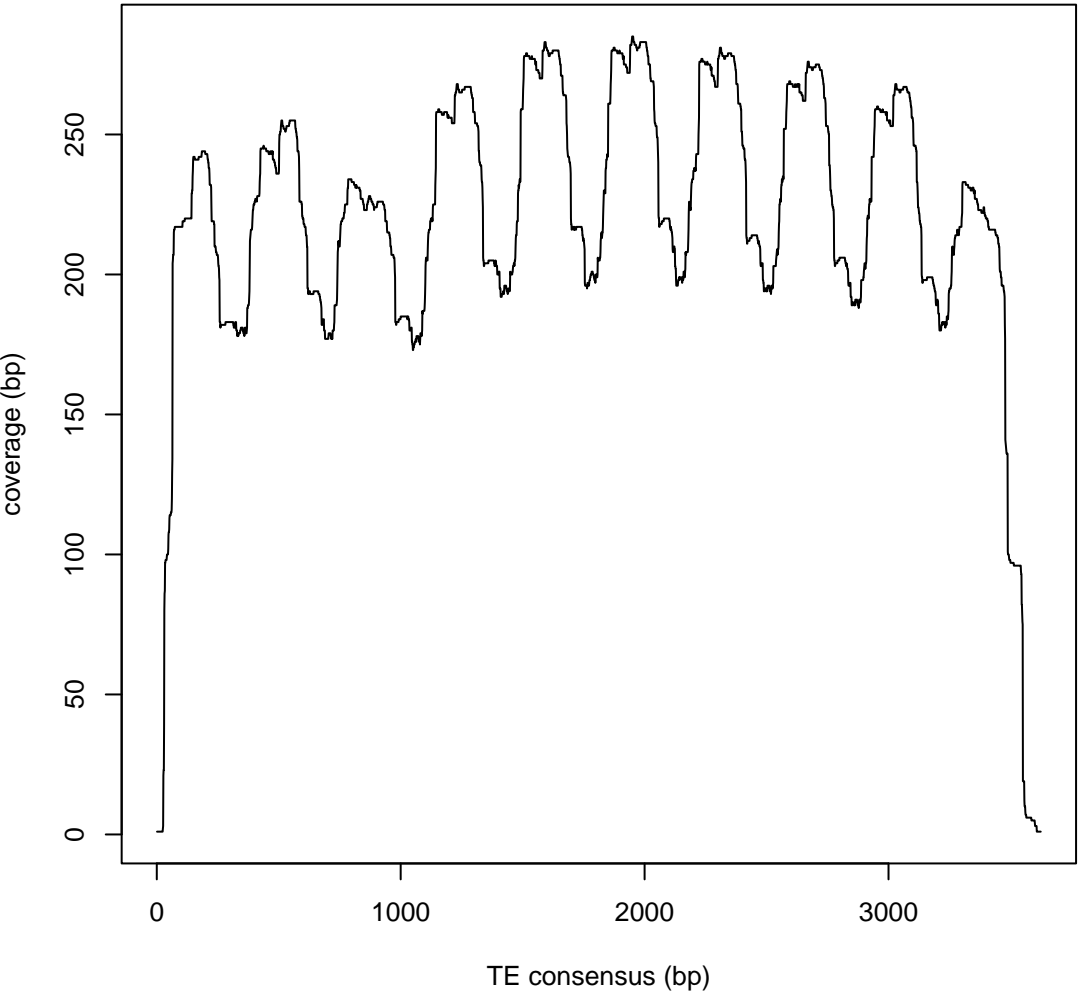


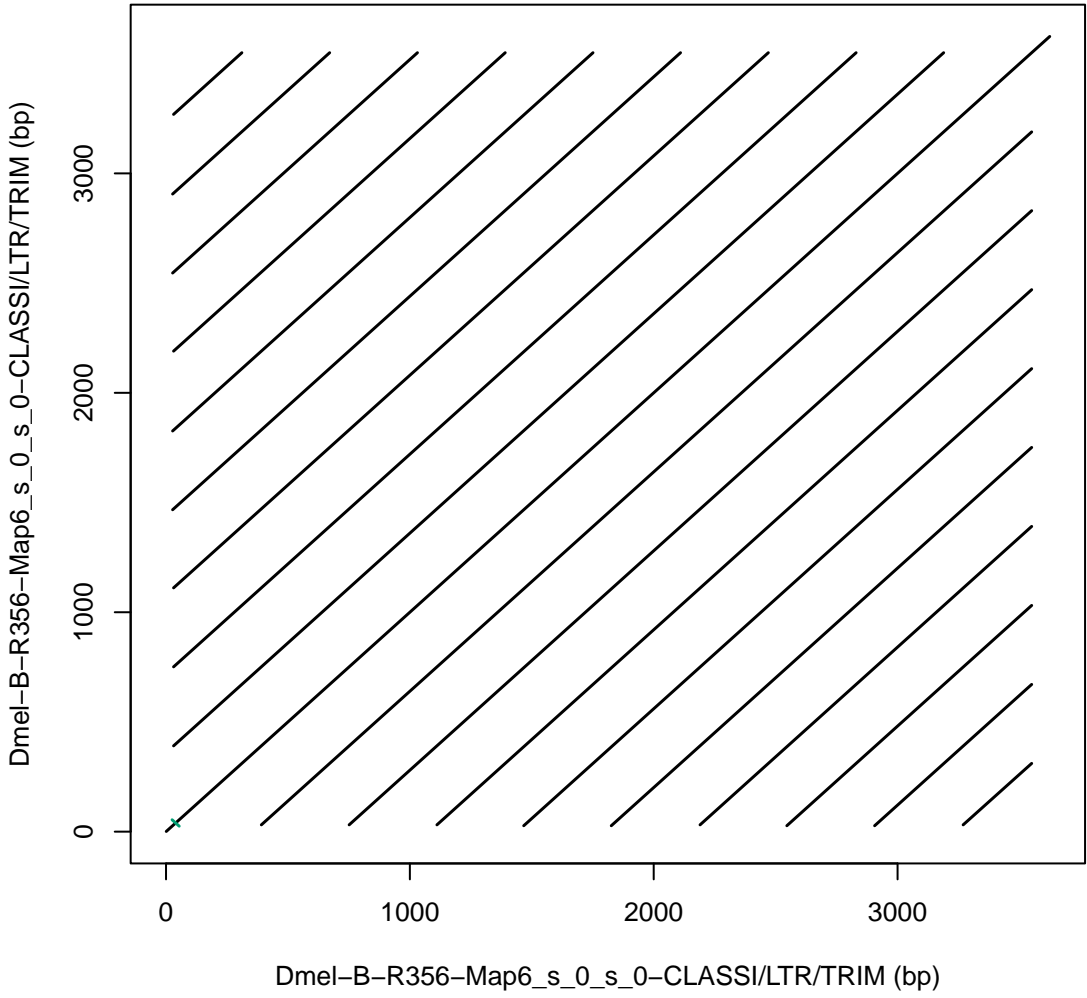
TE: Dmel-B-R356-Map6_s_0_s_0-CLASSI/LTR/TRIM
consensus size: 3624bp; fragments: 1223; full length: 1 (>=3261.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

