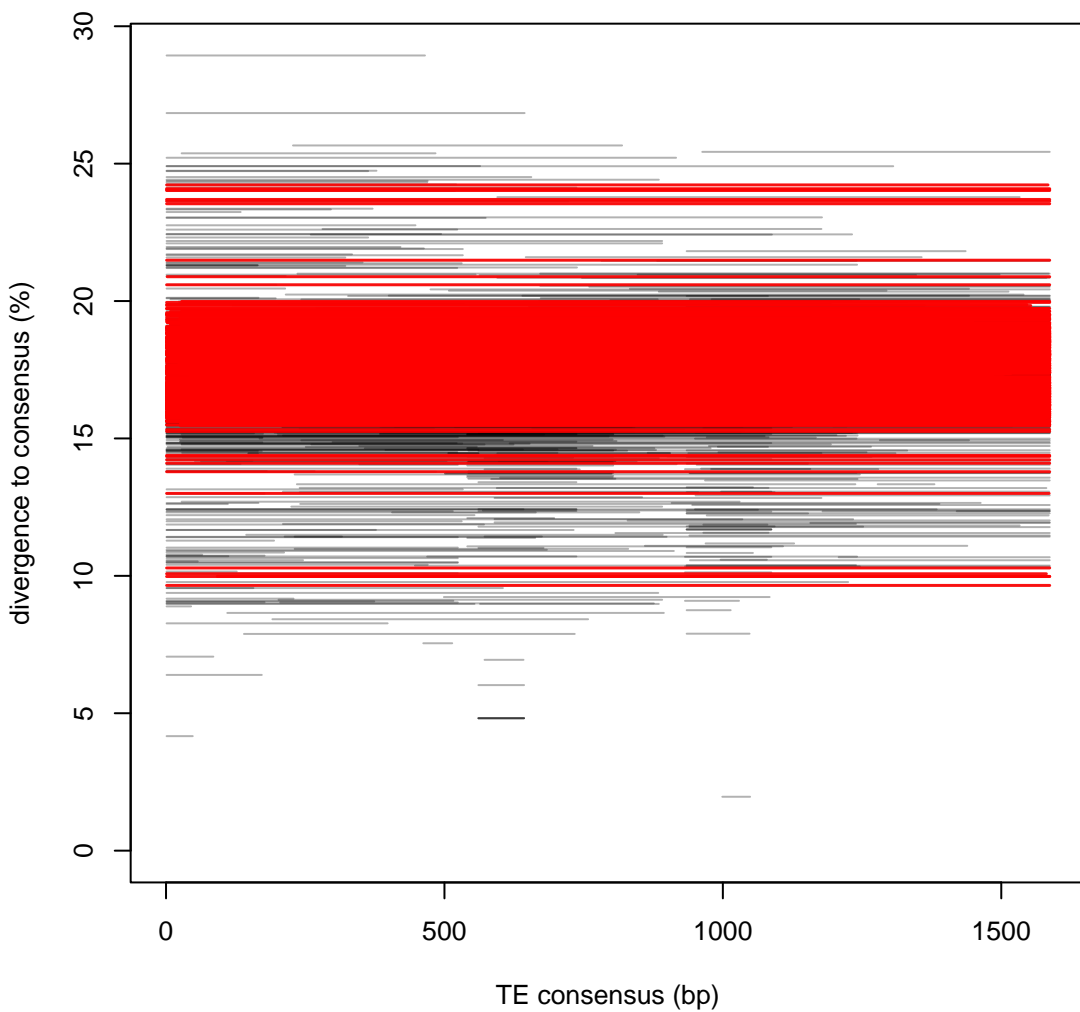
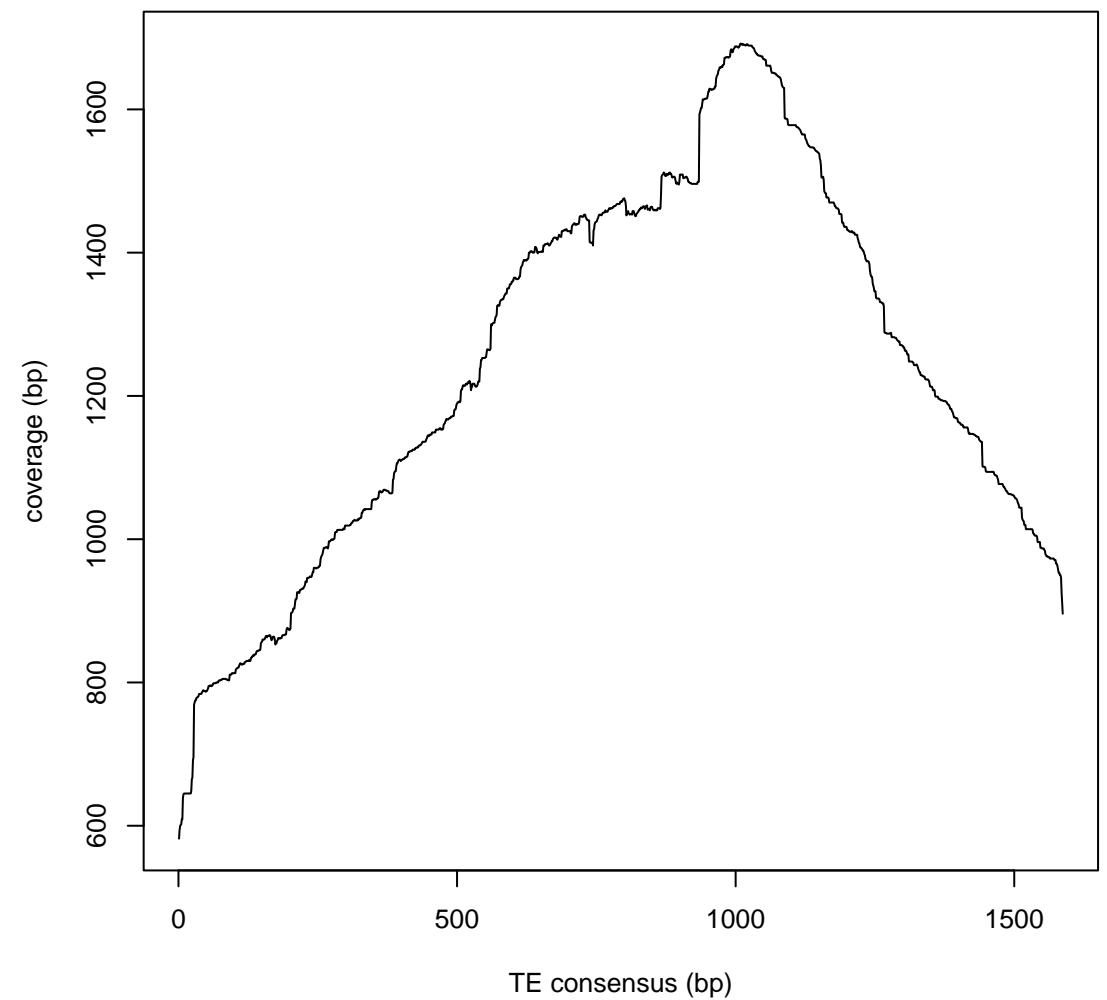


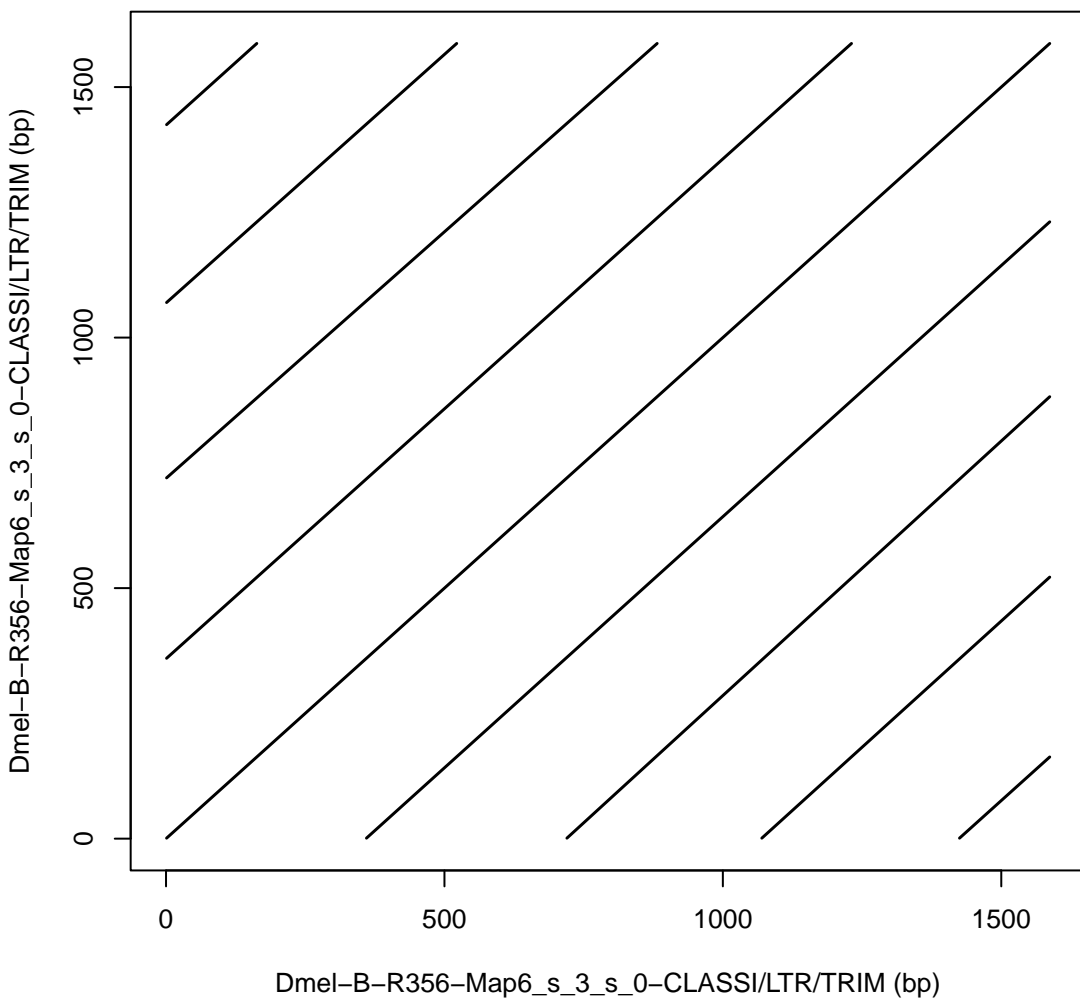
TE: Dmel-B-R356-Map6_s_3_s_0-CLASSI/LTR/TRIM
consensus size: 1587bp; fragments: 2054; full length: 352 (>=1428.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

