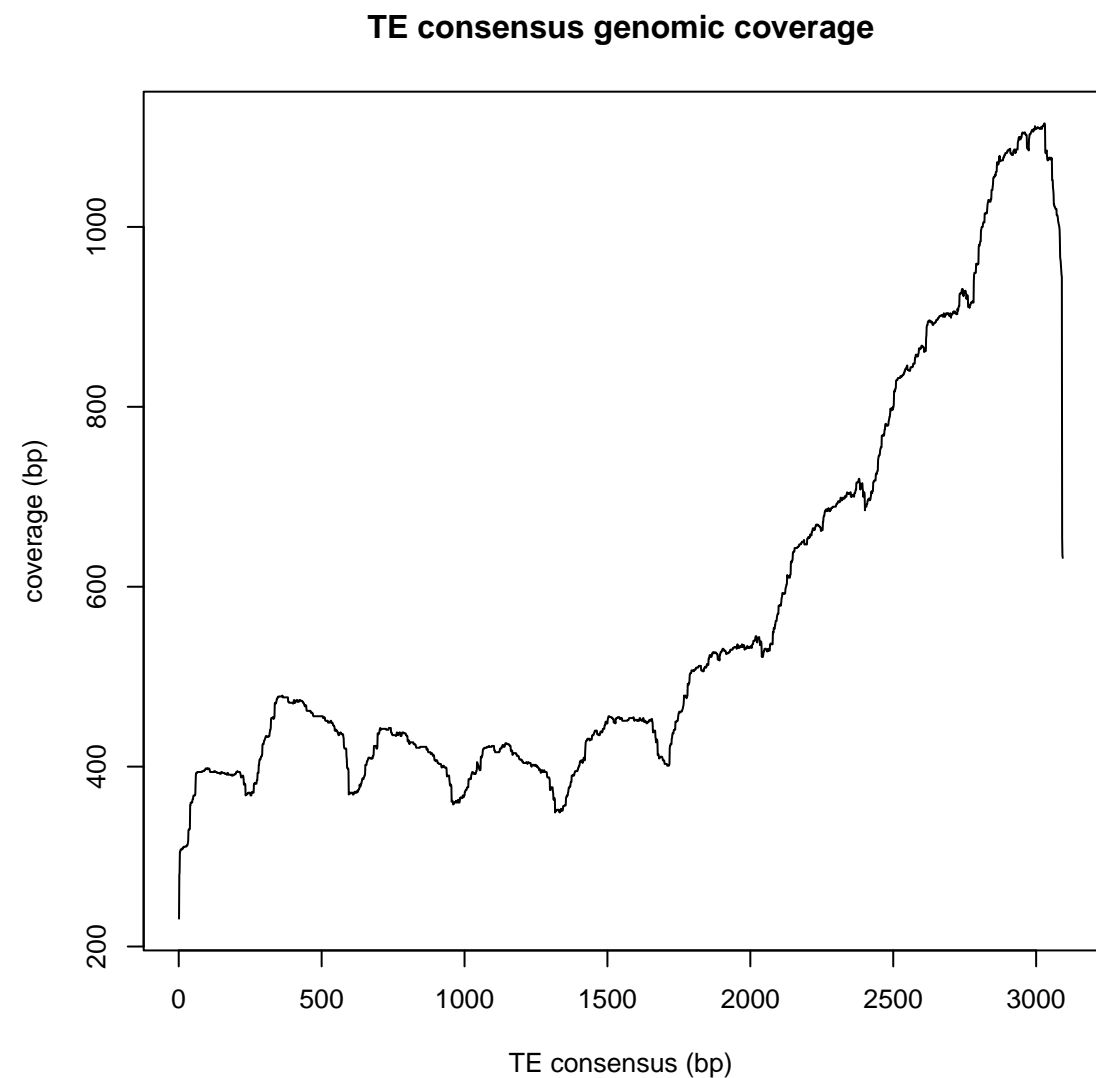
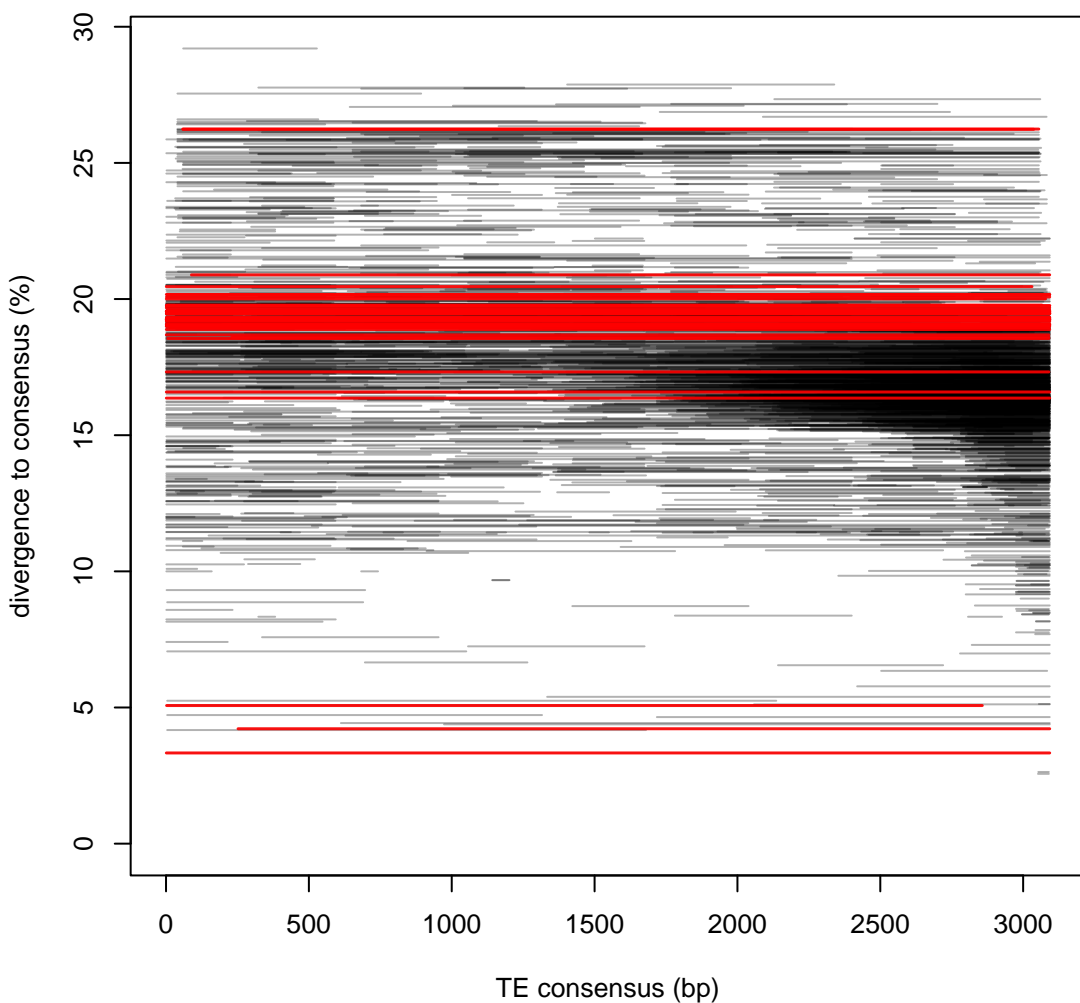
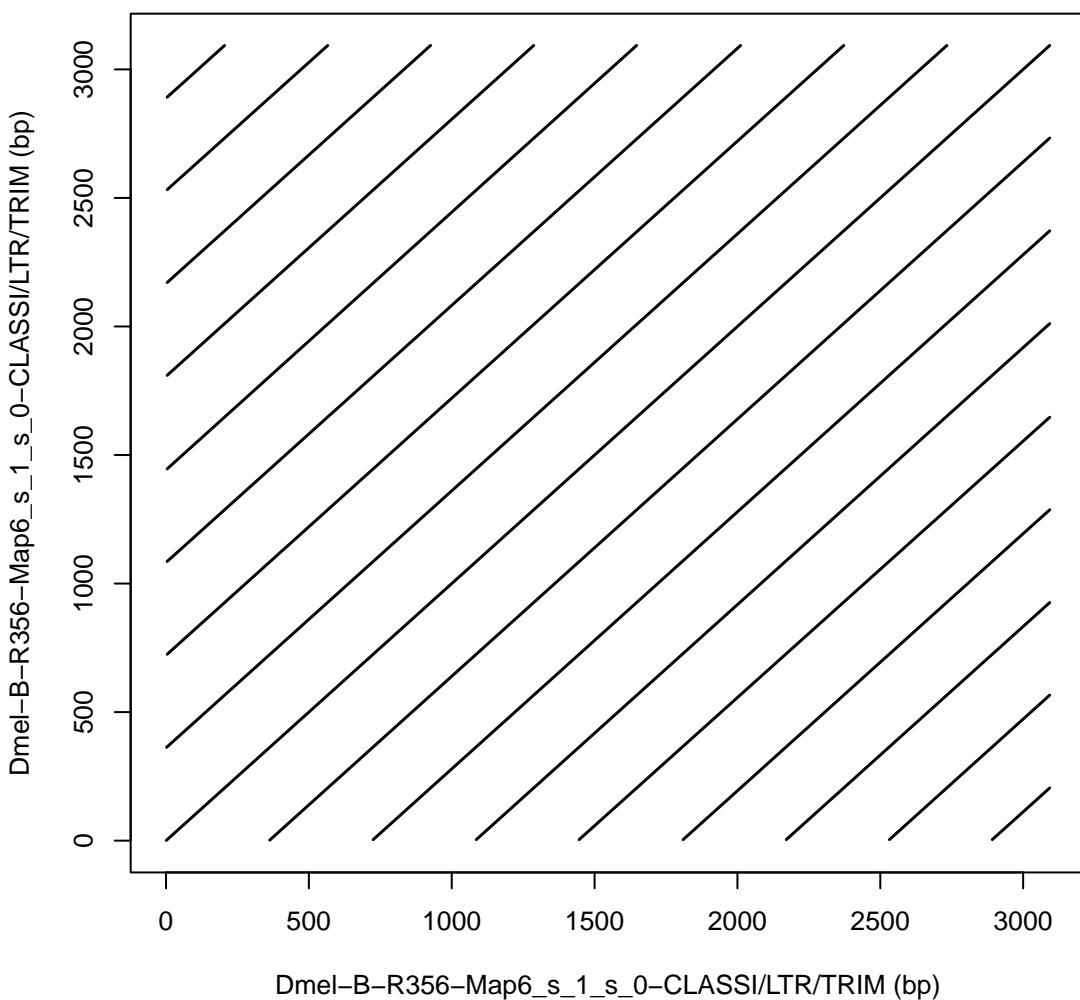


TE: Dmel-B-R356-Map6_s_1_s_0-CLASSI/LTR/TRIM
consensus size: 3093bp; fragments: 2154; full length: 48 (>=2783.7bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

