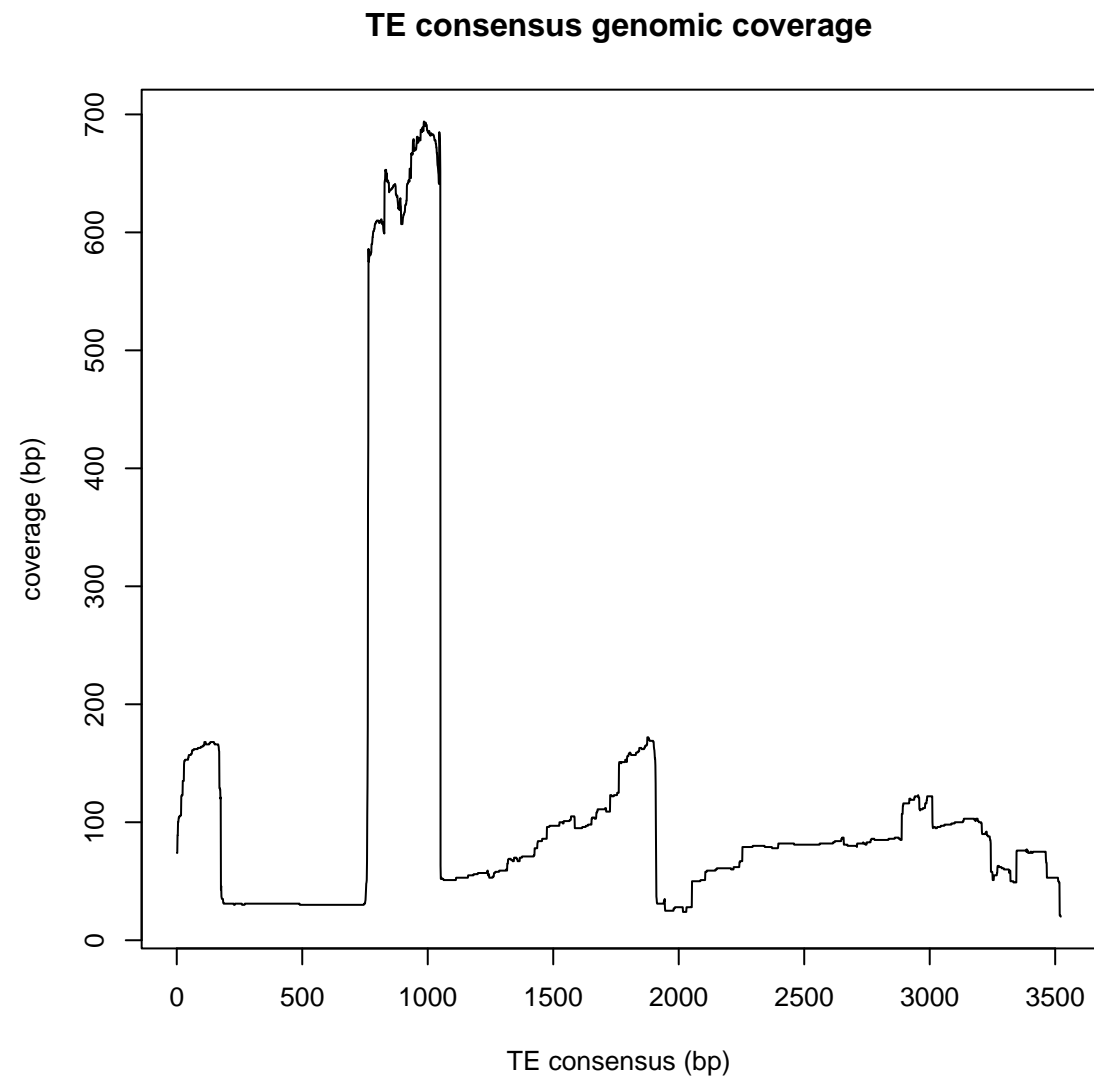
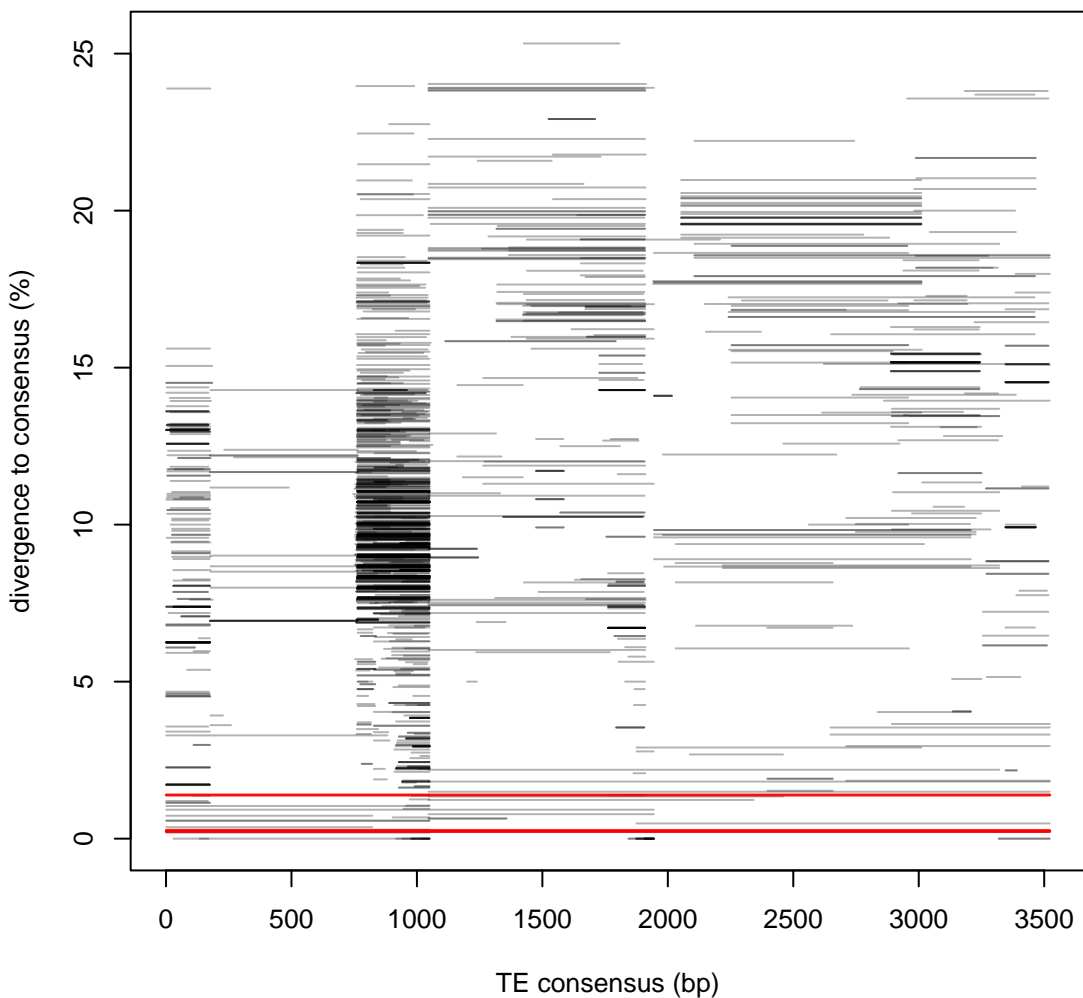
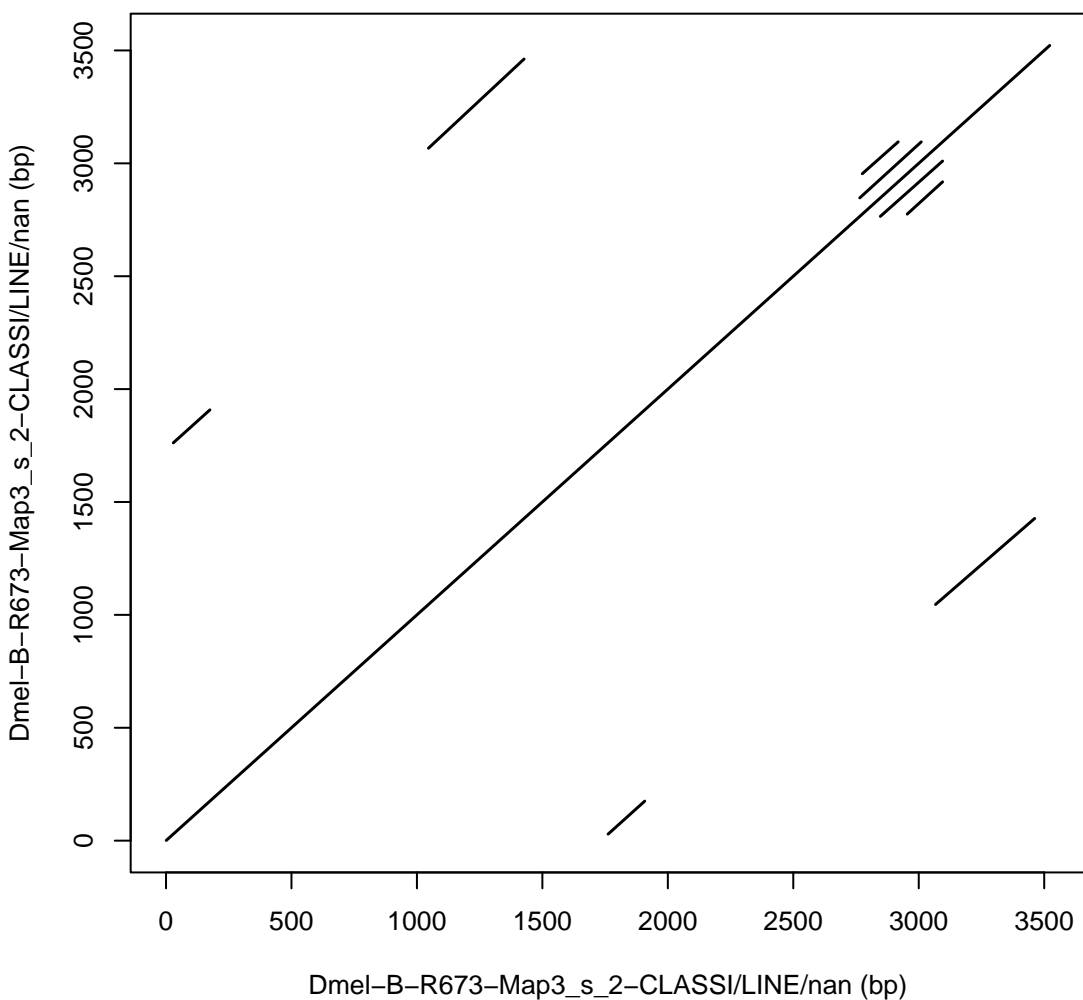


TE: Dmel-B-R673-Map3_s_2-CLASSI/LINE/nan
consensus size: 3522bp; fragments: 1446; full length: 3 (≥ 3169.8 bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

