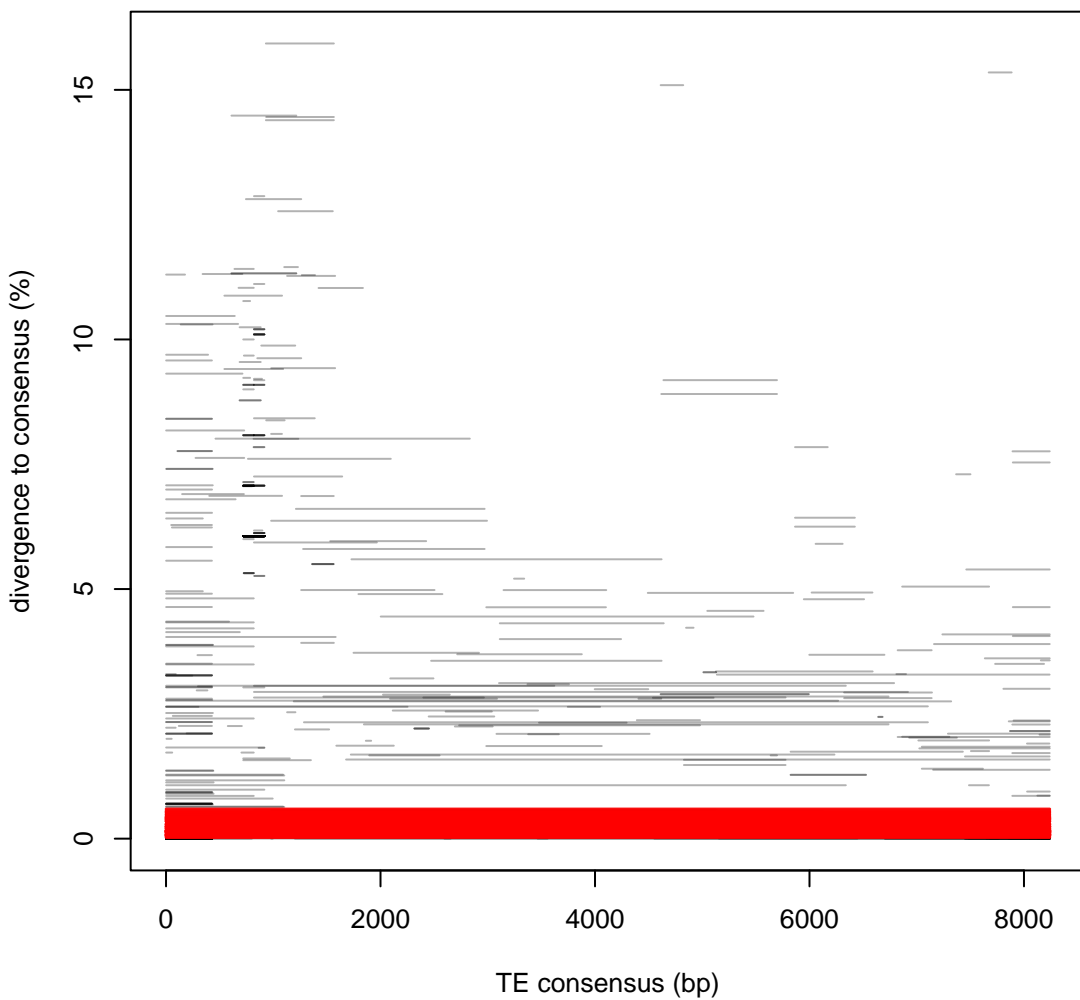
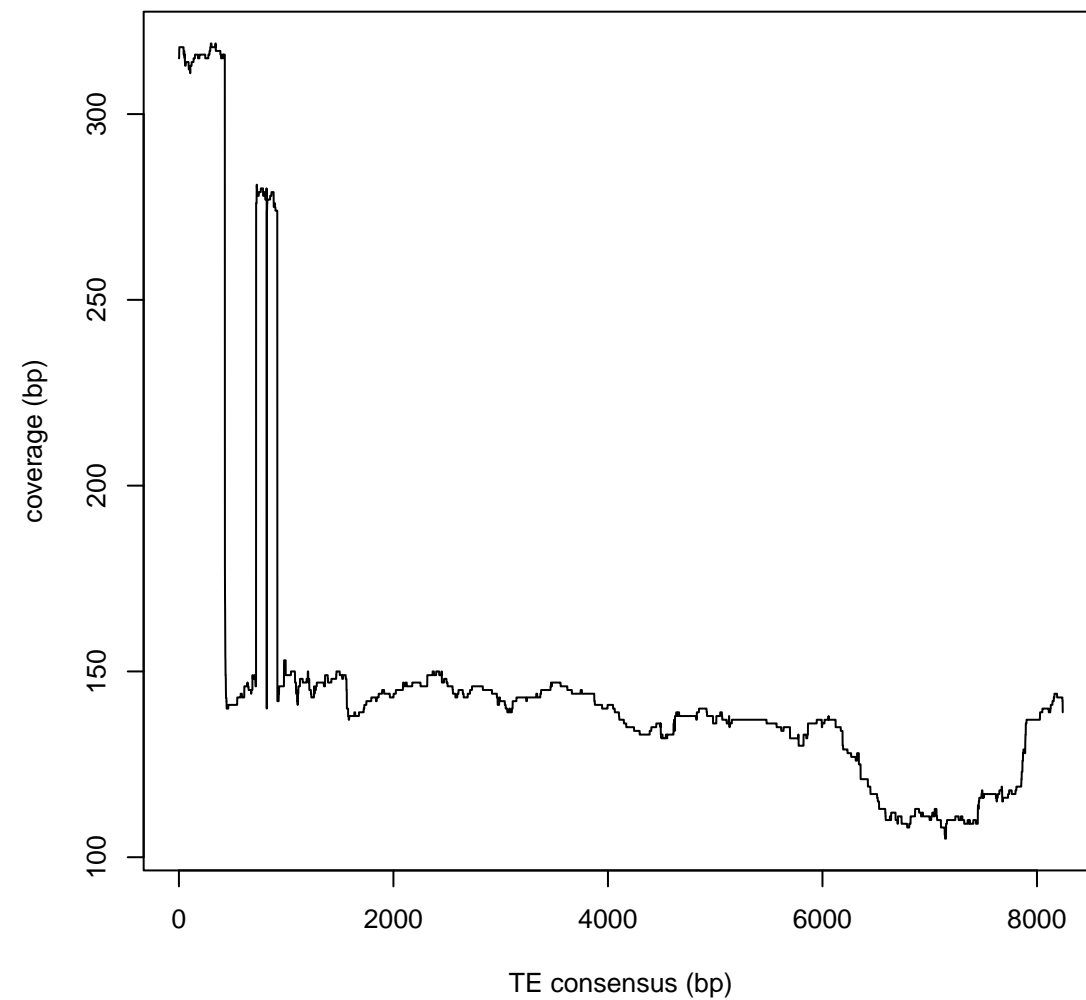


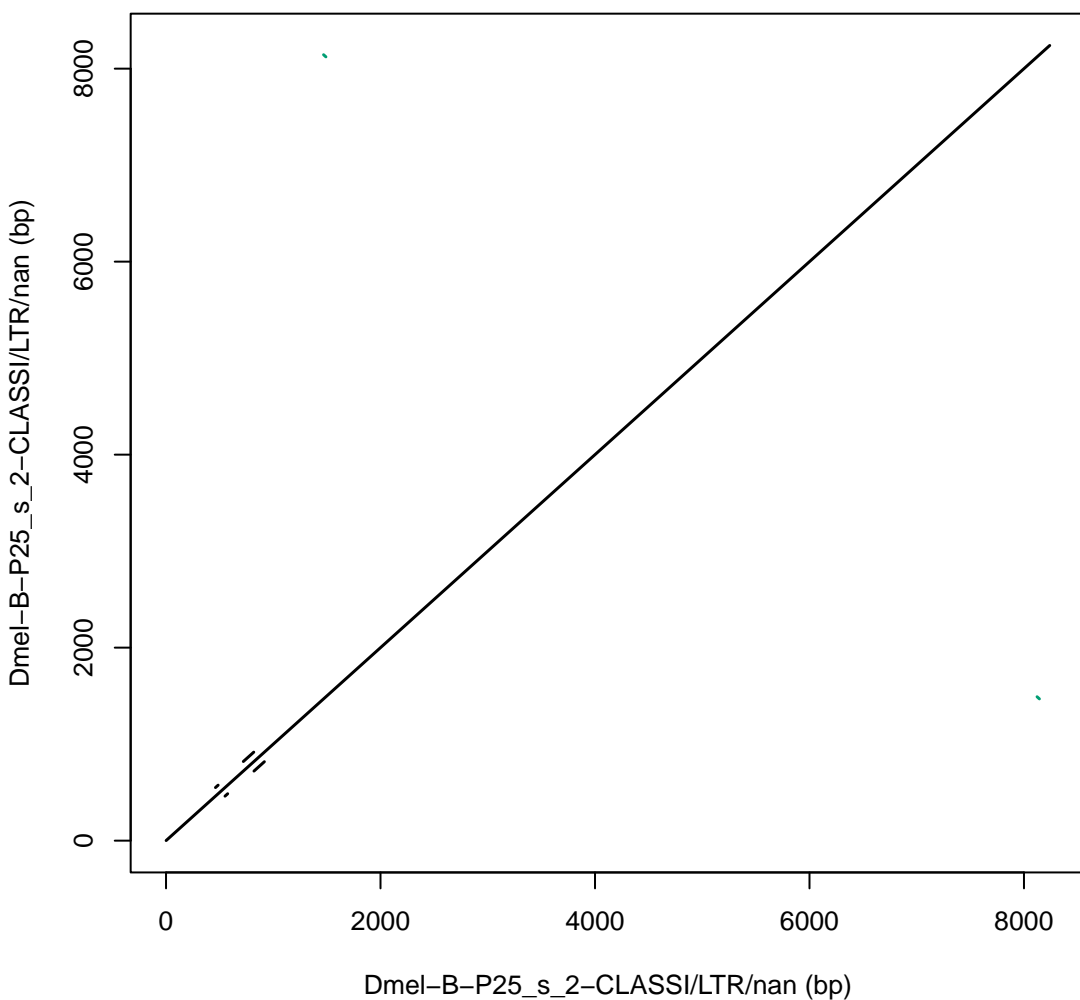
TE: Dmel-B-P25_s_2-CLASSI/LTR/nan
consensus size: 8240bp; fragments: 825; full length: 69 (>=7416bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

