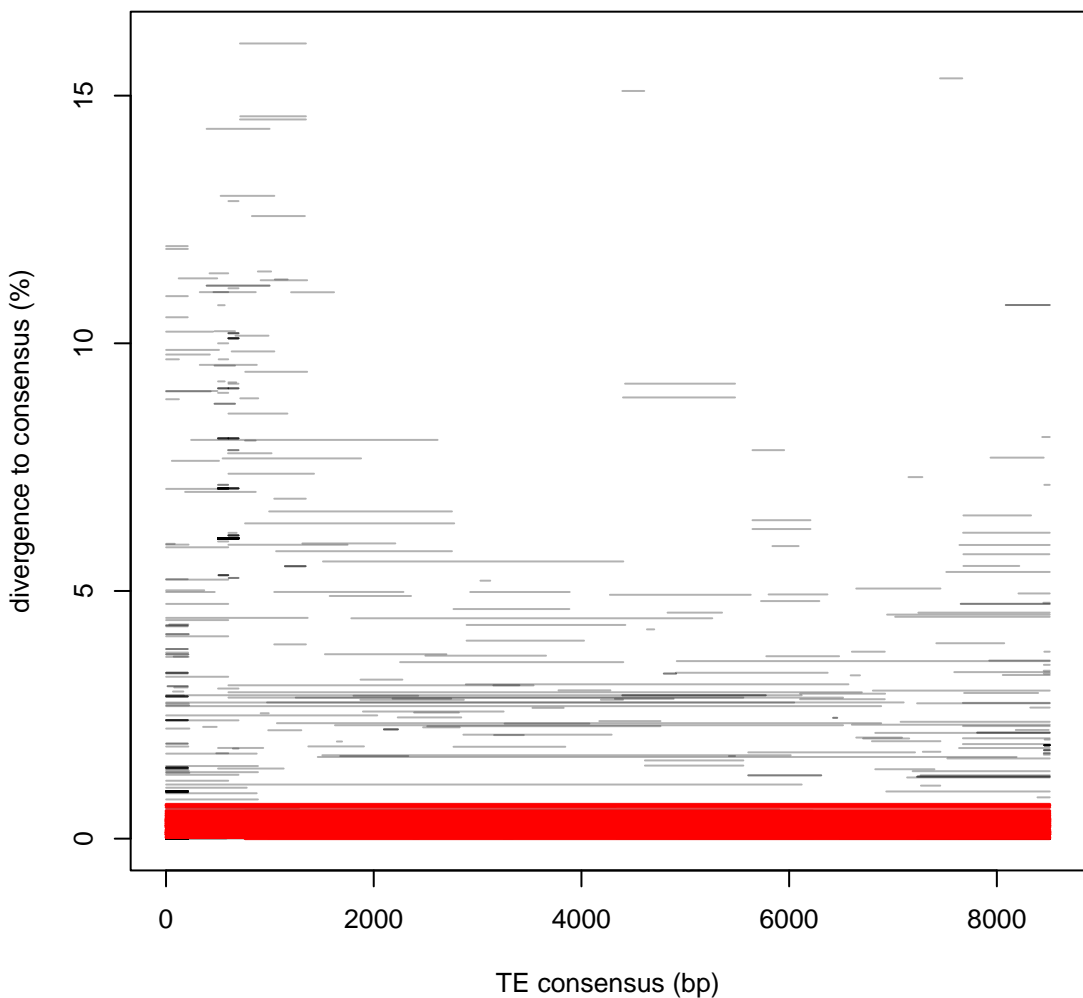
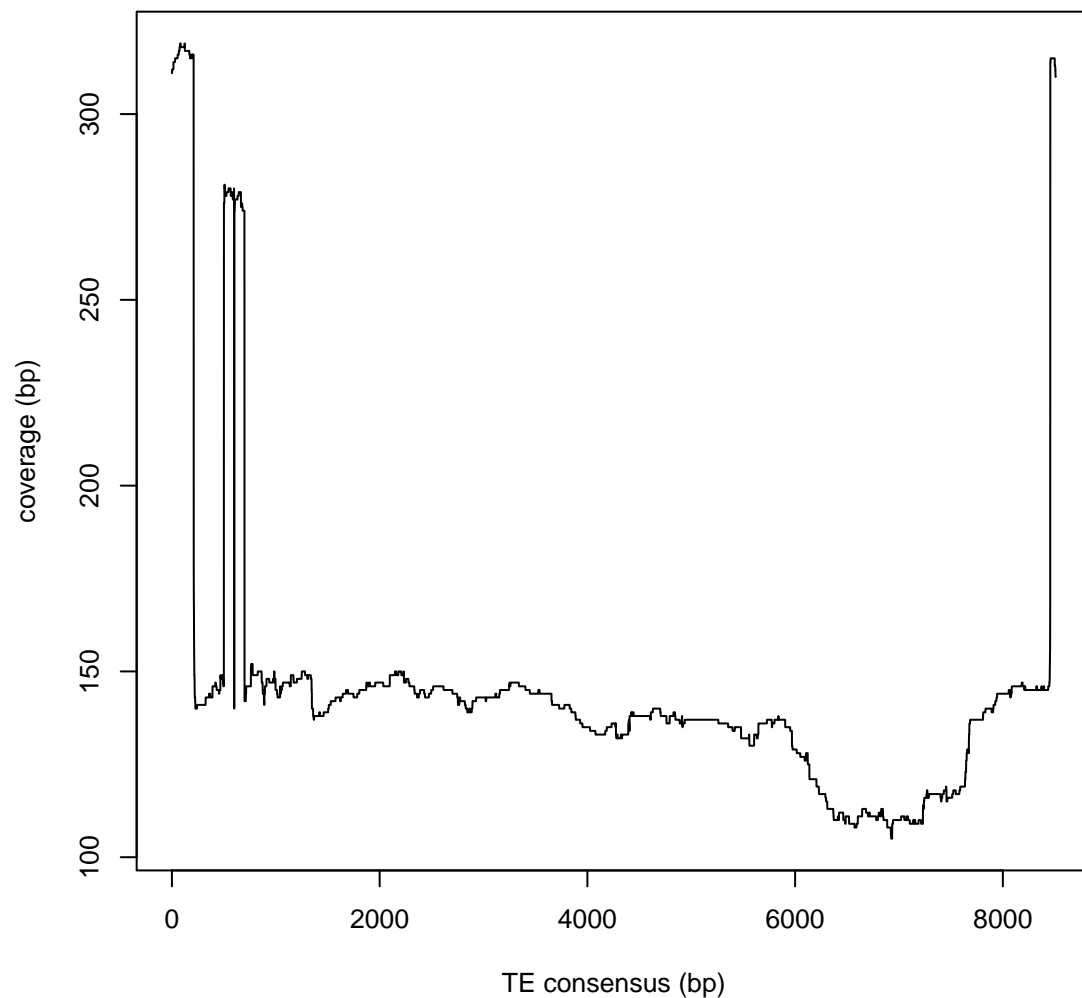


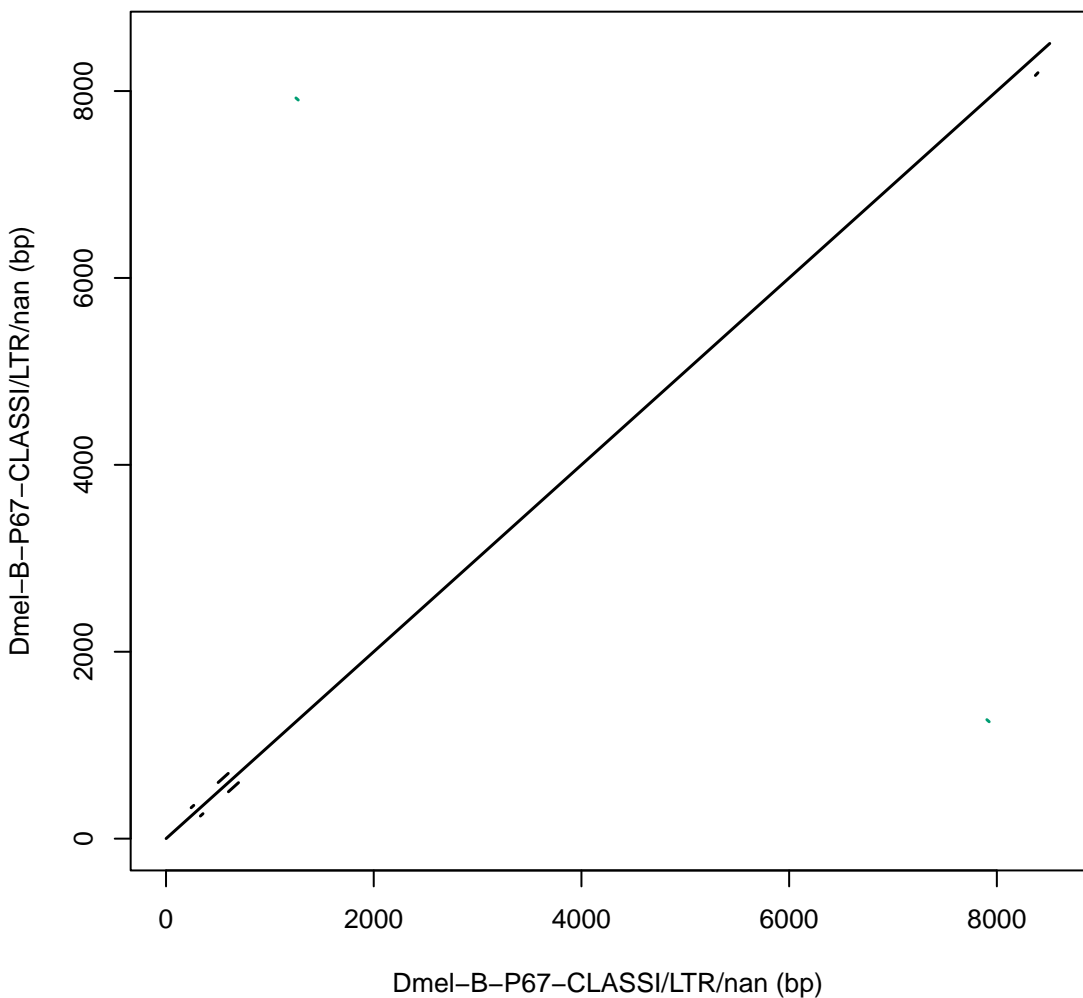
TE: Dmel-B-P67-CLASSI/LTR/nan
consensus size: 8509bp; fragments: 992; full length: 70 (≥ 7658.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

