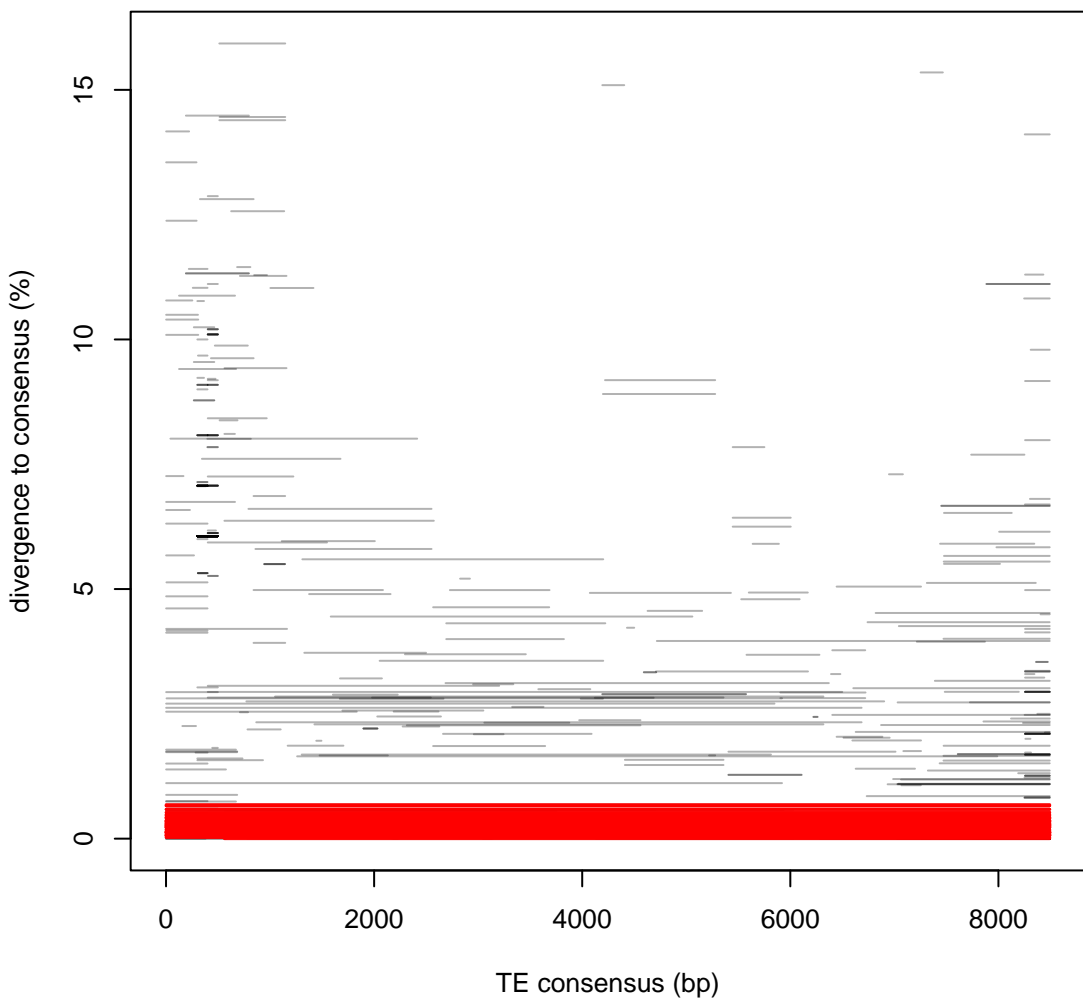
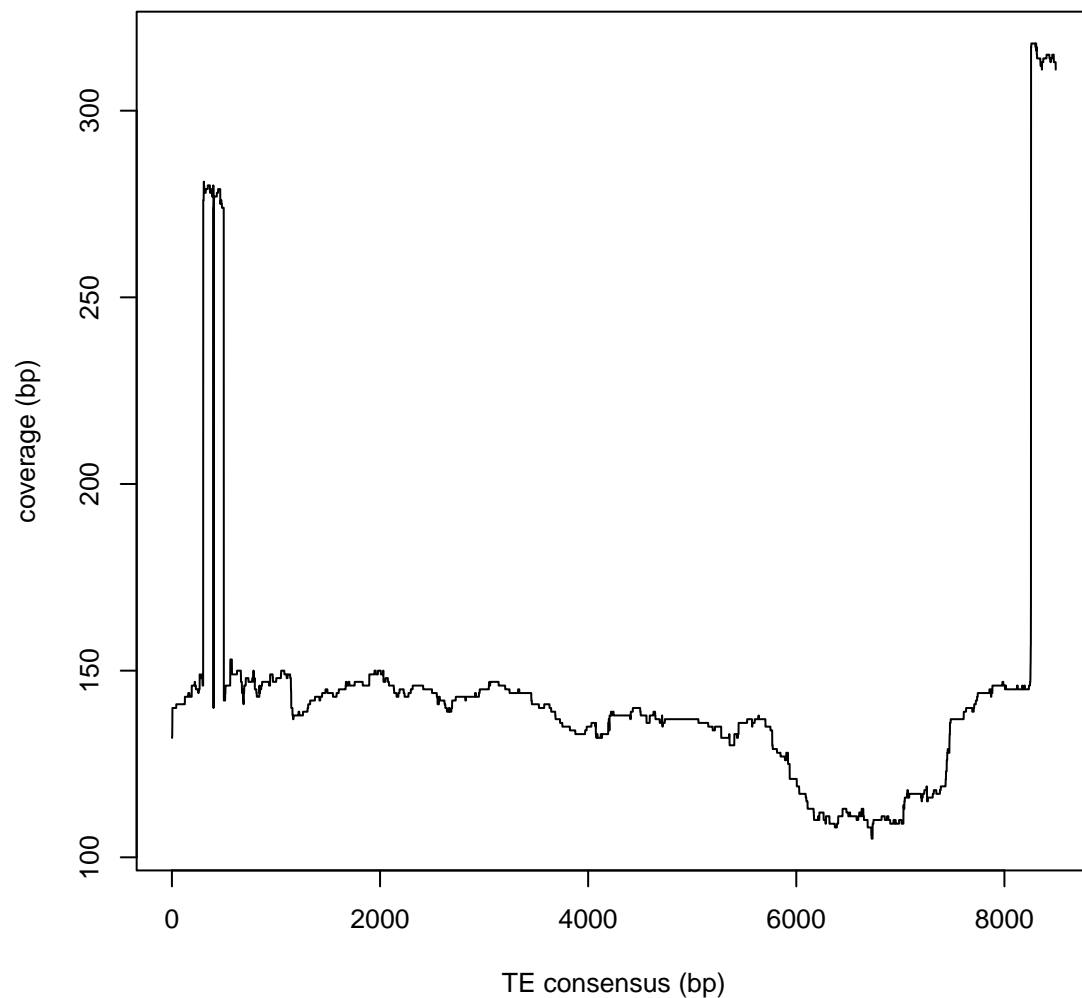


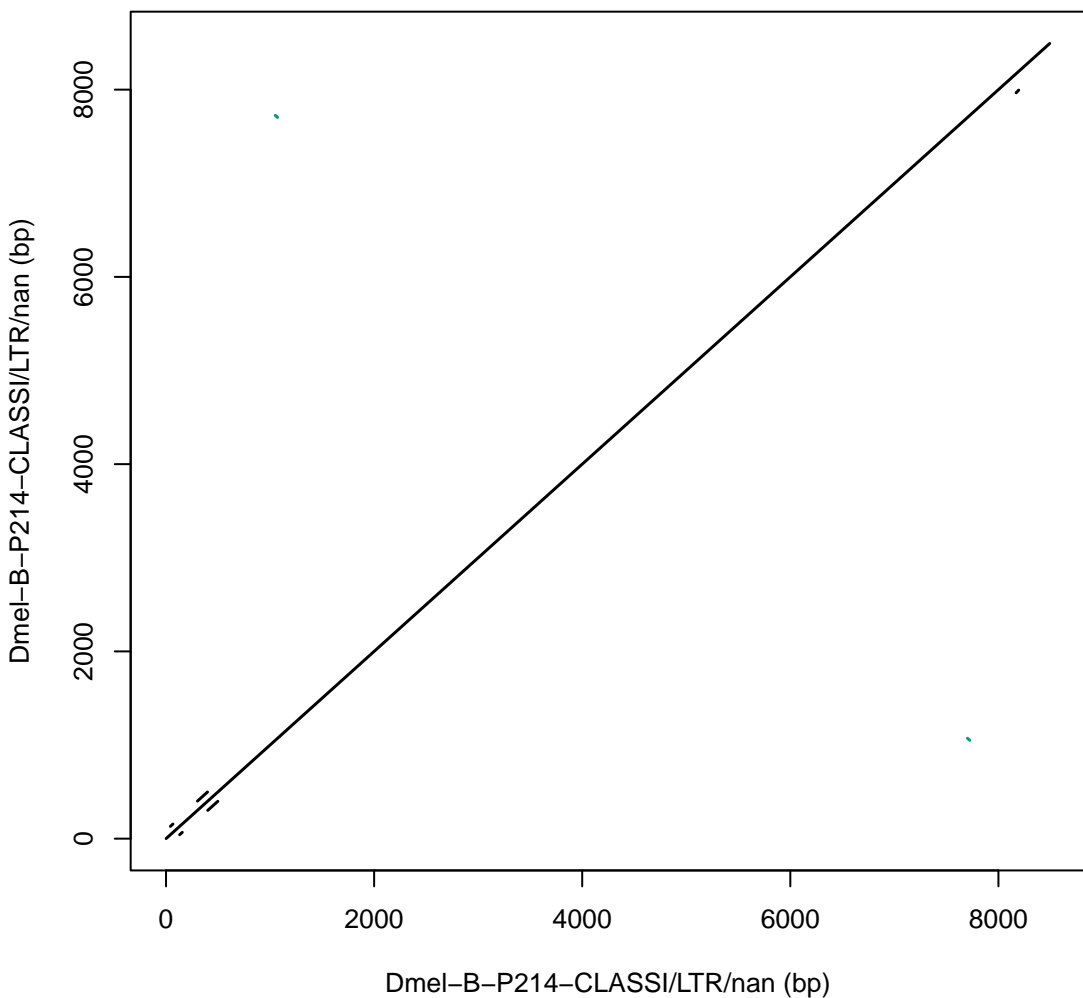
TE: Dmel-B-P214-CLASSI/LTR/nan
consensus size: 8493bp; fragments: 822; full length: 72 (≥ 7643.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

