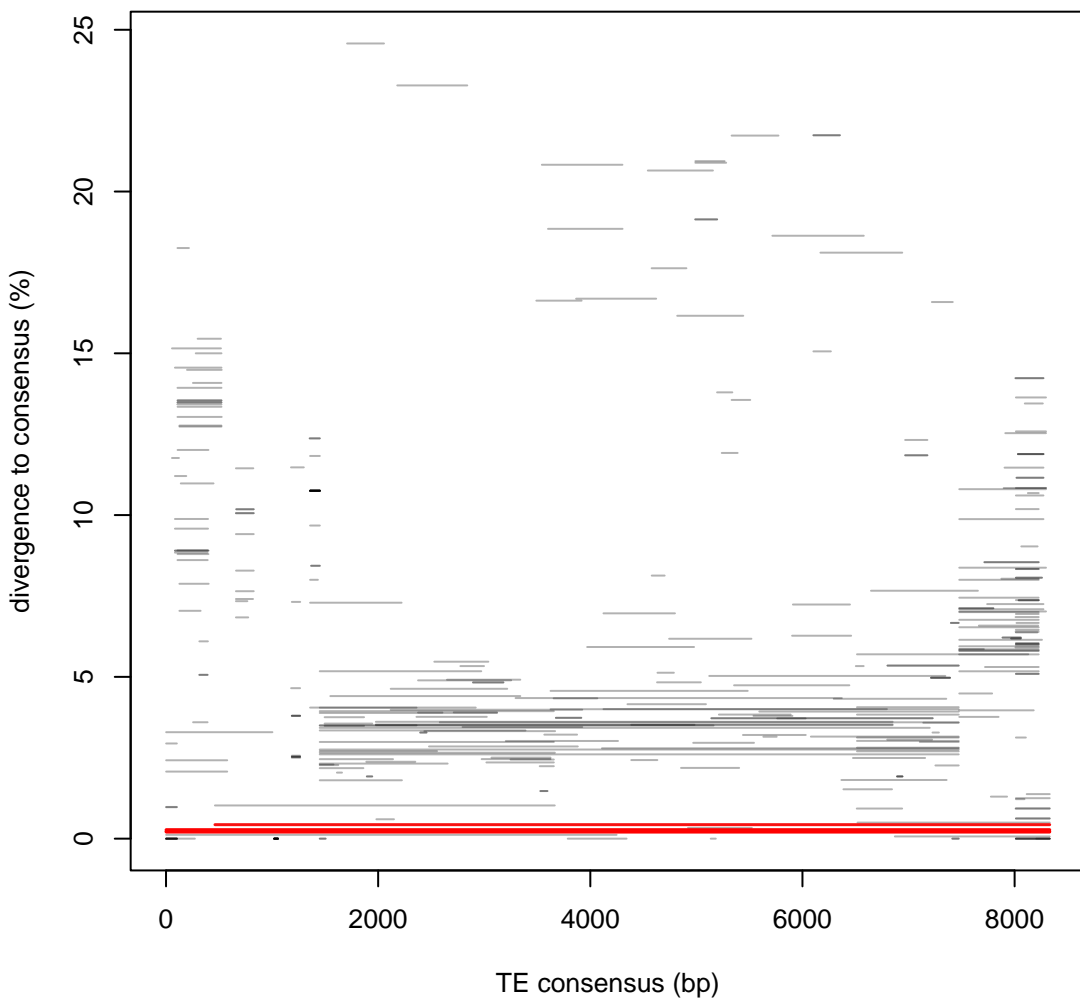
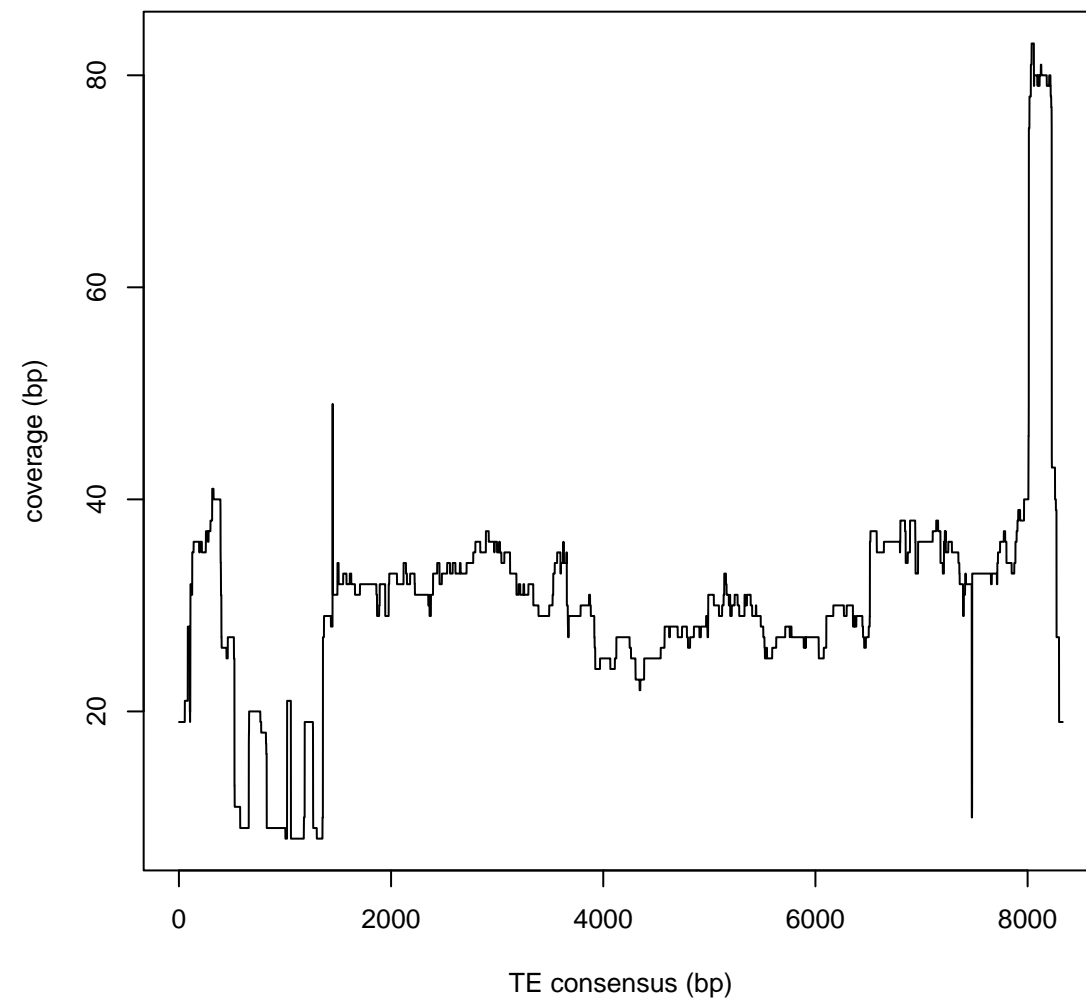


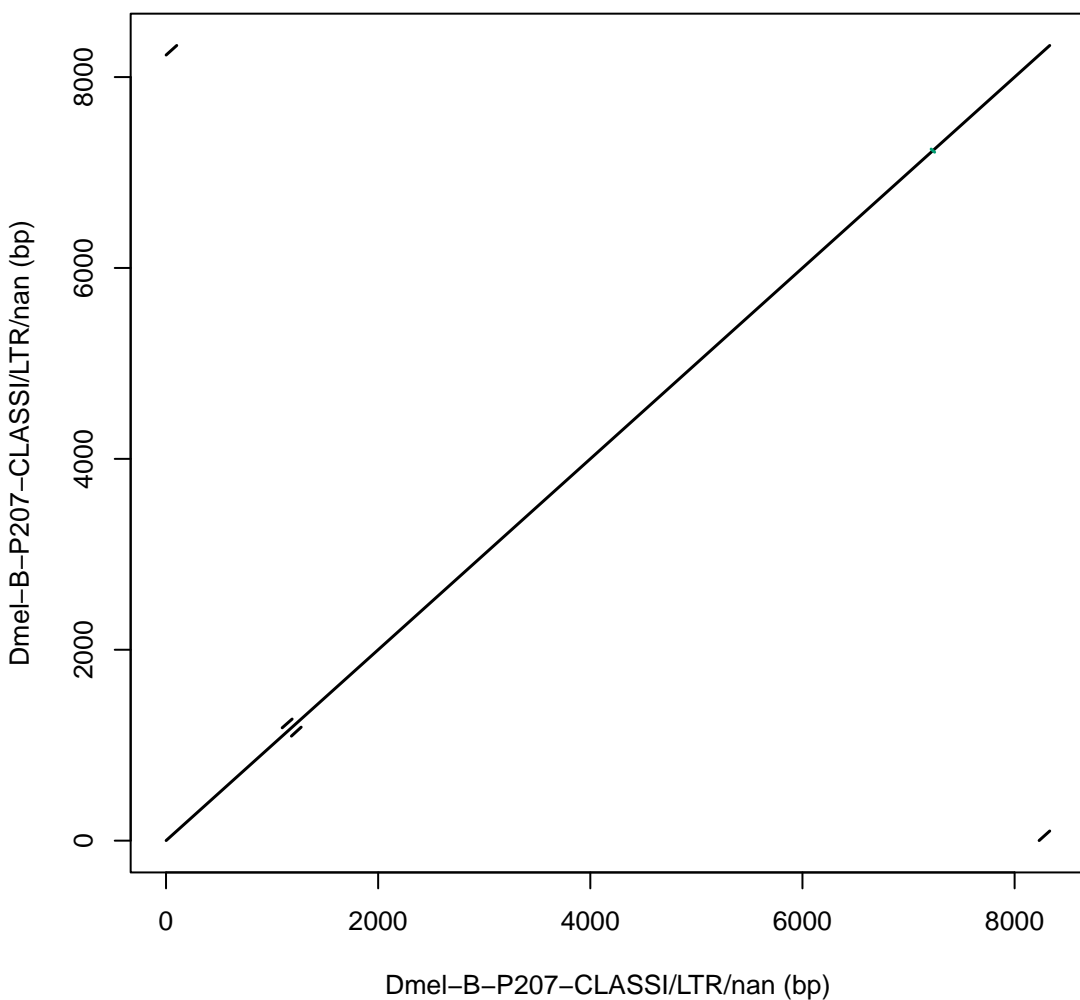
TE: Dmel-B-P207-CLASSI/LTR/nan
consensus size: 8331bp; fragments: 350; full length: 6 (≥ 7497.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

