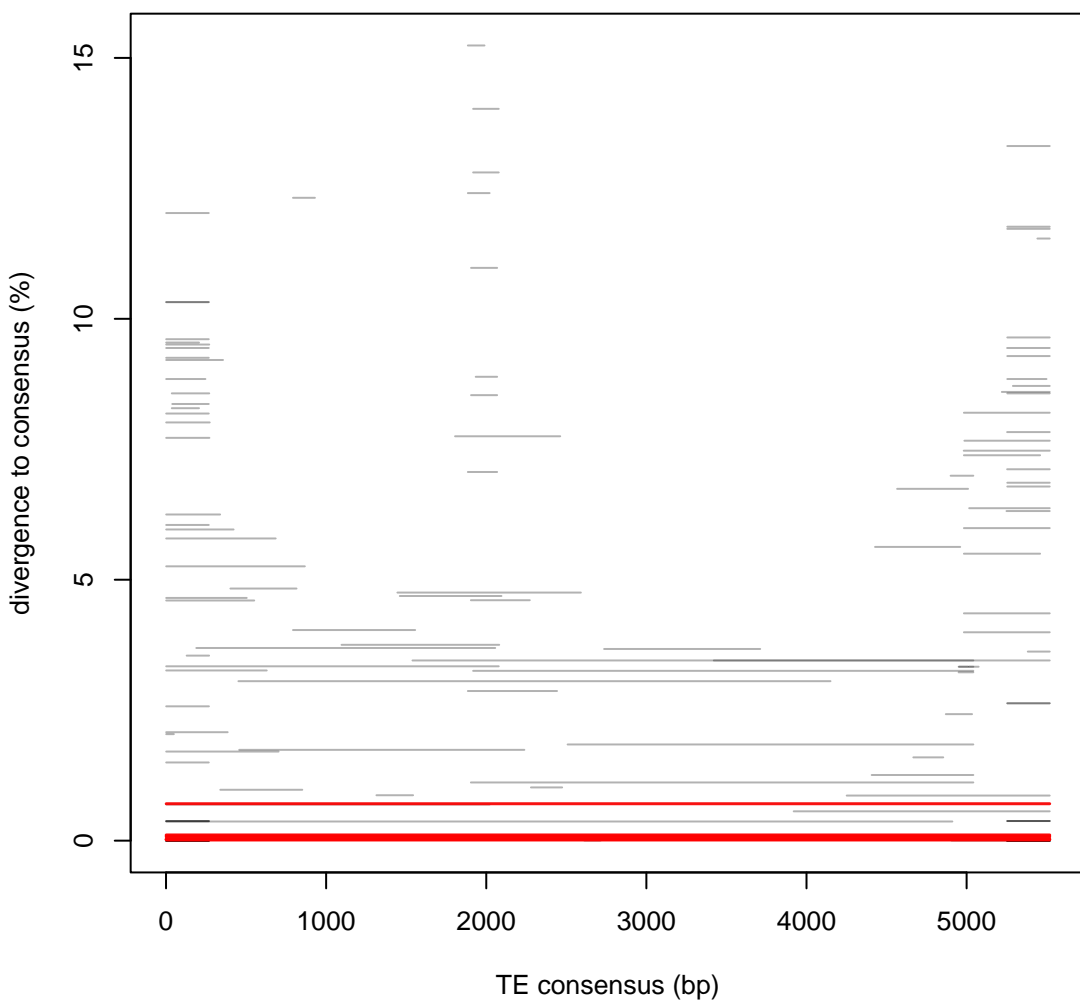
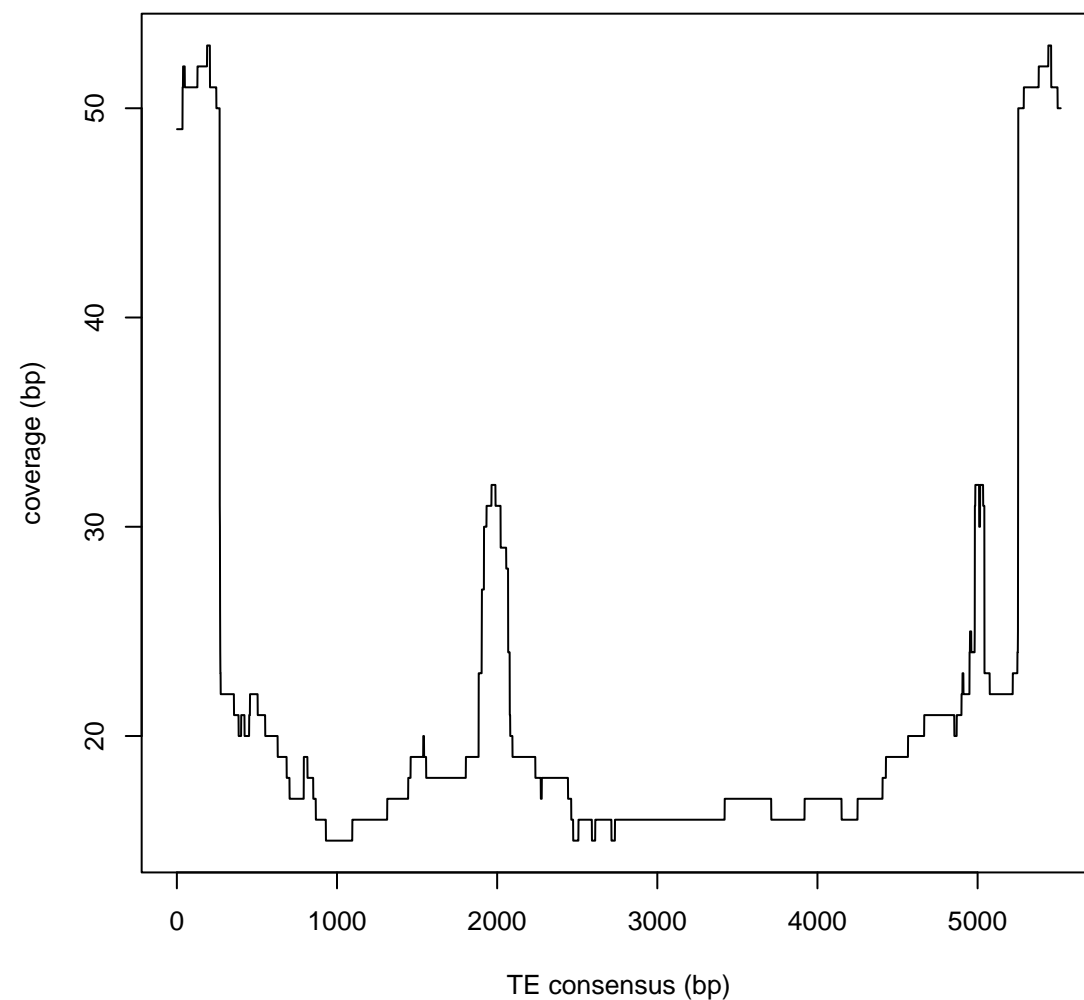


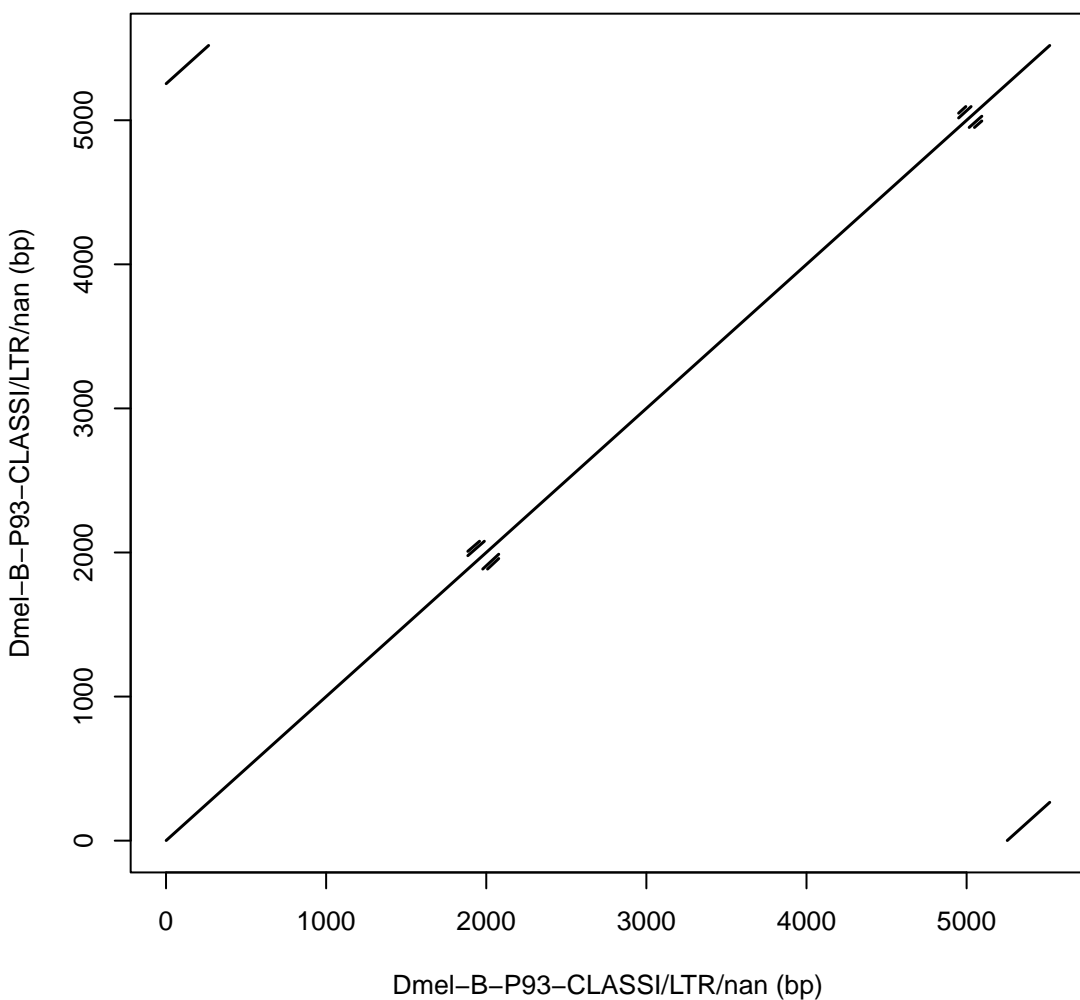
TE: Dmel-B-P93-CLASSI/LTR/nan
consensus size: 5519bp; fragments: 137; full length: 8 (>=4967.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

