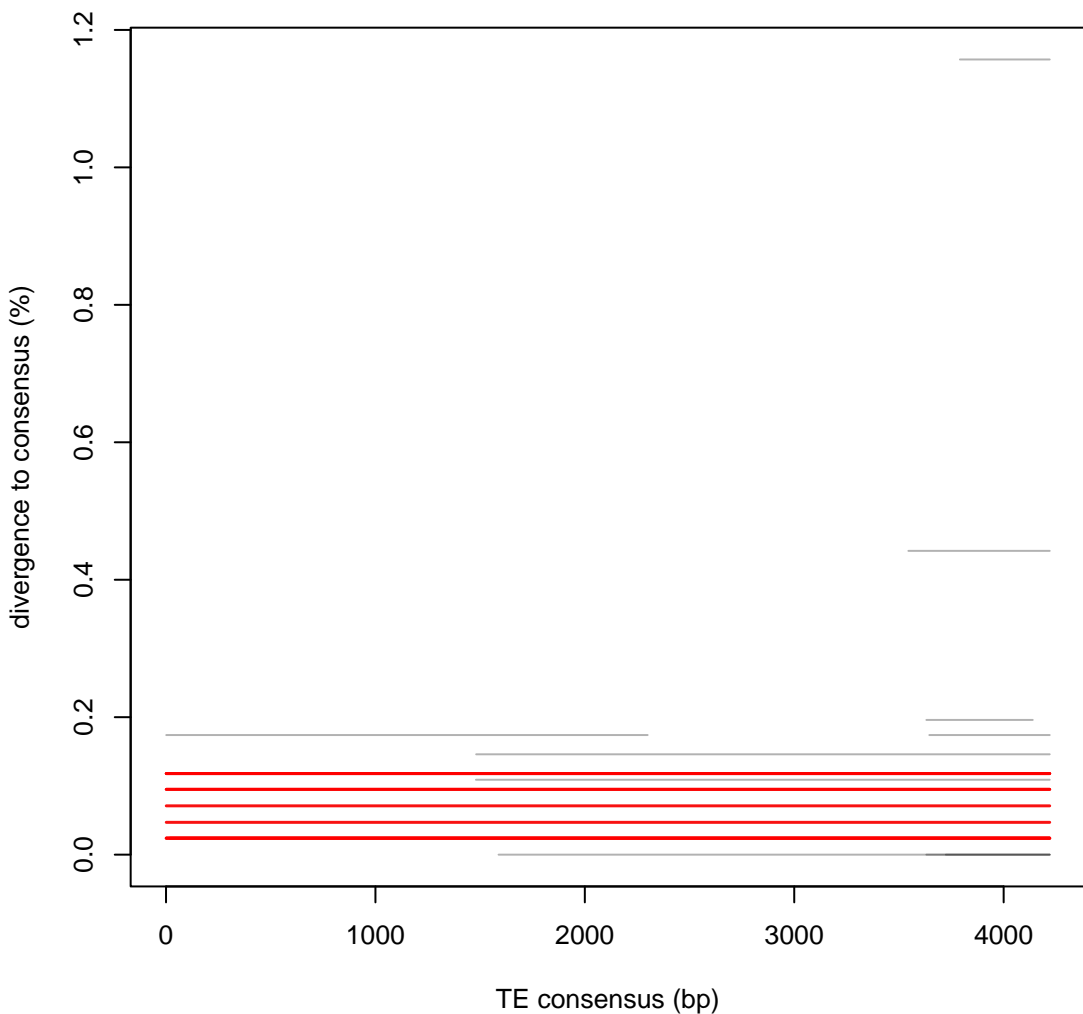
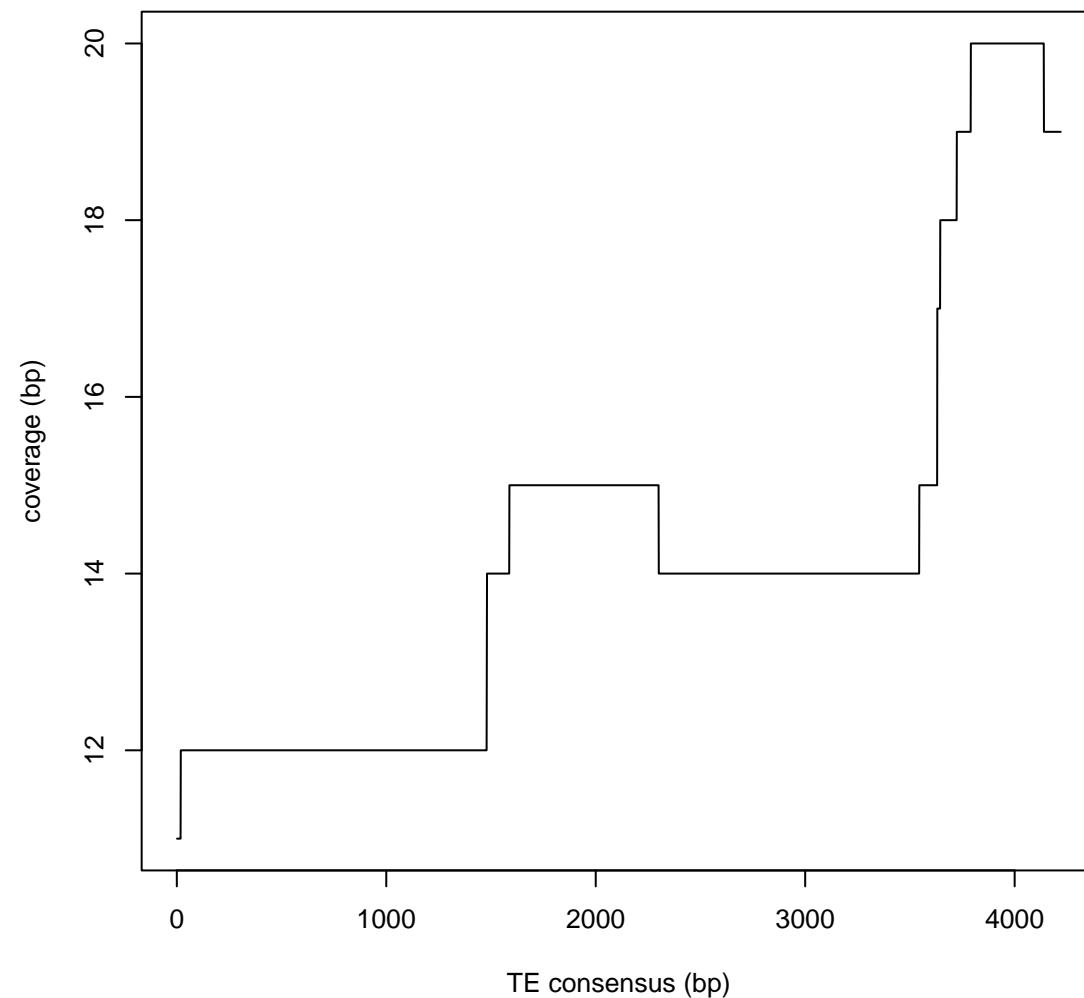


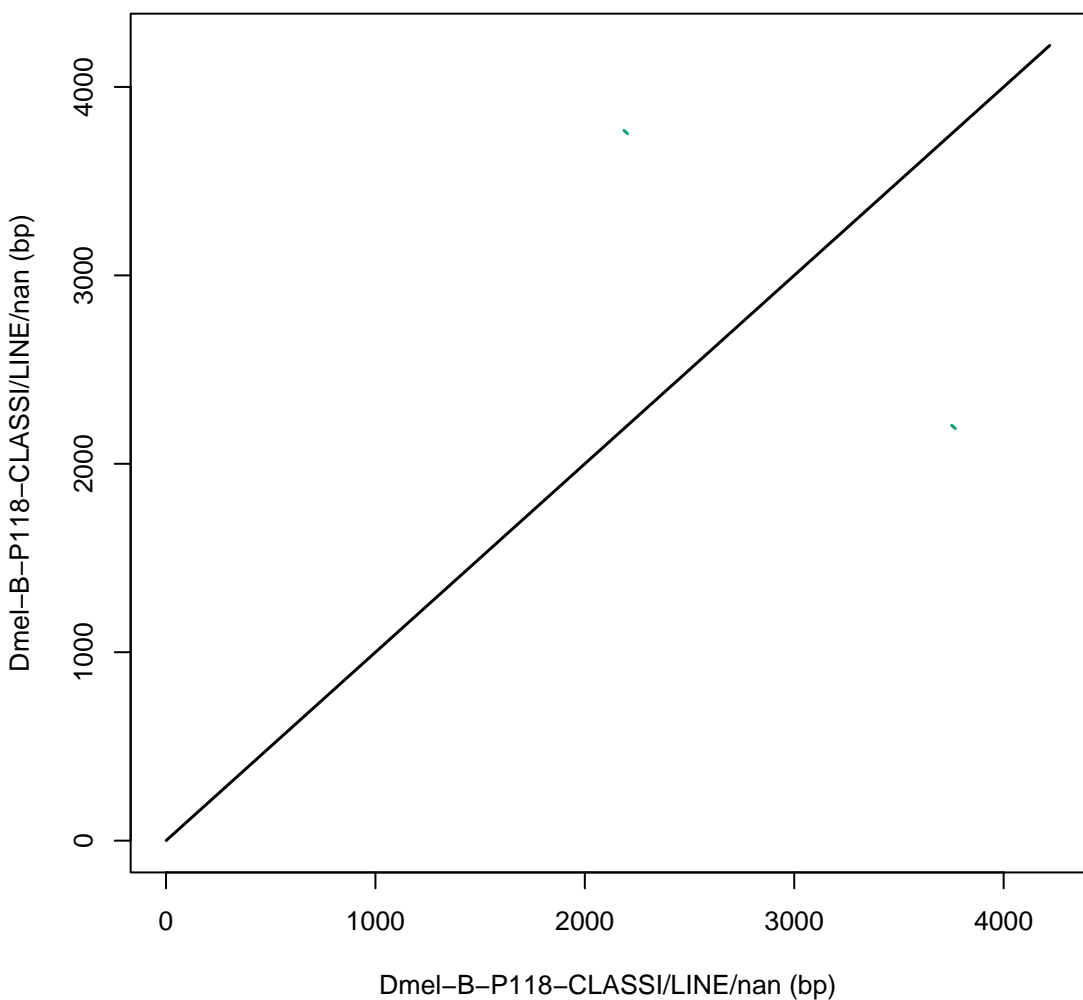
TE: Dmel-B-P118-CLASSI/LINE/nan
consensus size: 4220bp; fragments: 21; full length: 11 (≥ 3798 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

