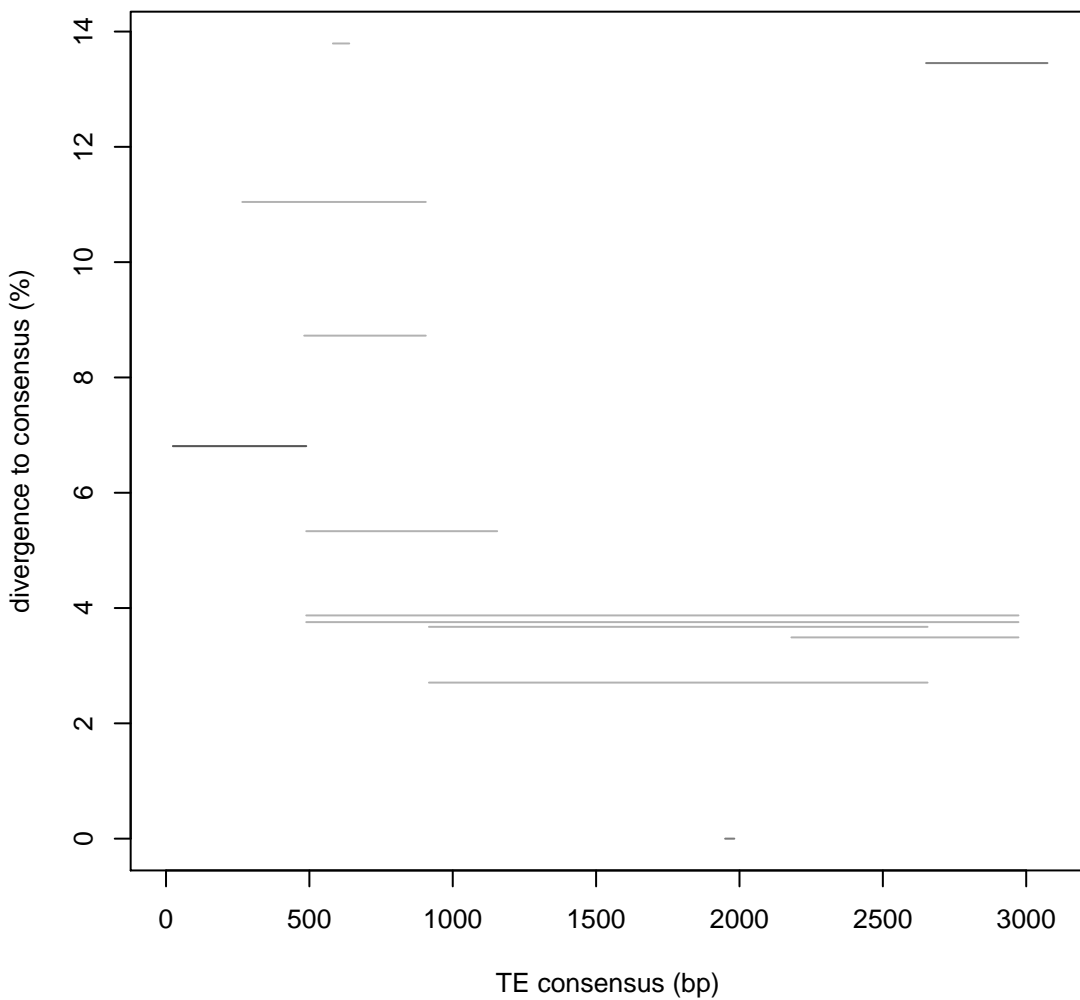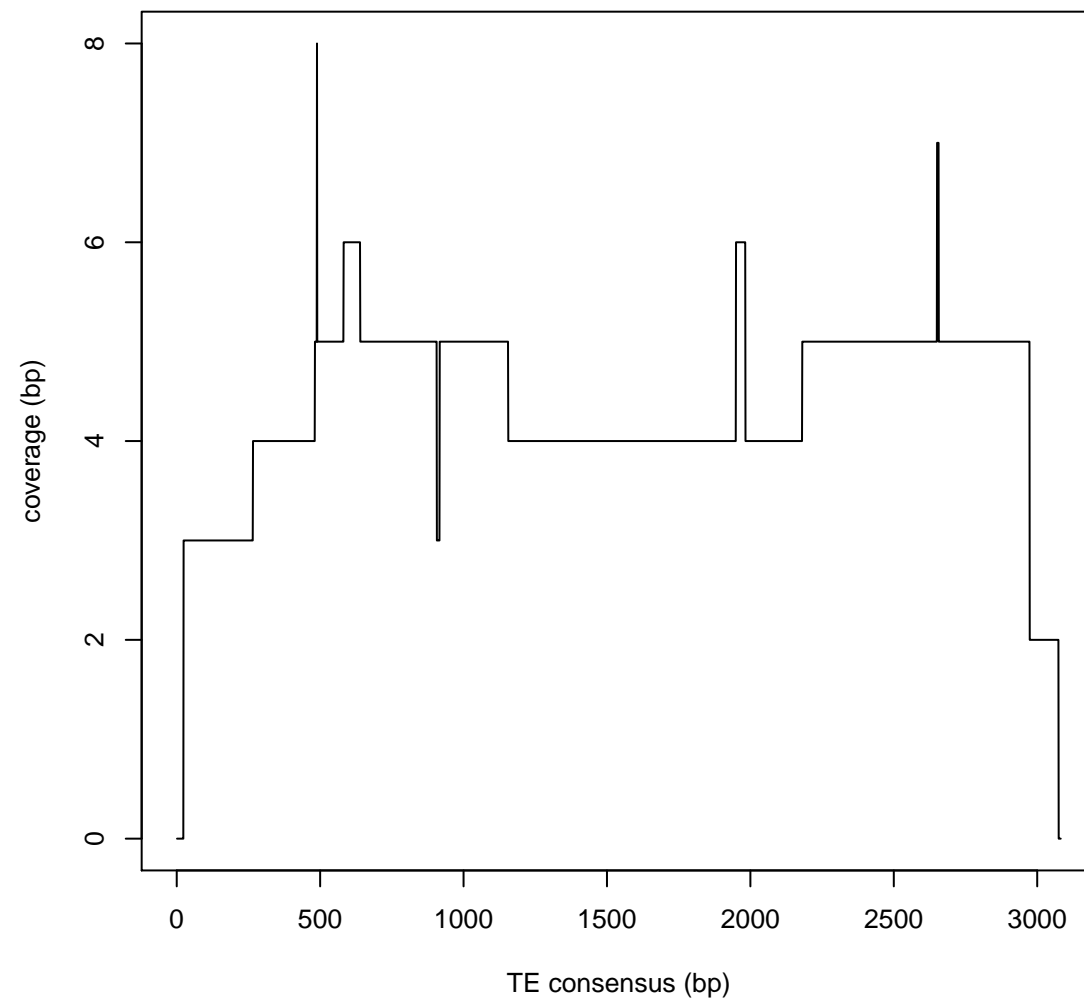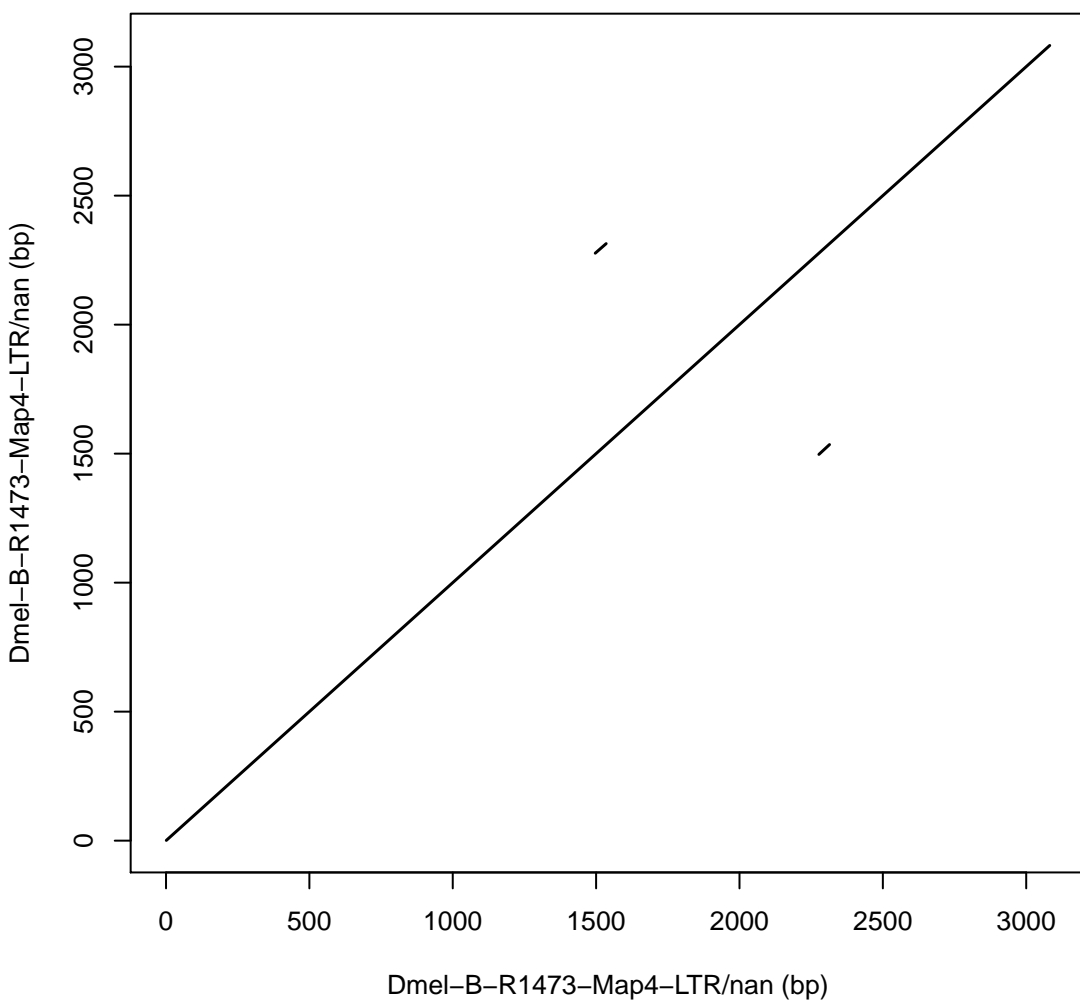**TE: Dmel−B−R1473−Map4−LTR/nan**
consensus size: 3082bp; fragments: 16; full length: 0 (>=2773.8bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected