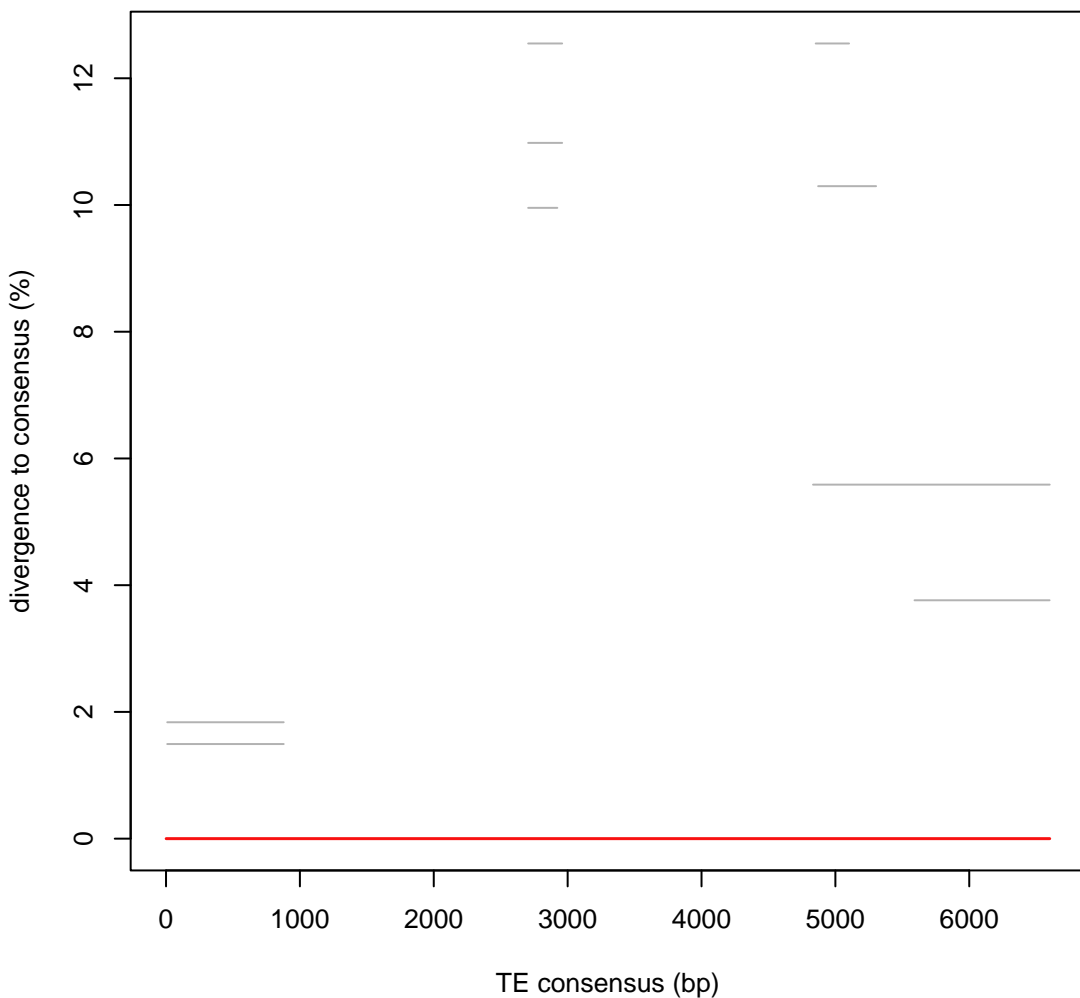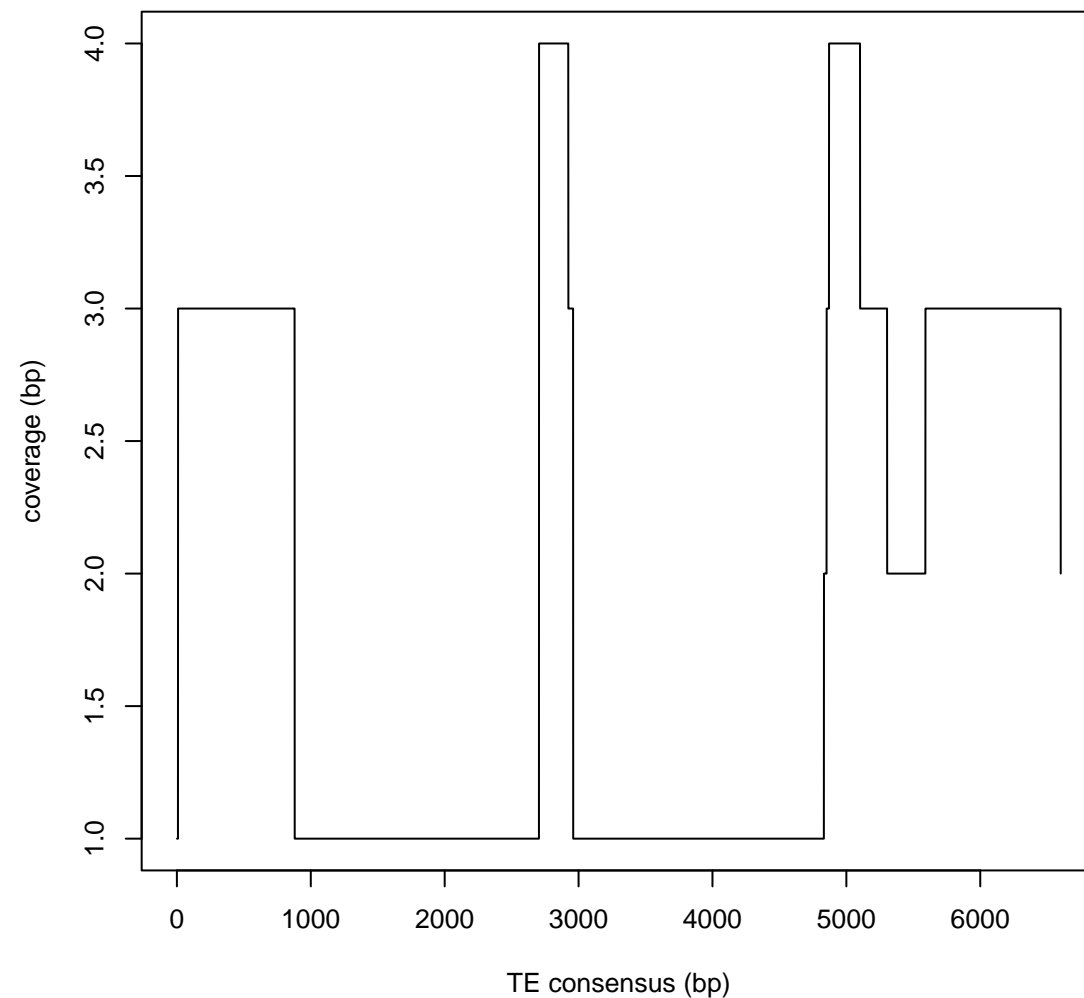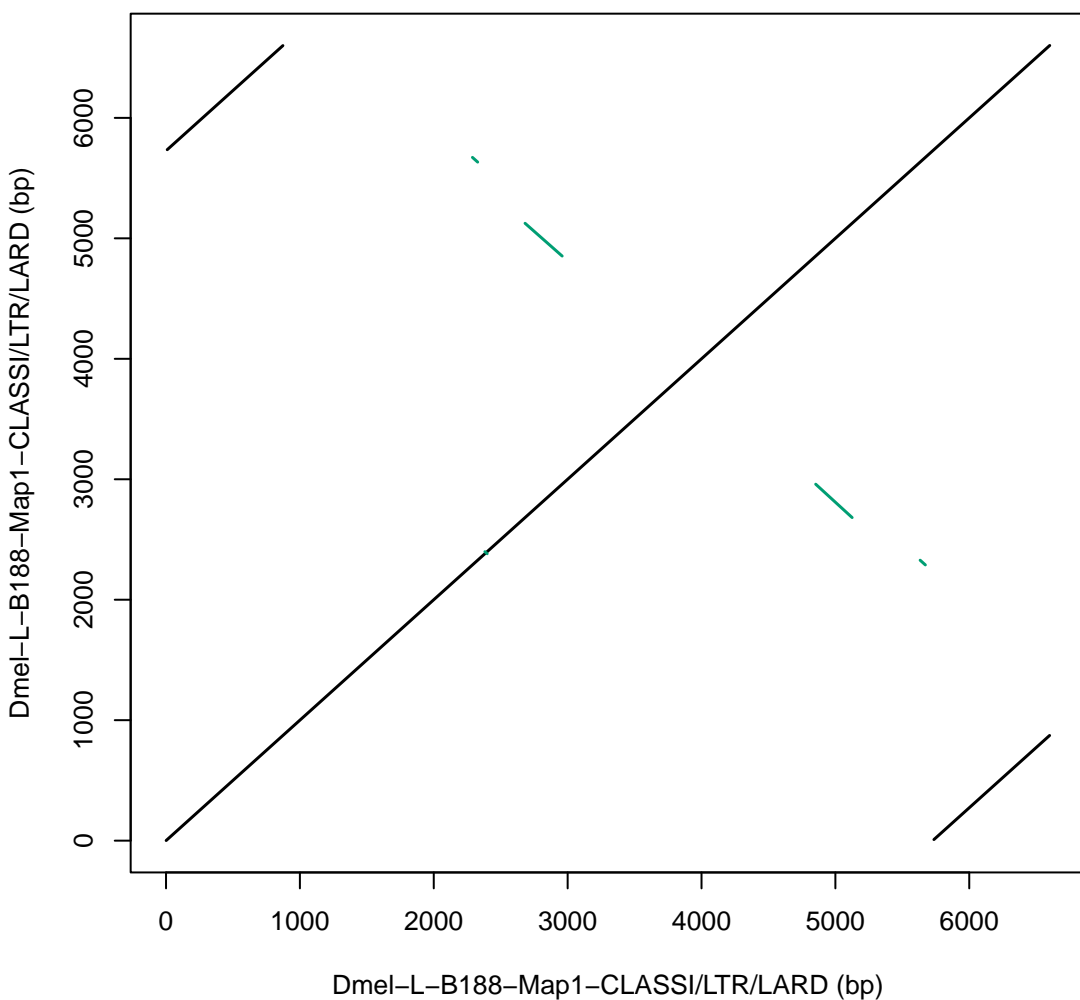**TE: Dmel−L−B188−Map1−CLASSI/LTR/LARD**
consensus size: 6601bp; fragments: 10; full length: 1 (>=5940.9bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected