TE: Dmel−B−R1844−Map4−CLASSI/LINE/nan
consensus size: 1778bp; fragments: 144; full length: 1 (>=1600.2bp)

divergence to consensus (%)
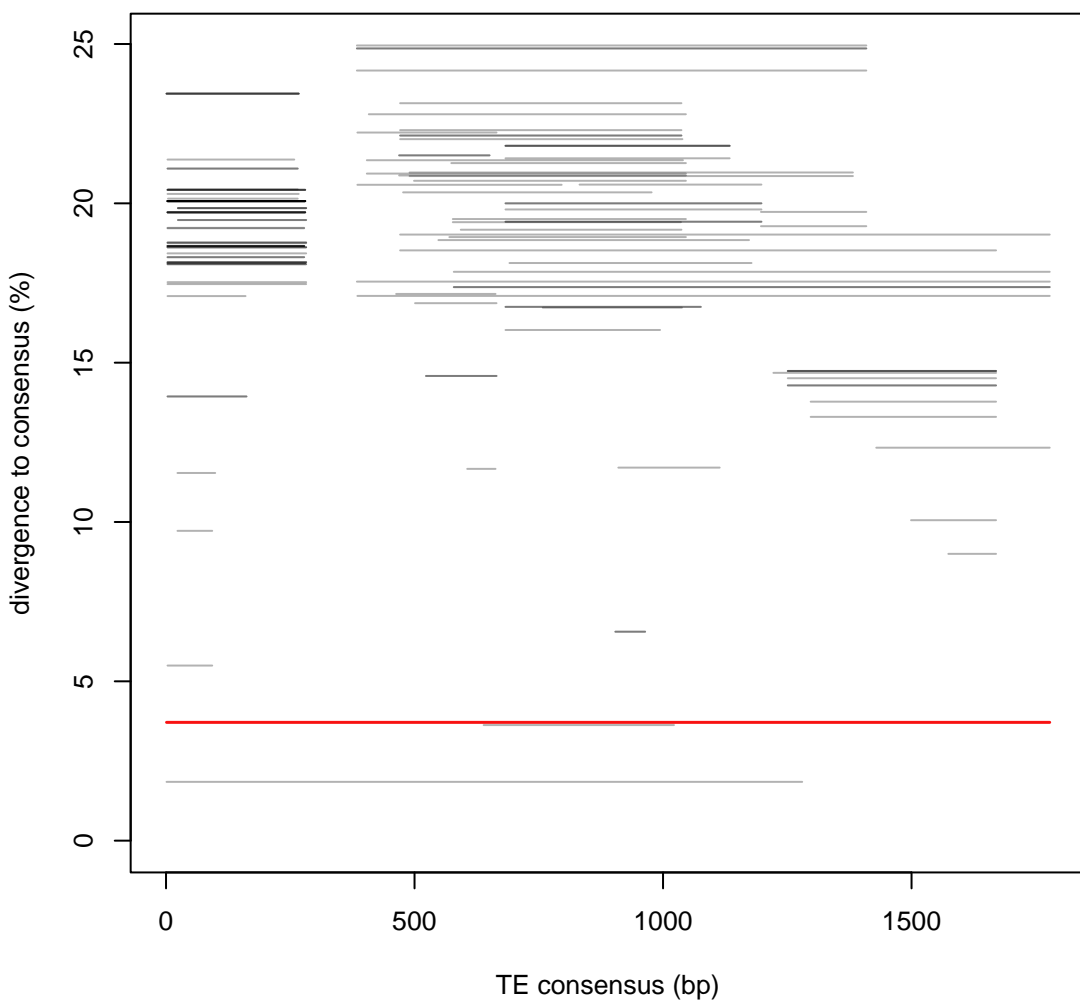
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
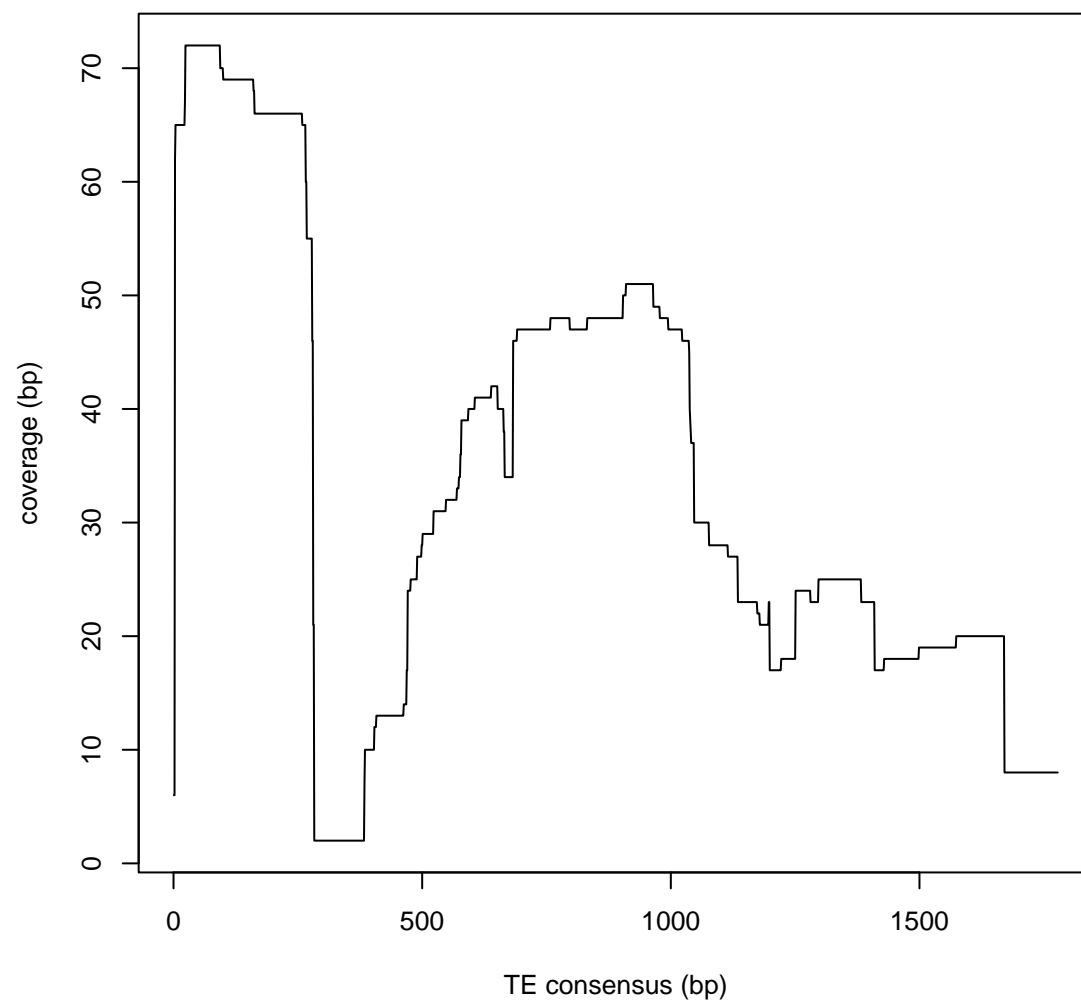
TE consensus (bp)

TE consensus self dotplot (blastn)

Dmel−B−R1844−Map4−CLASSI/LINE/nan (bp)

Dmel−B−R1844−Map4−CLASSI/LINE/nan (bp)
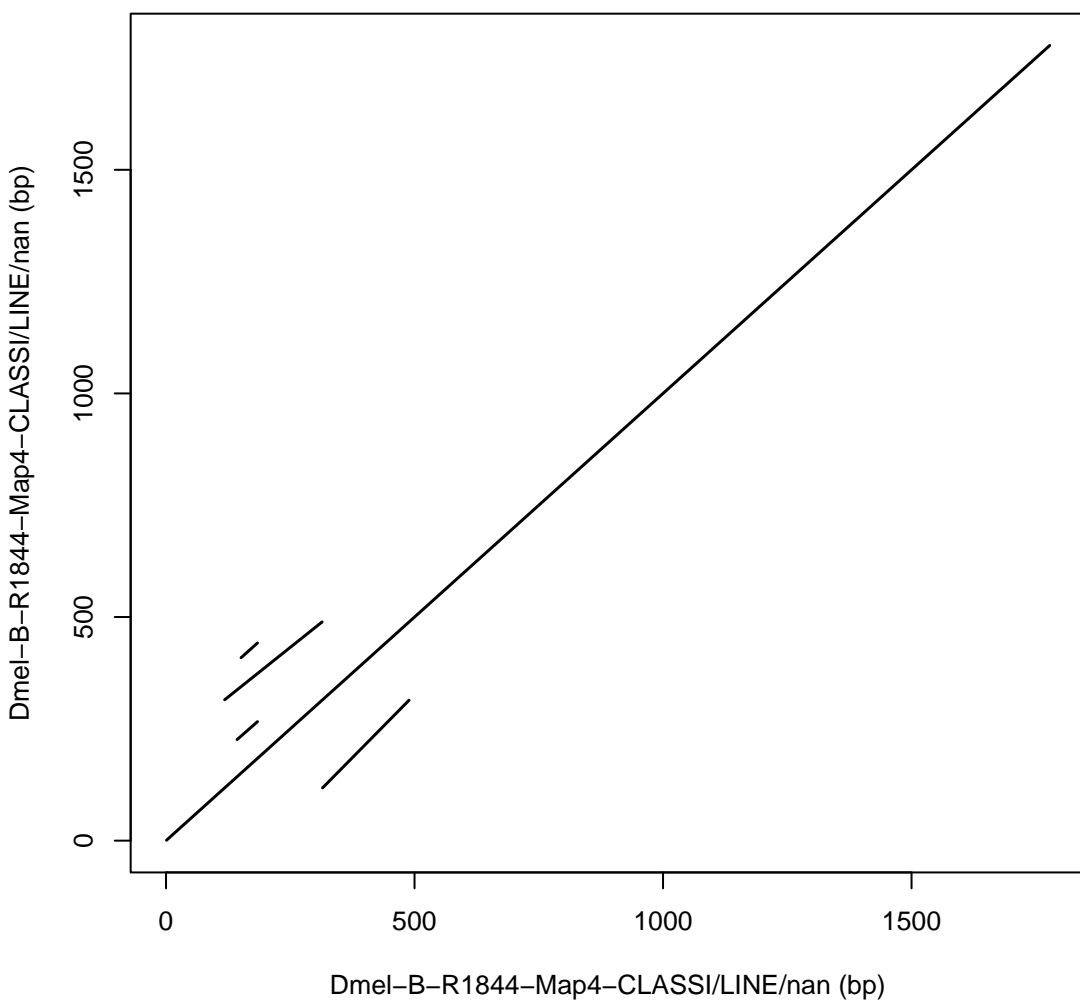
TE consensus structure and protein hits

no orf >400 bp detected

Dmel−B−R1844−Map4−CLASSI/LINE/nan (bp)