TE: Dmel−B−P63.94−Map20−TIR/nan
consensus size: 2203bp; fragments: 886; full length: 0 (>=1982.7bp)

divergence to consensus (%)
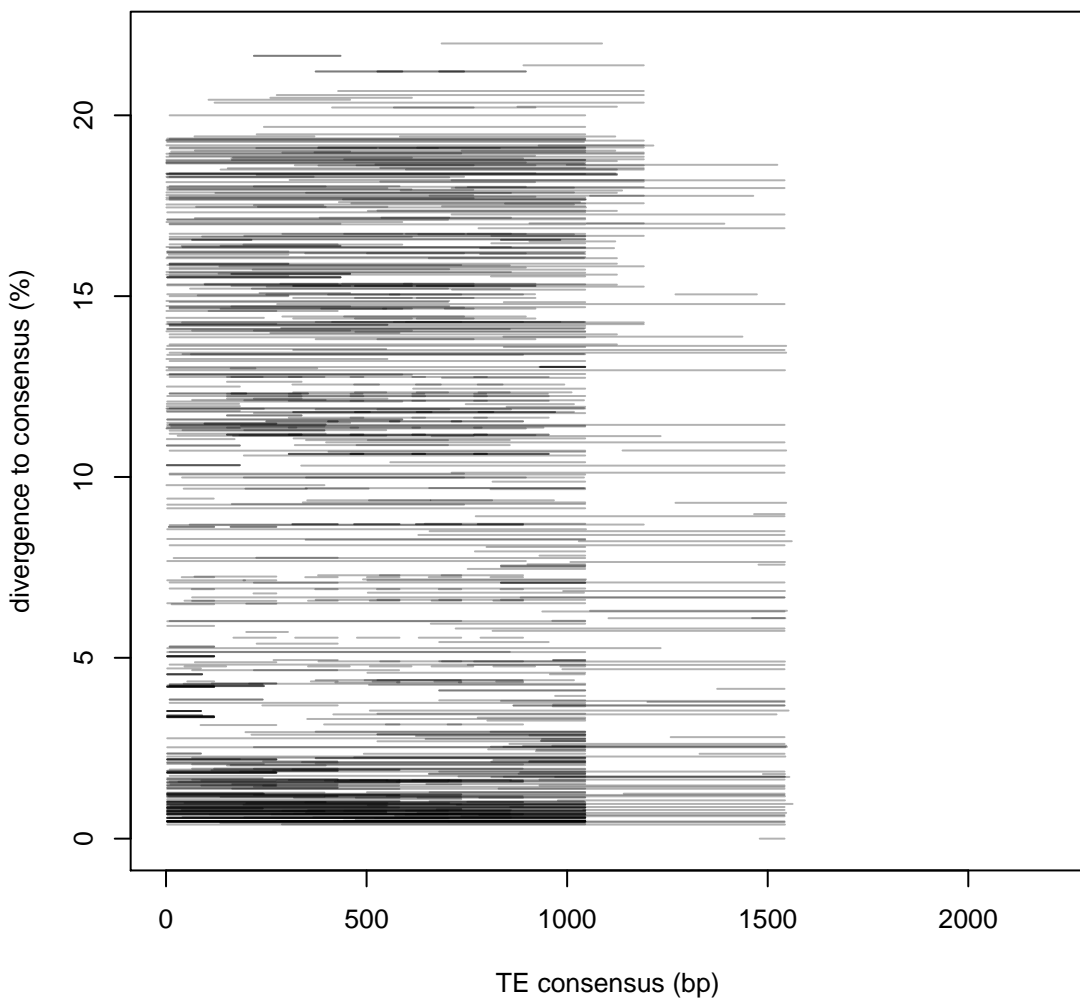
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
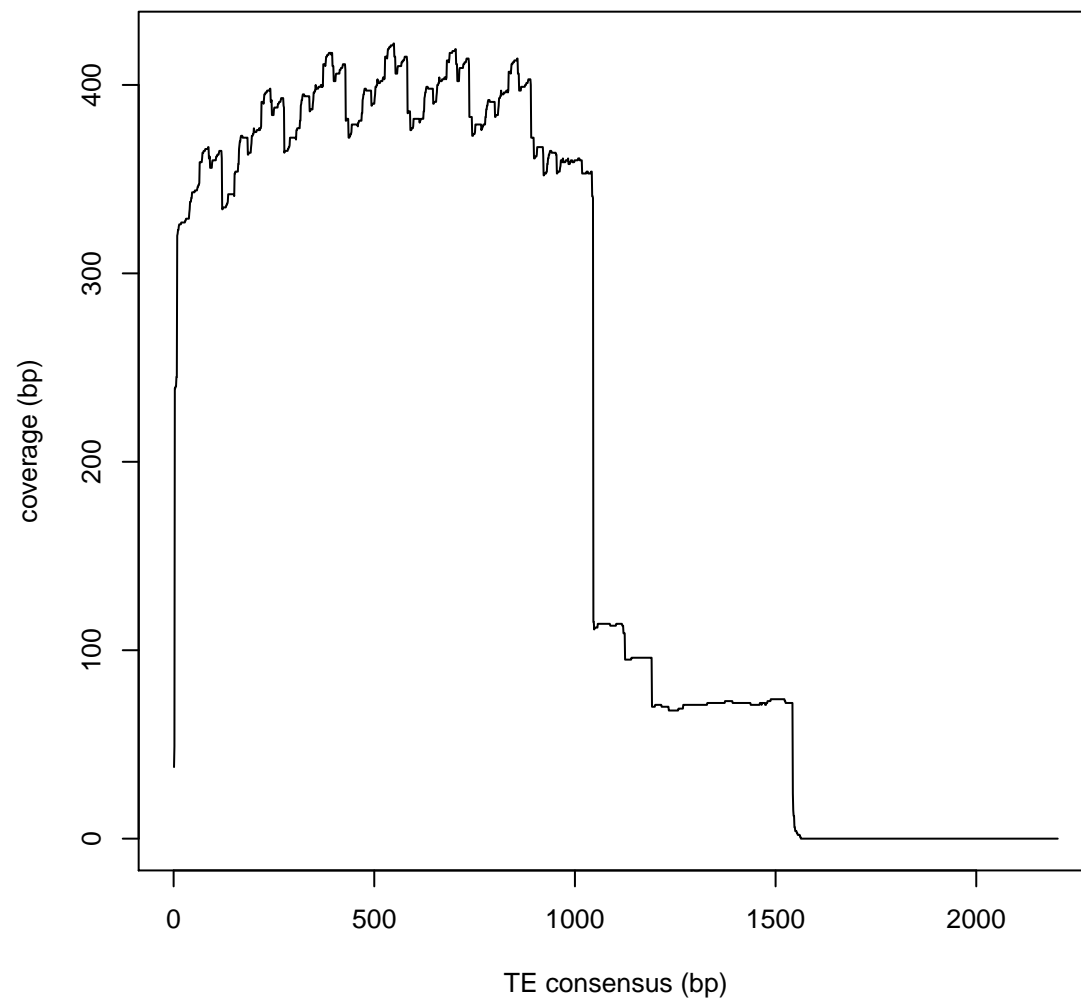
TE consensus (bp)

TE consensus self dotplot (blastn)

Dmel−B−P63.94−Map20−TIR/nan (bp)

Dmel−B−P63.94−Map20−TIR/nan (bp)
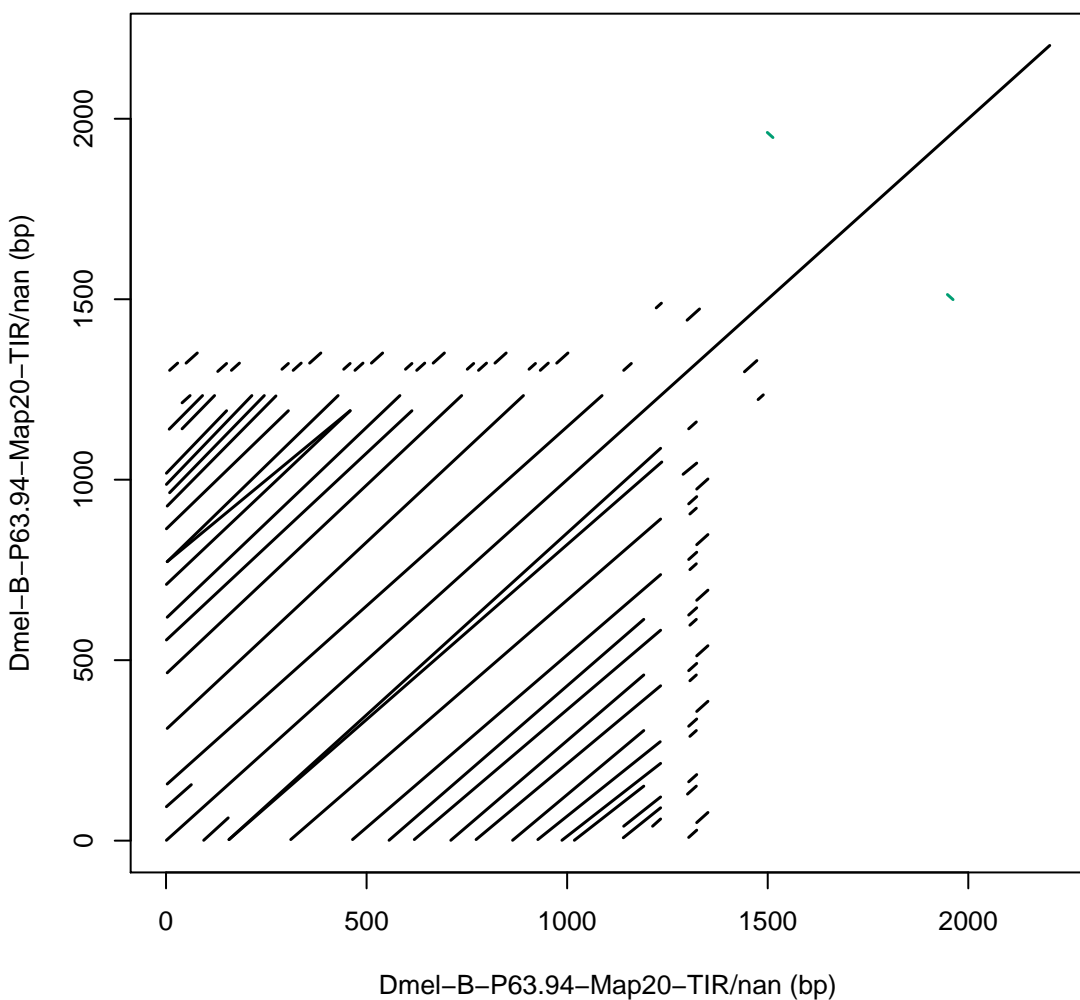
TE consensus structure and protein hits

no orf >400 bp detected

Dmel−B−P63.94−Map20−TIR/nan (bp)