TE: seq_7741
consensus size: 3630bp; fragments: 14238; full length: 556 (>=3267bp)

divergence to consensus (%)
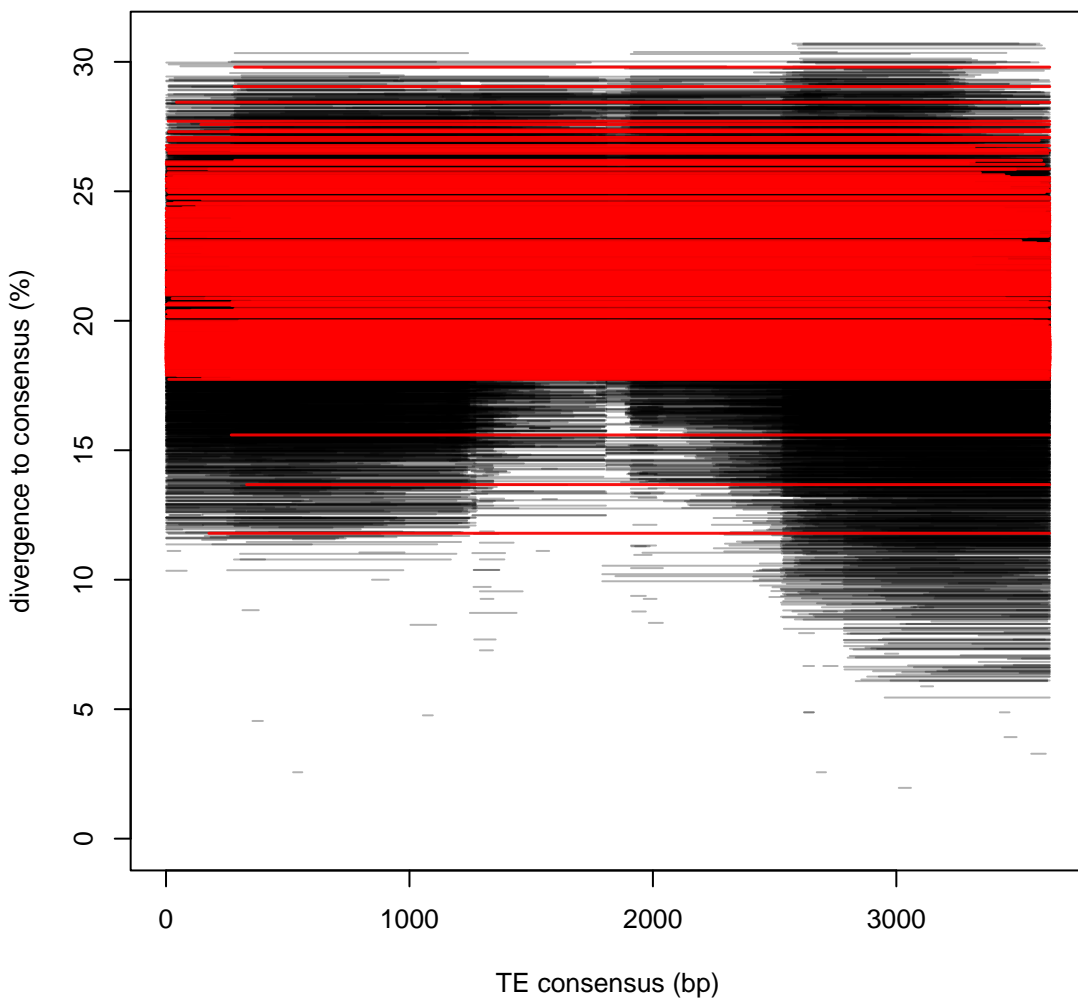
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
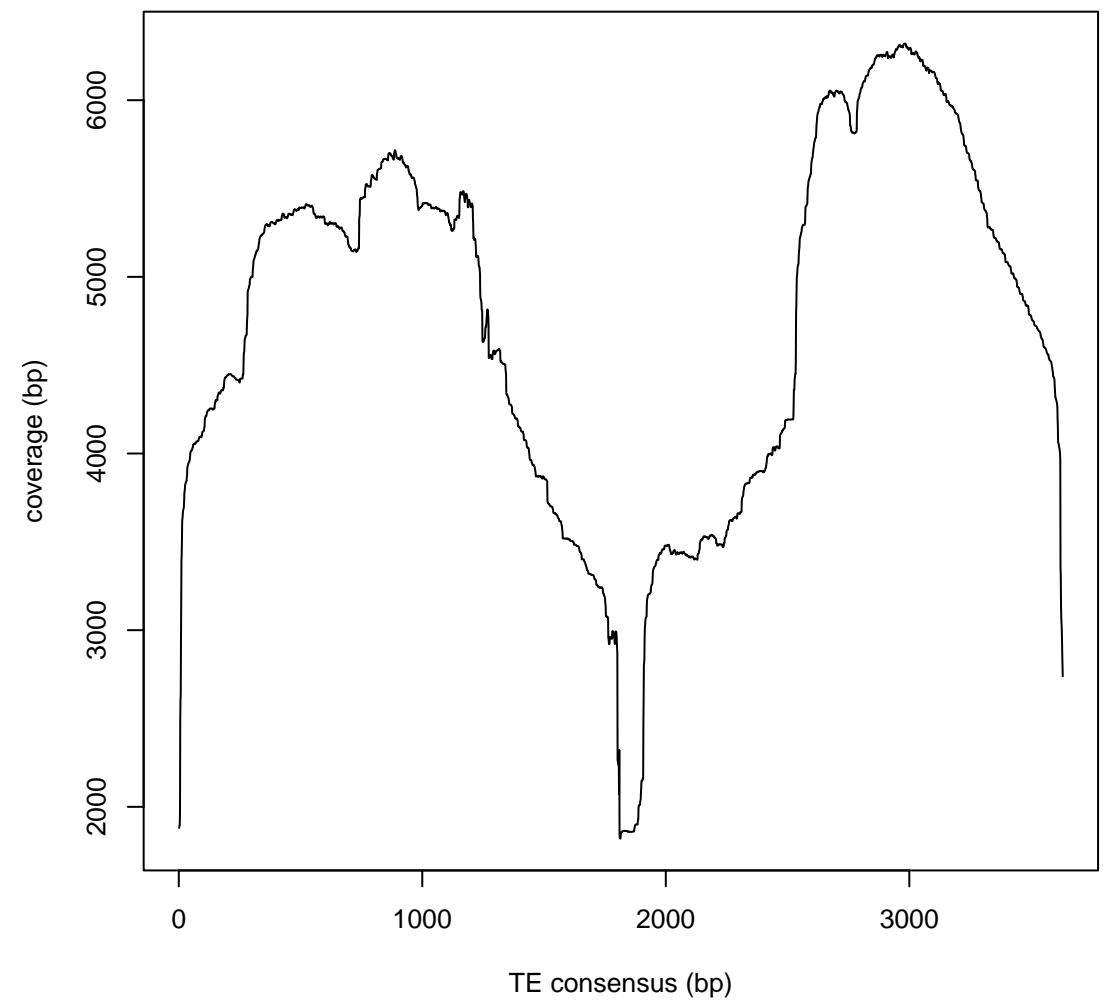
TE consensus (bp)

TE consensus self dotplot (blastn)

seq_7741 (bp)

seq_7741 (bp)
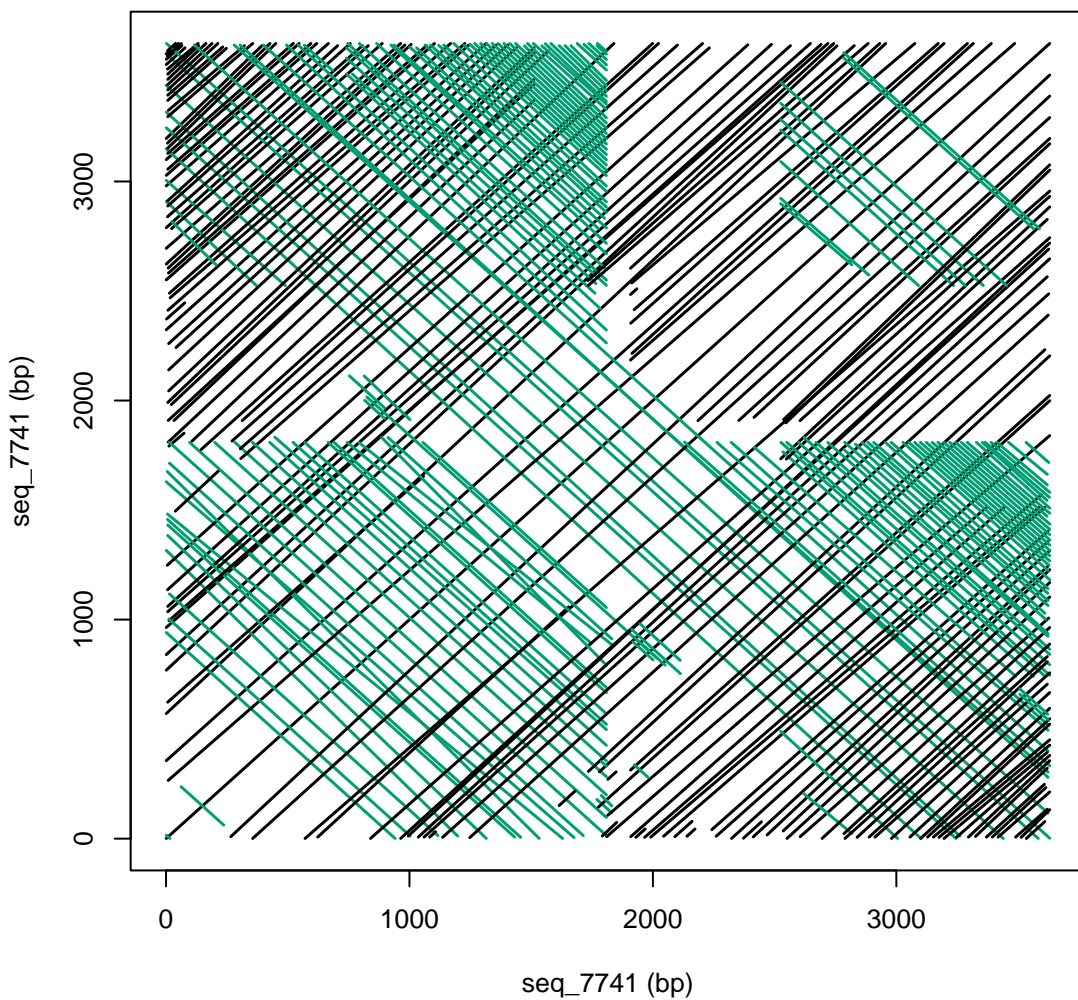
TE consensus structure and protein hits

no orf >400 bp detected

seq_7741 (bp)