TE: Dmel–B–G519–Map5–Unclassified/nan
consensus size: 2748bp; fragments: 16; full length: 0 (>=2473.2bp)

divergence to consensus (%)
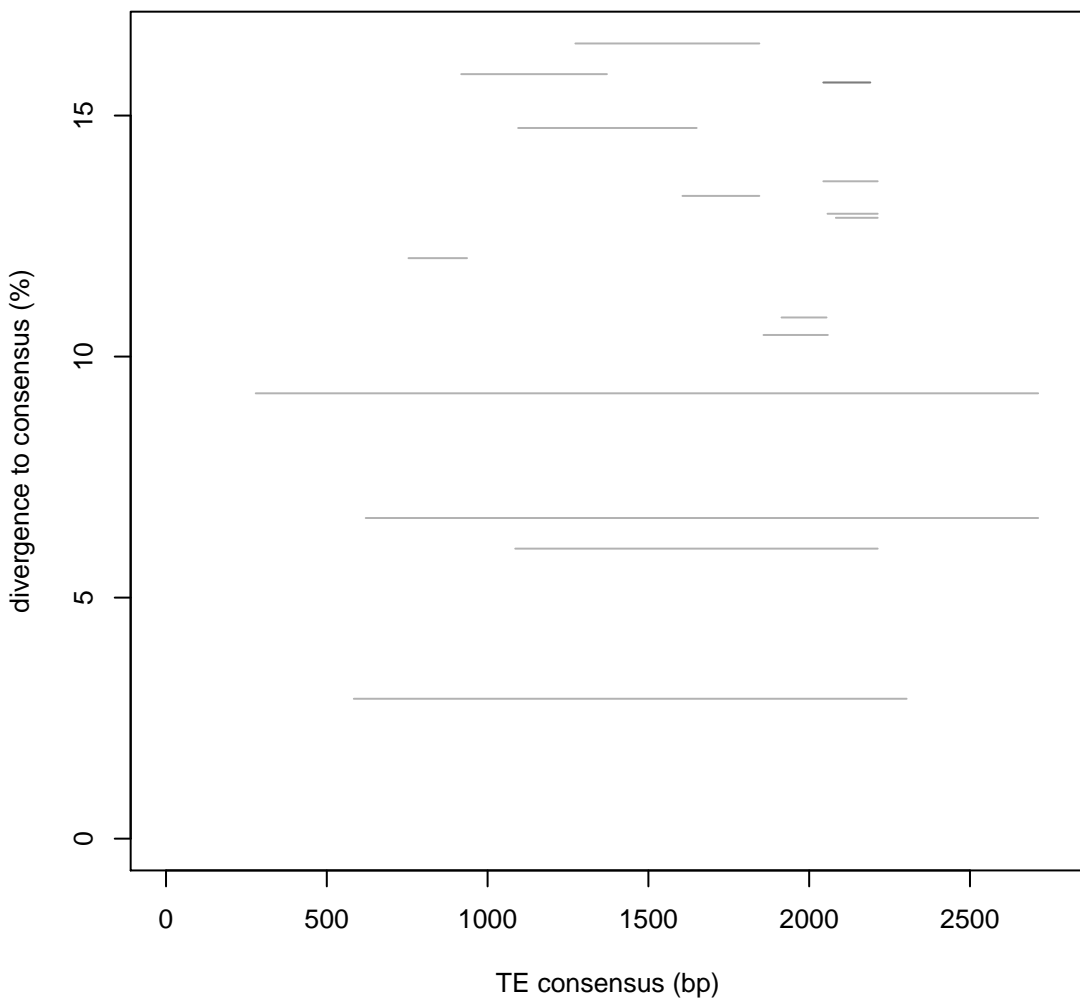
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
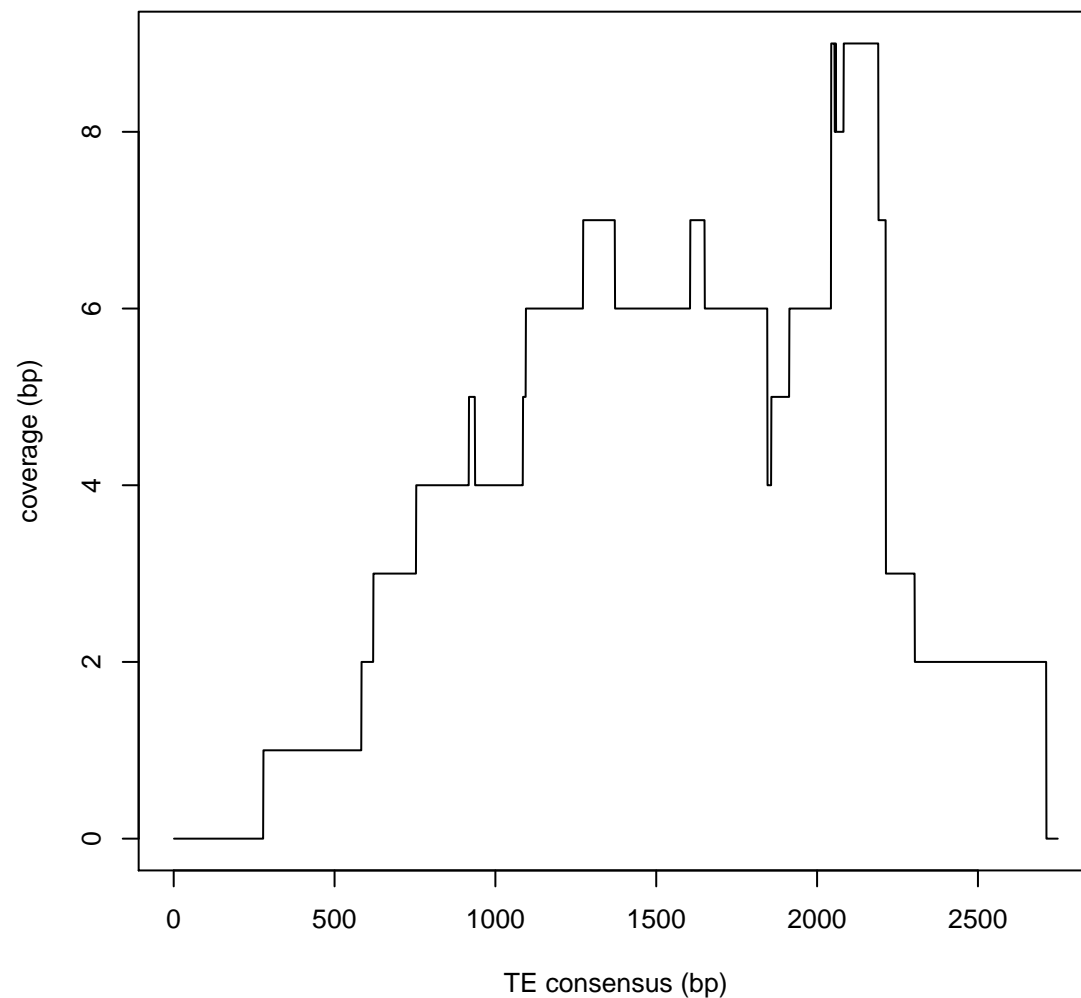
TE consensus (bp)

TE consensus self dotplot (blastn)

Dmel–B–G519–Map5–Unclassified/nan (bp)

Dmel–B–G519–Map5–Unclassified/nan (bp)
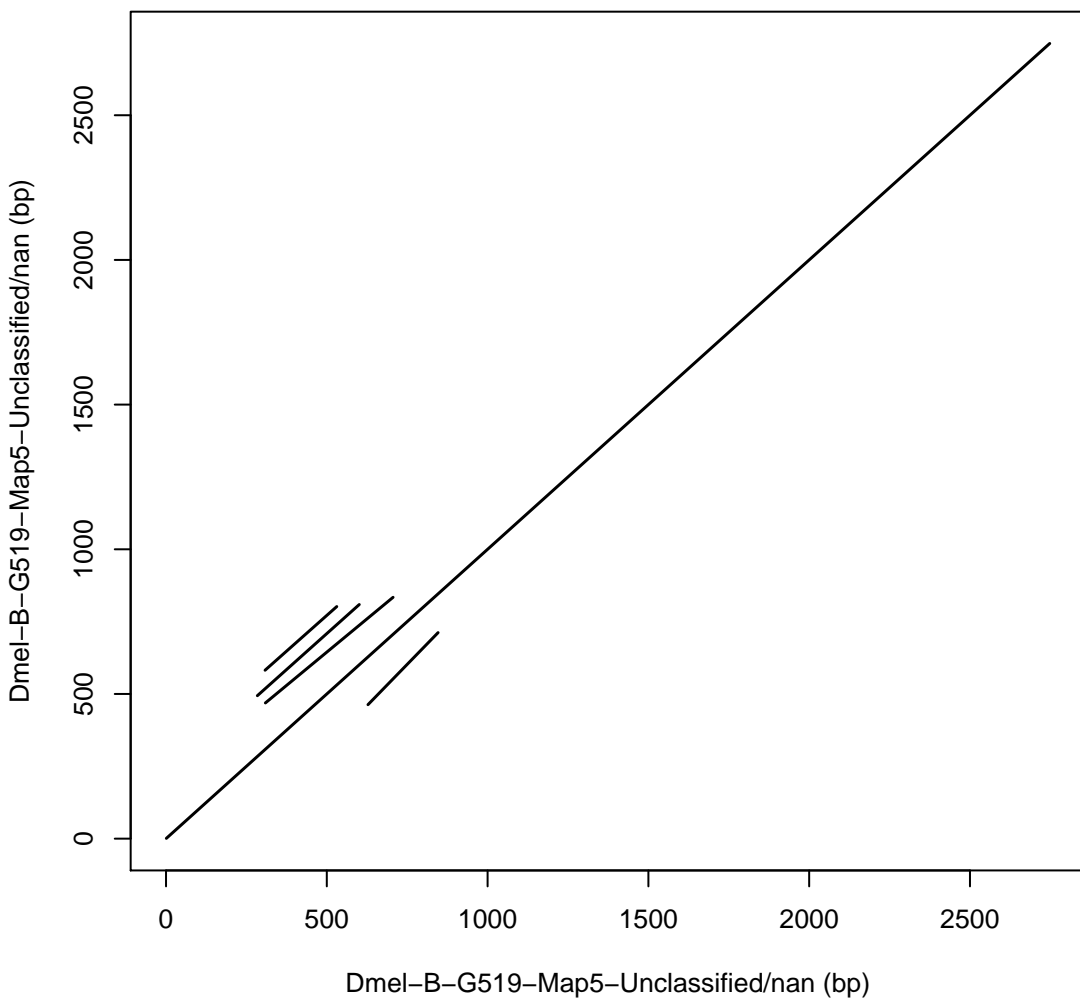
TE consensus structure and protein hits

DOC_pol LINE/I–Jockey

Dmel–B–G519–Map5–Unclassified/nan (bp)