TE: DmeI−B−R312−Map4−LINE/nan
consensus size: 3194bp; fragments: 202; full length: 3 (>=2874.6bp)

divergence to consensus (%)
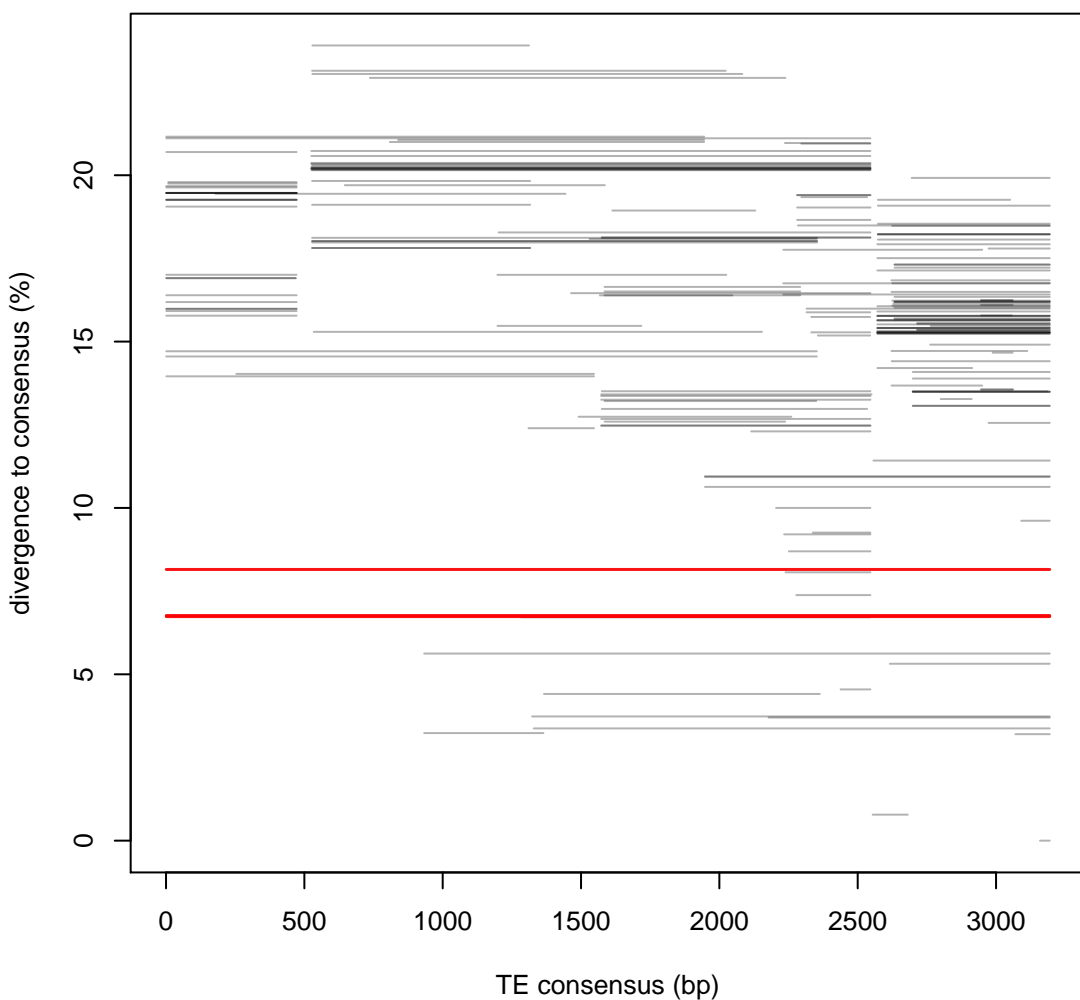
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
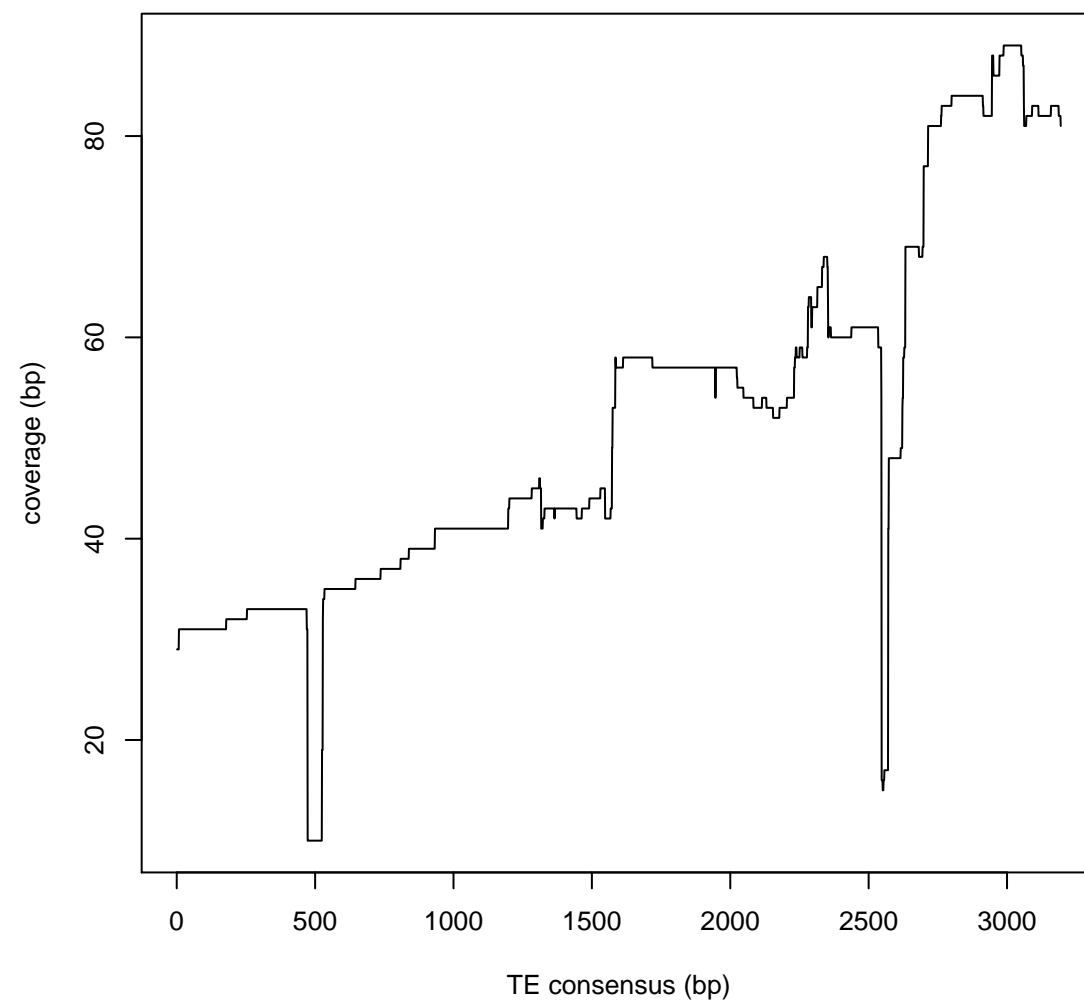
TE consensus (bp)

TE consensus self dotplot (blastn)

Dmel−B−R312−Map4−LINE/nan (bp)

Dmel−B−R312−Map4−LINE/nan (bp)
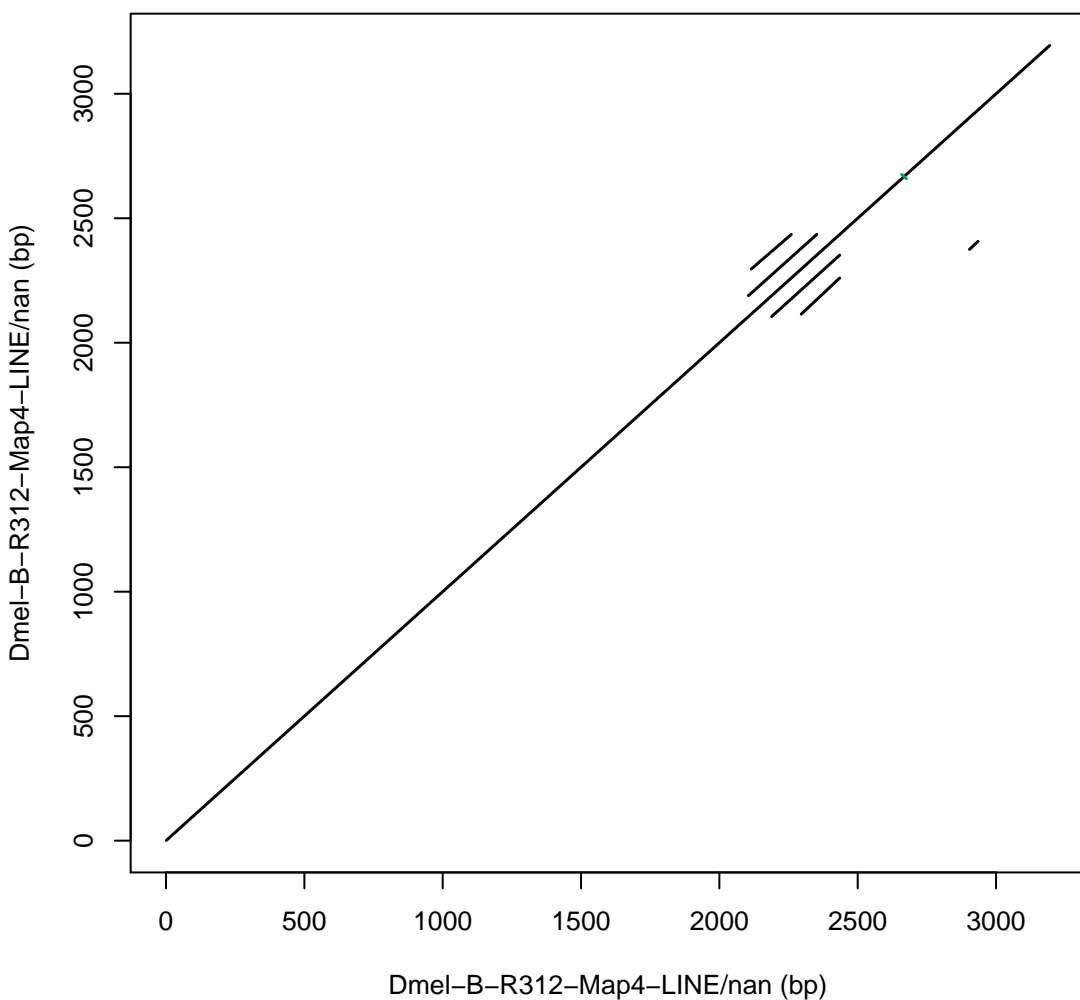
TE consensus structure and protein hits

HETA_gag LINE/I−Jockey

Gypsy−8_PSH_pol LTR/Gypsy

Dmel−B−R312−Map4−LINE/nan (bp)