TE: Dmel−L−B38−Map2_reversed−LINE
consensus size: 2859bp; fragments: 448; full length: 0 (>=2573.1bp)

divergence to consensus (%)
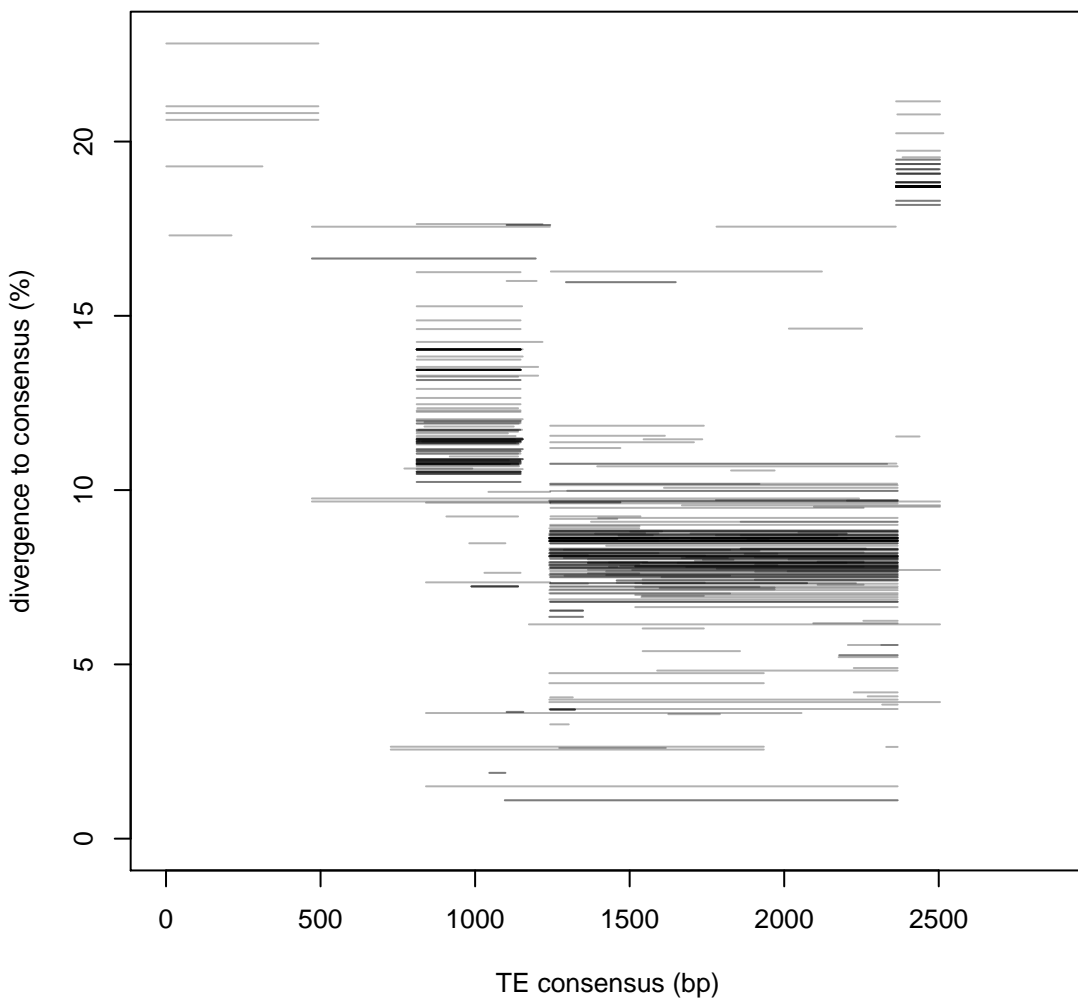
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
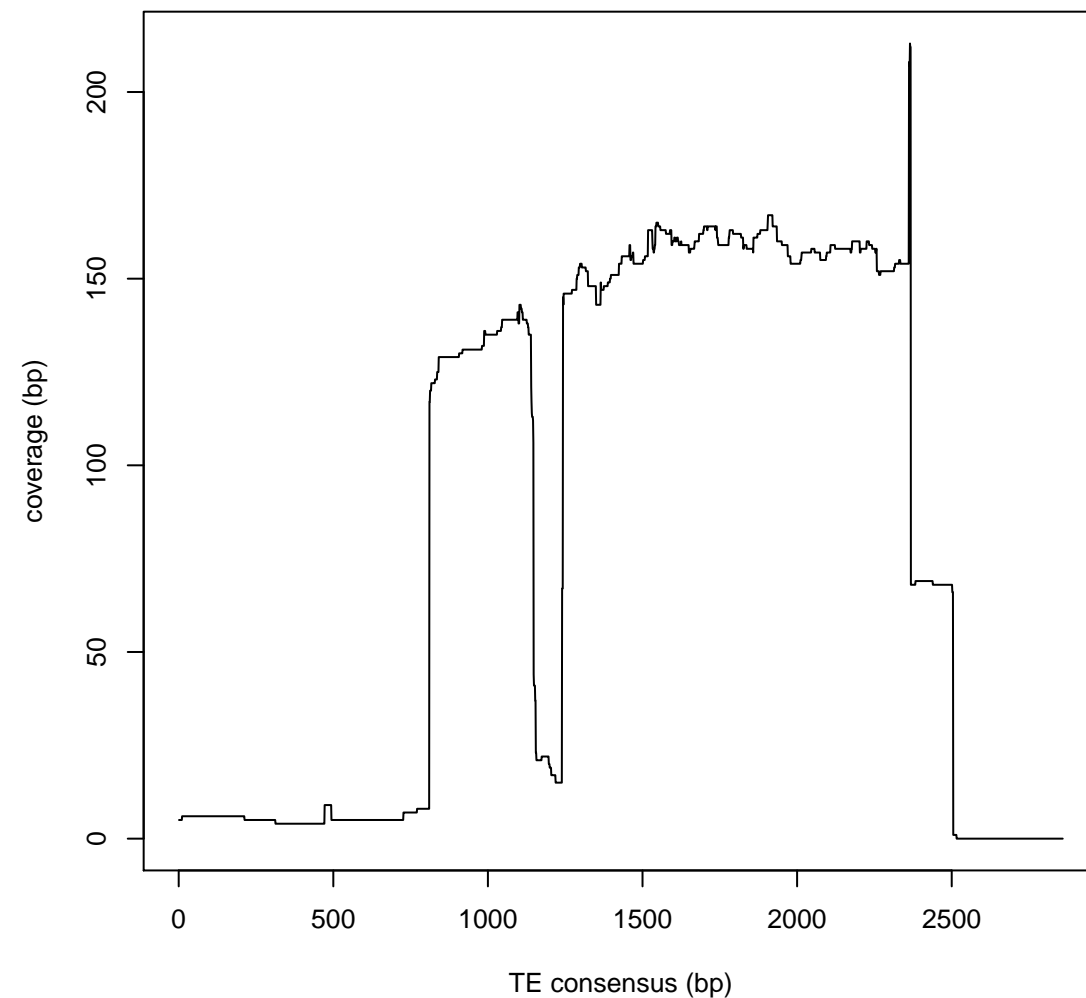
TE consensus (bp)

TE consensus self dotplot (blastn)

Dmel−L−B38−Map2_reversed−LINE (bp)

Dmel−L−B38−Map2_reversed−LINE (bp)
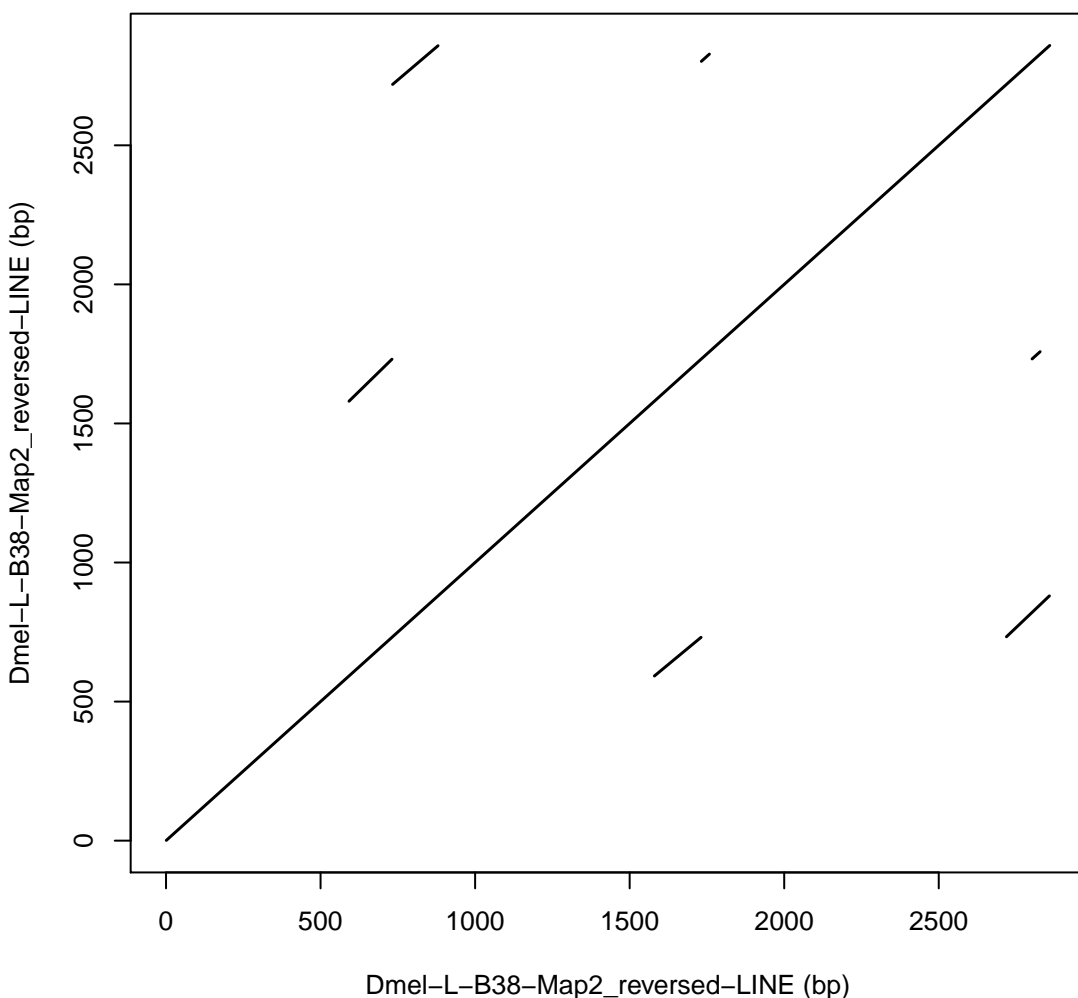
TE consensus structure and protein hits

BEL_pol LTR/Pao
BELPADRE_pol LTR/Pao
BEL−1_DGri−I_pol LTR/Pao
BEL−8_DMo_p2 LTR/Pao
BEL−6_DWil_pol LTR/P

Dmel−L−B38−Map2_reversed−LINE (bp)