TE: Dmel−L−B123−Map1−LTR/Gypsy
consensus size: 7911bp; fragments: 406; full length: 1 (>=7119.9bp)

divergence to consensus (%)
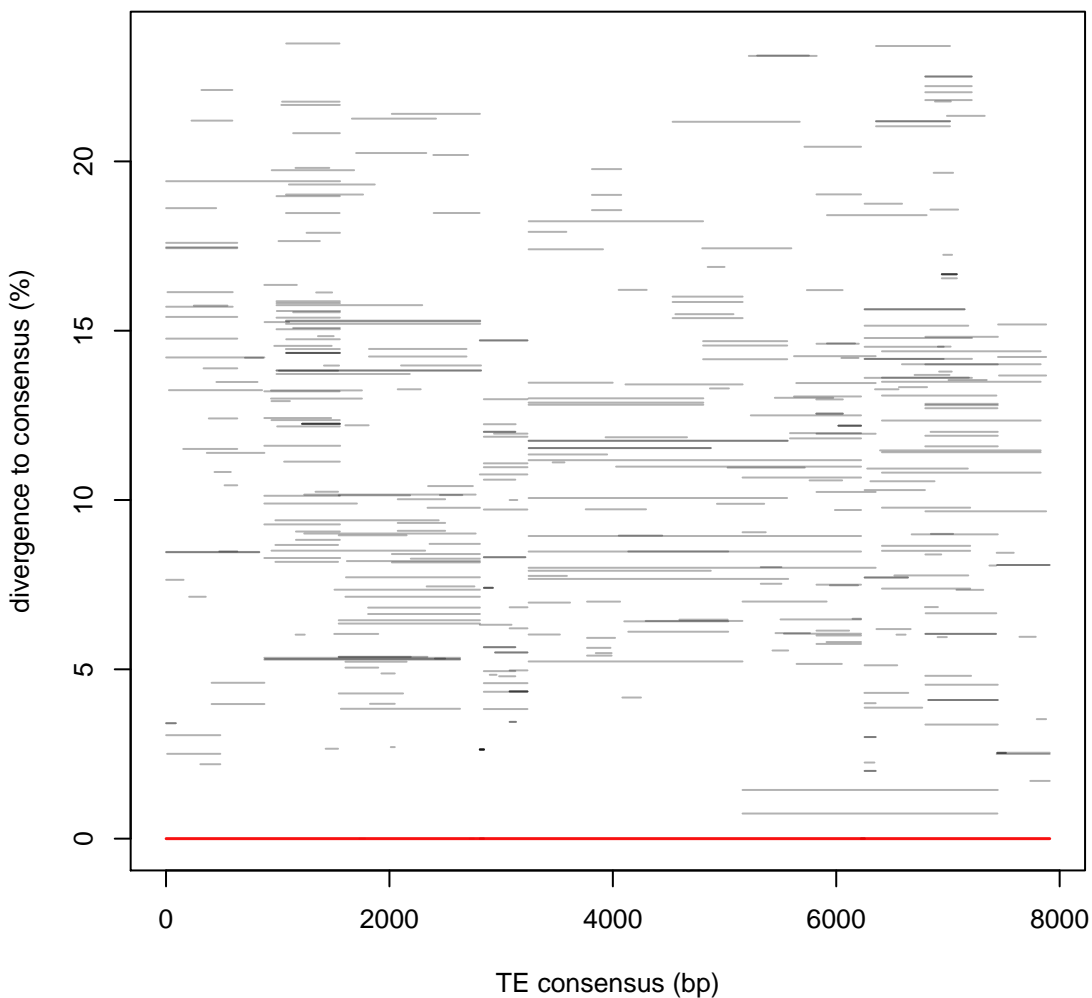
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
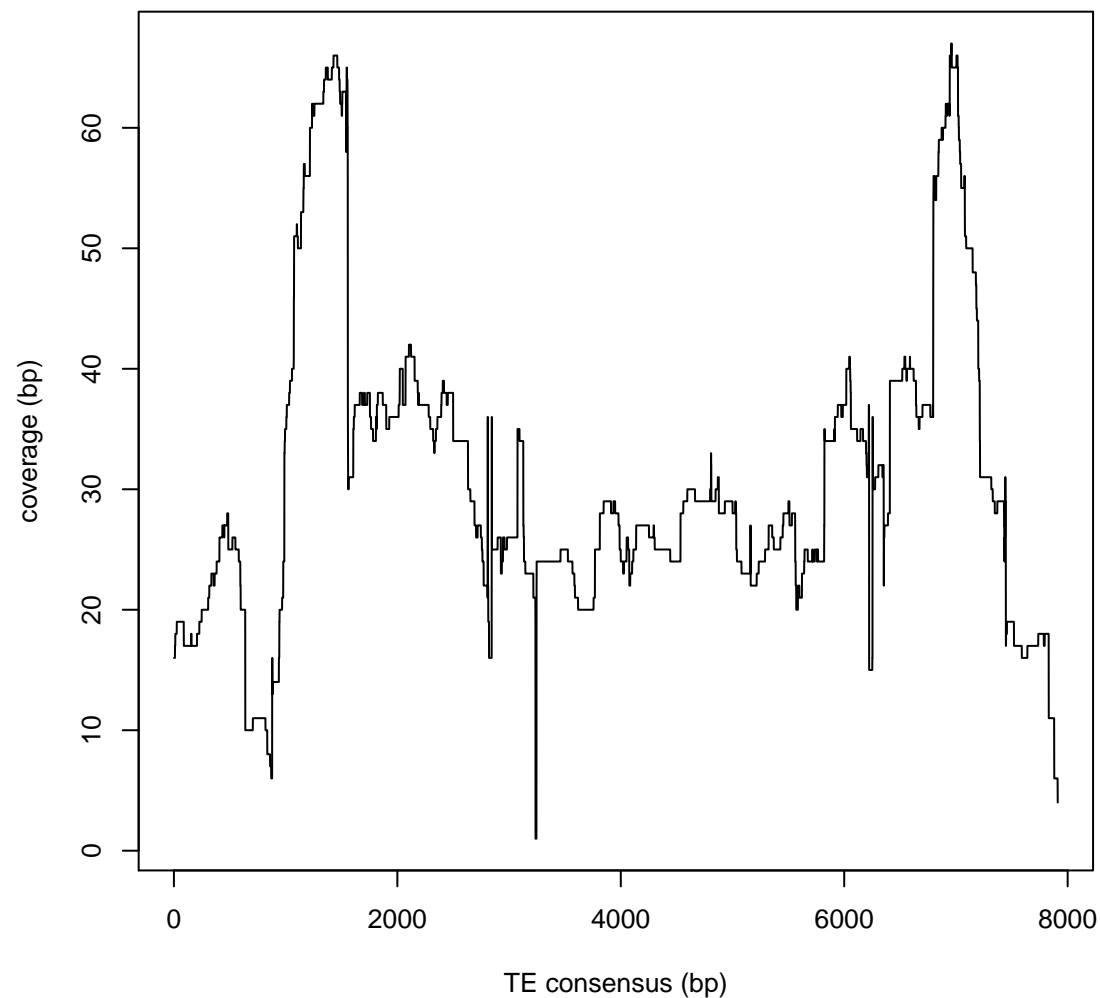
TE consensus (bp)

TE consensus self dotplot (blastn)

Dmel−L−B123−Map1−LTR/Gypsy (bp)

Dmel−L−B123−Map1−LTR/Gypsy (bp)
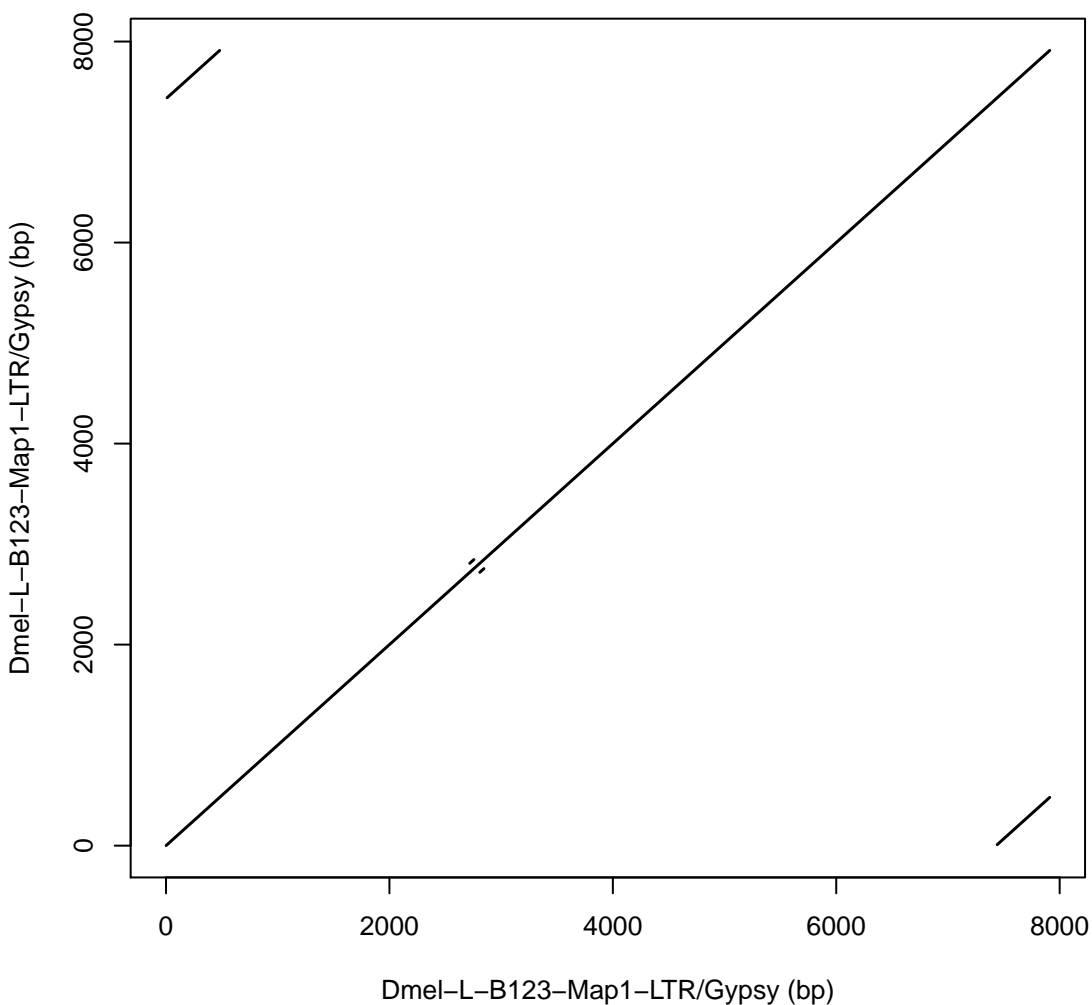
TE consensus structure and protein hits

Gypsy12_gag LTR/Gypsy
Gypsy−21_DAn_pol LTR/Gypsy
Gypsy−21_DAn_pol LTR/Gypsy
Copia−4_DGri LTR/Copia
Gypsy−21_DAn_pol LTR/Gypsy
Gypsy−10_DTa_pol LTR/Gypsy

Dmel−L−B123−Map1−LTR/Gypsy (bp)