TE: Dmel−L−B447−Map1−CLASSI/LTR/Gypsy
consensus size: 2922bp; fragments: 178; full length: 1 (>=2629.8bp)

divergence to consensus (%)
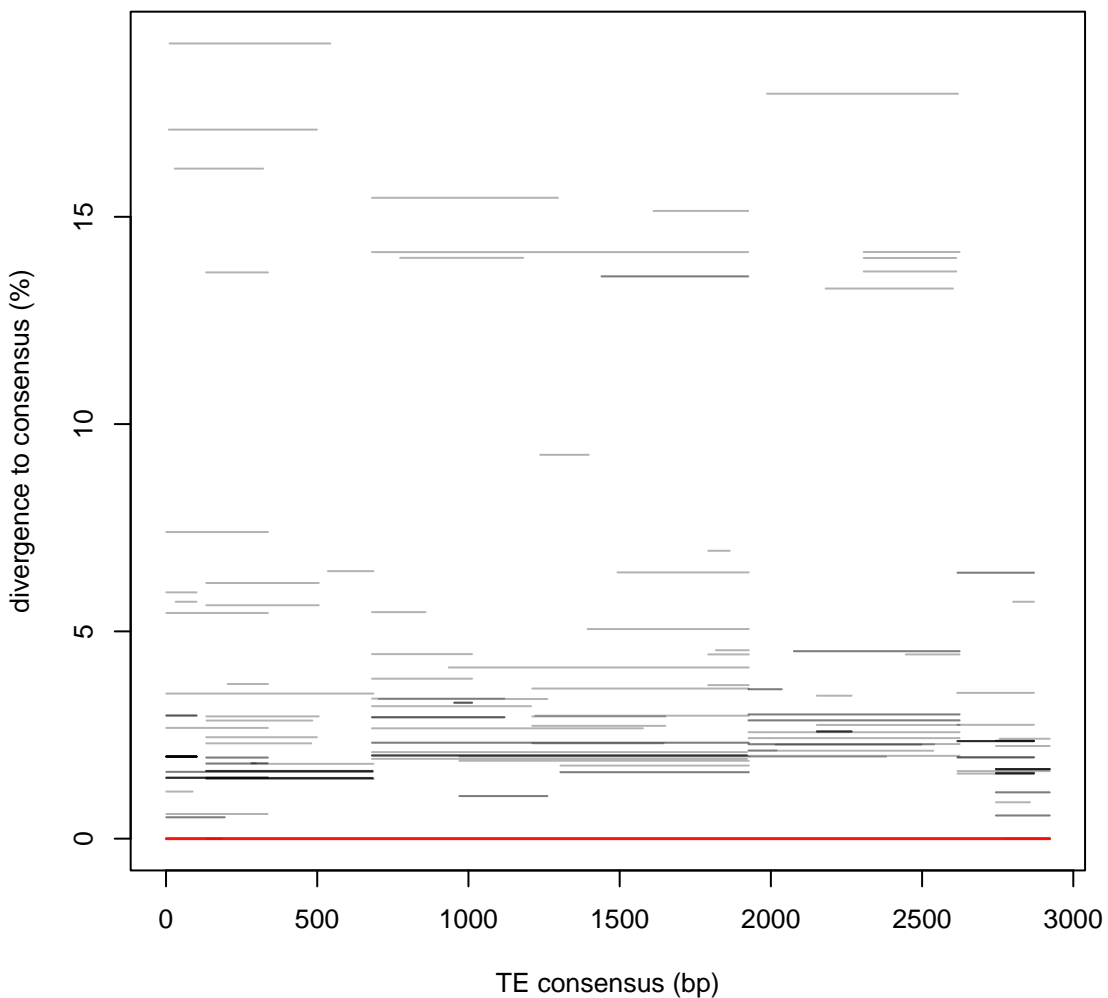
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
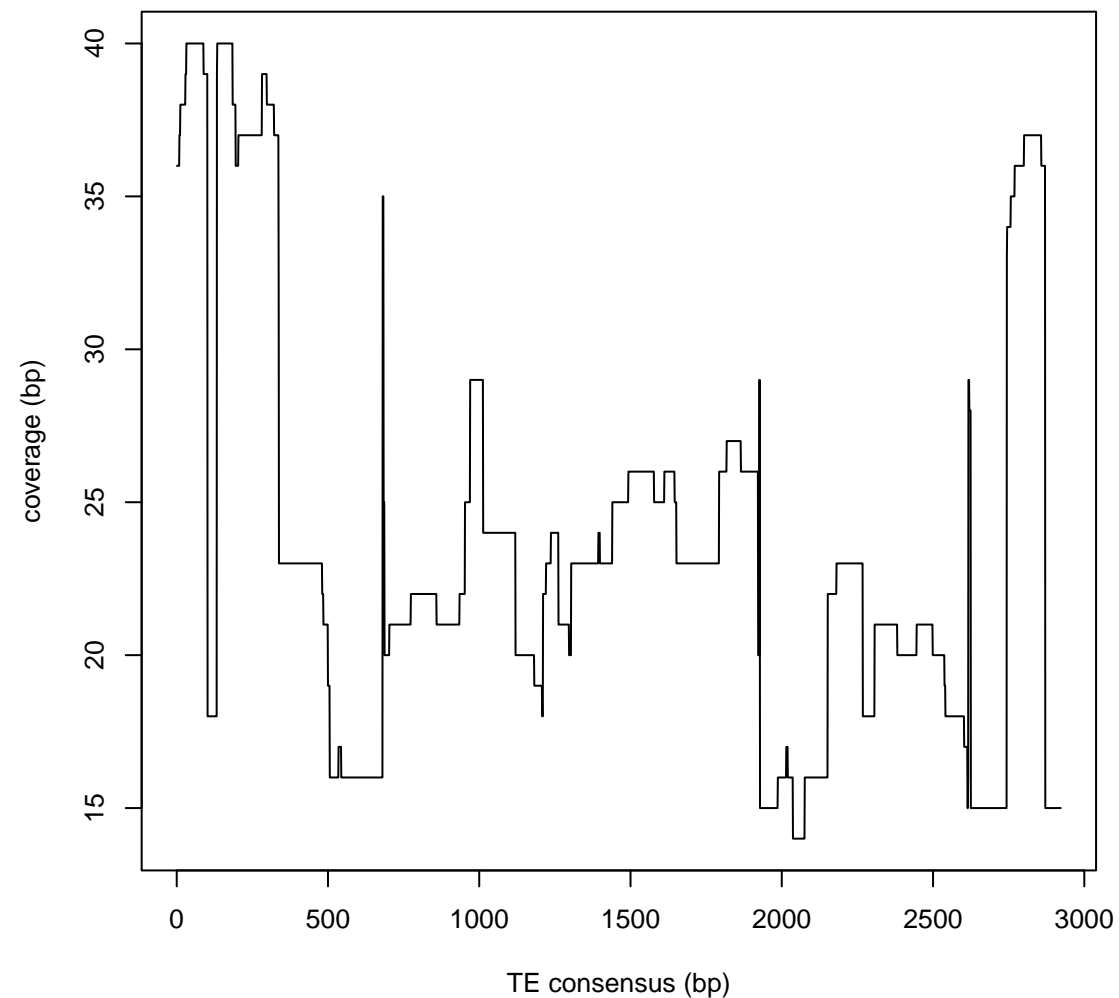
TE consensus (bp)

TE consensus self dotplot (blastn)

Dmel−L−B447−Map1−CLASSI/LTR/Gypsy (bp)

Dmel−L−B447−Map1−CLASSI/LTR/Gypsy (bp)
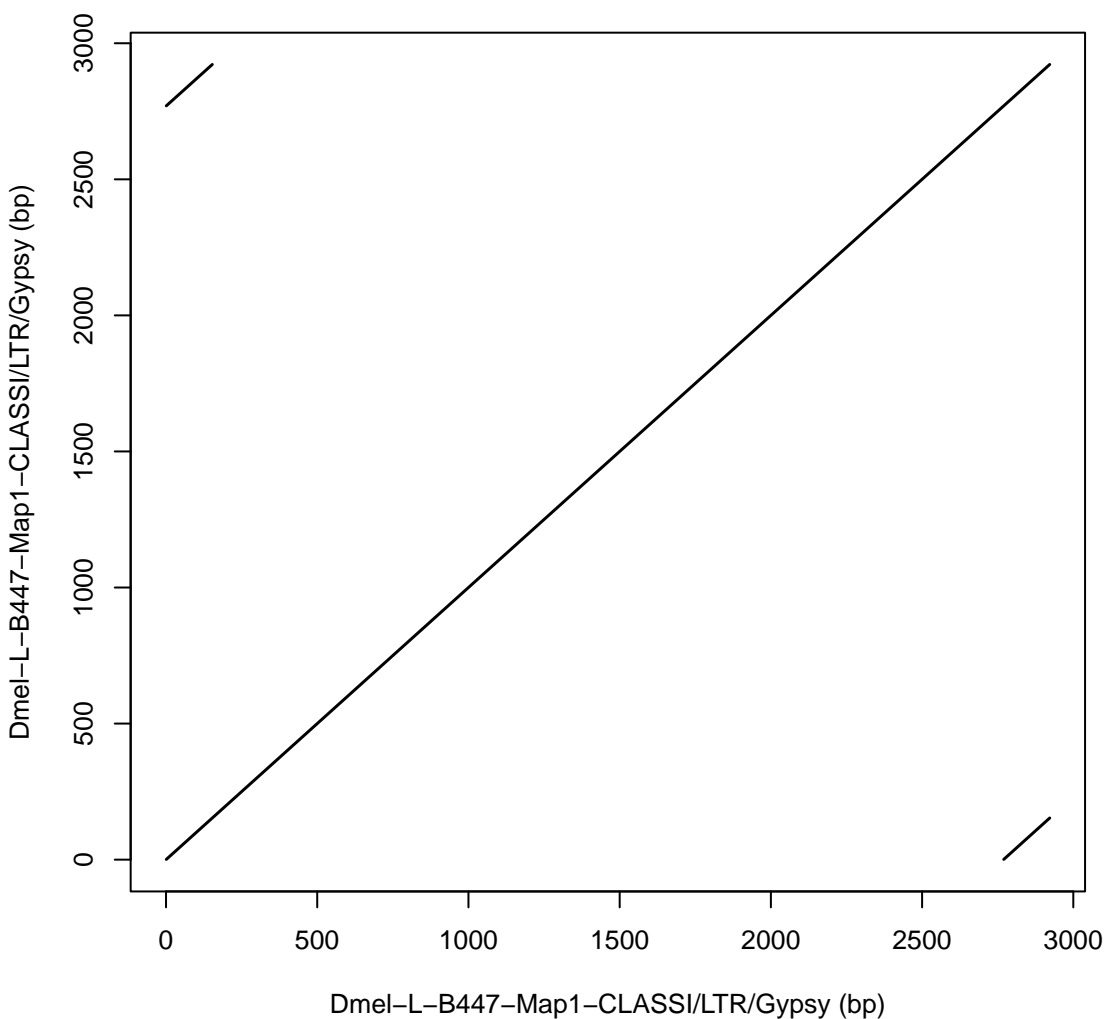
TE consensus structure and protein hits

Gypsy−6_DBp LTR/Gypsy

Gypsy−30_DWil LTR/Gypsy

Gypsy−4_DWil LTR/Gypsy

Dmel−L−B447−Map1−CLASSI/LTR/Gypsy (bp)