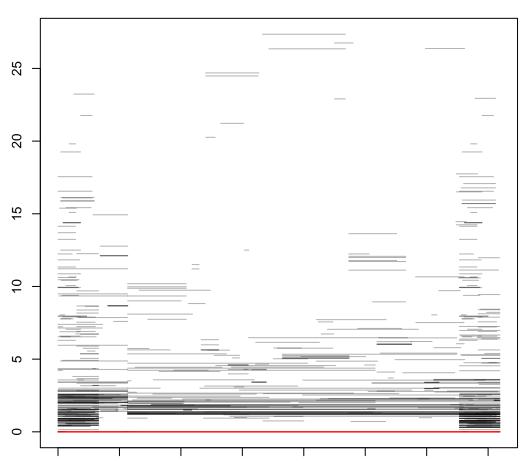
TE: Dmel-L-B159-Map1-CLASSI/LTR/Gypsy consensus size: 7191bp; fragments: 625; full length: 1 (>=6471.9bp)



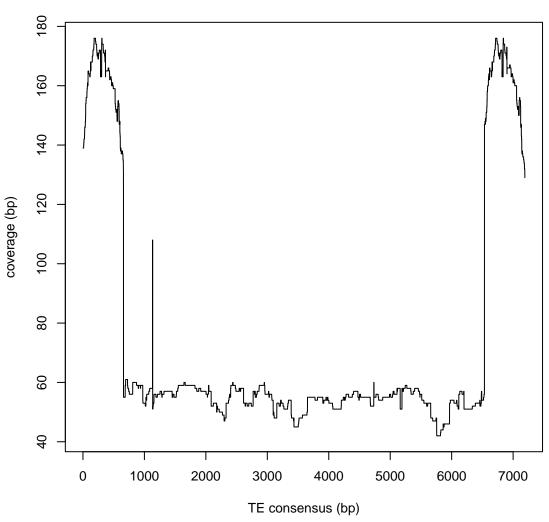
divergence to consensus (%)

0

1000

2000

TE consensus genomic coverage



TE consensus self dotplot (blastn)

TE consensus (bp)

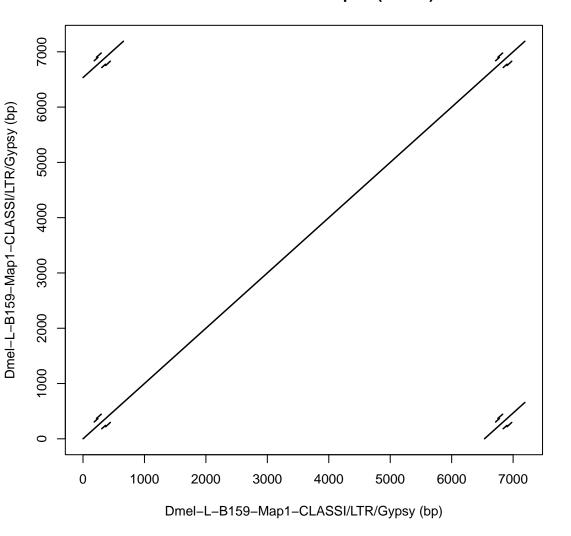
3000

4000

5000

6000

7000



TE consensus structure and protein hits

