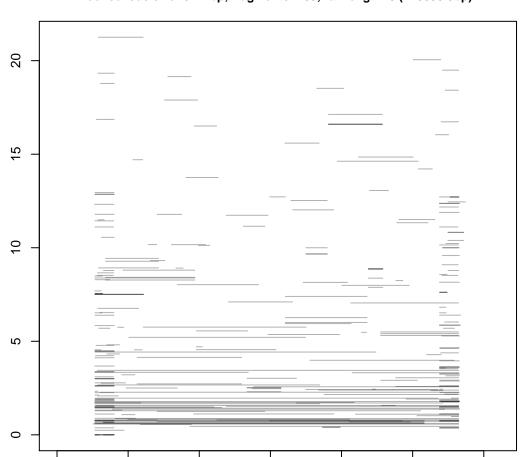
TE: Dmel-B-P226.10-Map3-LTR/Gypsy consensus size: 6212bp; fragments: 289; full length: 0 (>=5590.8bp)



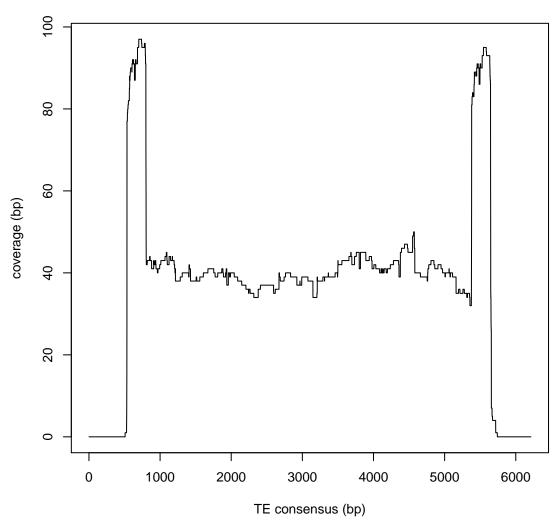
divergence to consensus (%)

0

1000

2000

TE consensus genomic coverage



TE consensus self dotplot (blastn)

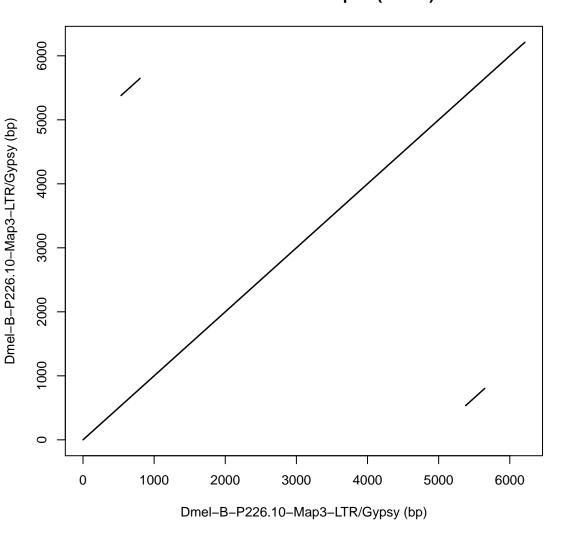
TE consensus (bp)

3000

4000

5000

6000



TE consensus structure and protein hits

