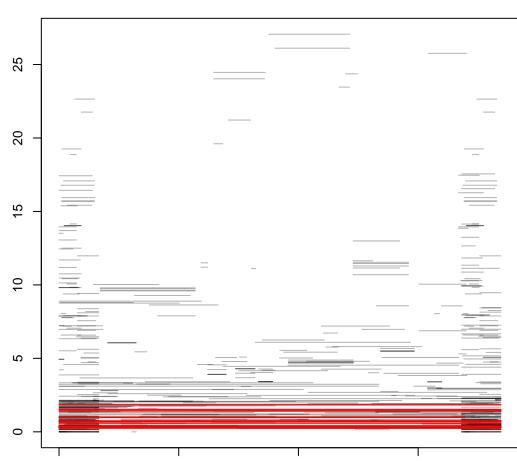
TE: Dmel-B-G1493-Map4\_reversed-CLASSI/LTR/Gypsy consensus size: 7382bp; fragments: 583; full length: 16 (>=6643.8bp)

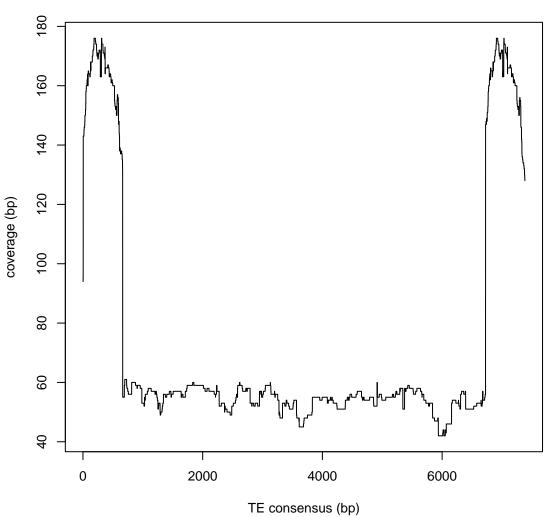


divergence to consensus (%)

0

2000

## TE consensus genomic coverage

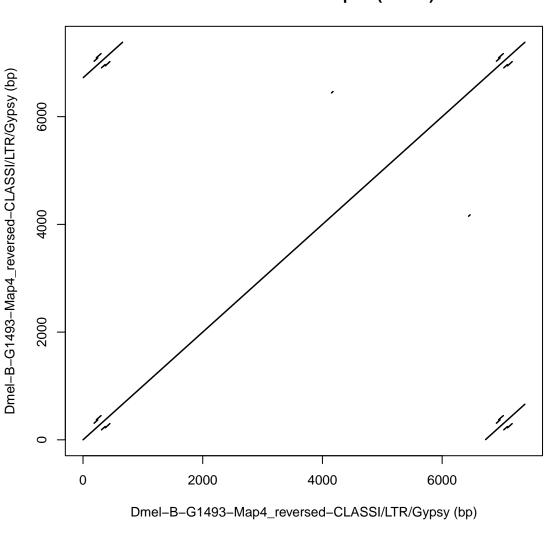


TE consensus self dotplot (blastn)

TE consensus (bp)

4000

6000



TE consensus structure and protein hits

