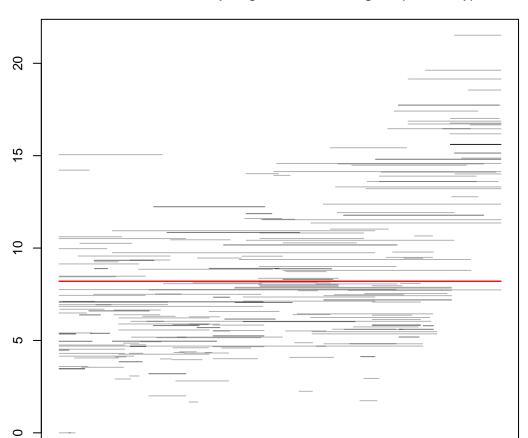
TE: Dmel-B-G745-Map4-CLASSI/LINE/nan consensus size: 2969bp; fragments: 255; full length: 1 (>=2672.1bp)



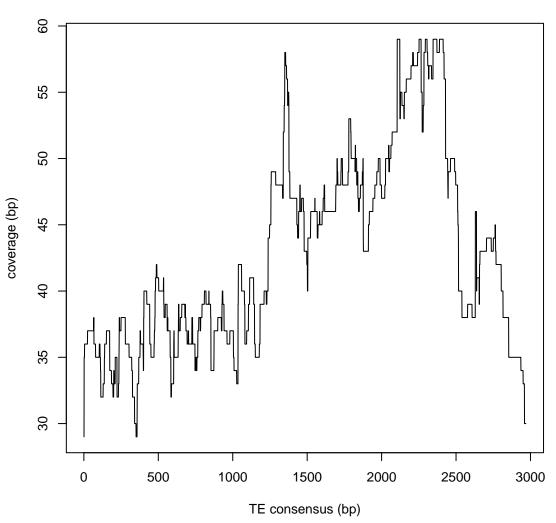
divergence to consensus (%)

0

500

1000

## TE consensus genomic coverage



## TE consensus self dotplot (blastn)

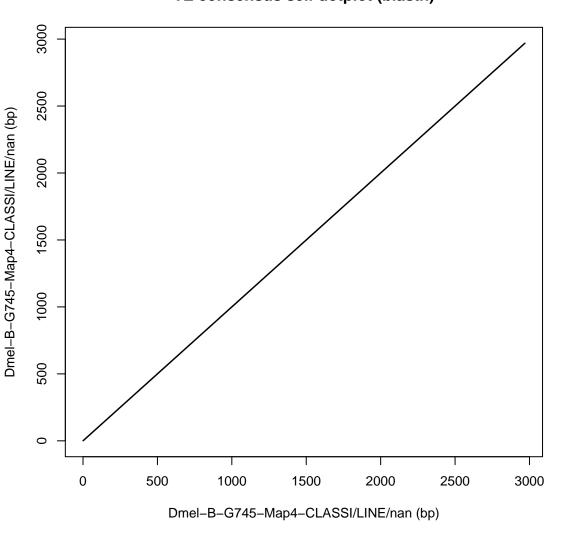
1500

TE consensus (bp)

2000

2500

3000



## TE consensus structure and protein hits

