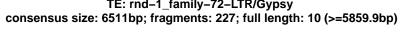
TE: rnd-1_family-72-LTR/Gypsy consensus size: 6511bp; fragments: 227; full length: 10 (>=5859.9bp)





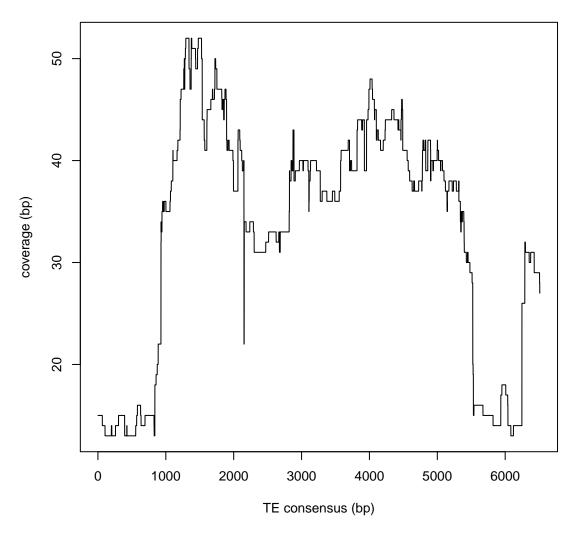
divergence to consensus (%)

0

1000

2000

TE consensus genomic coverage



TE consensus self dotplot (blastn)

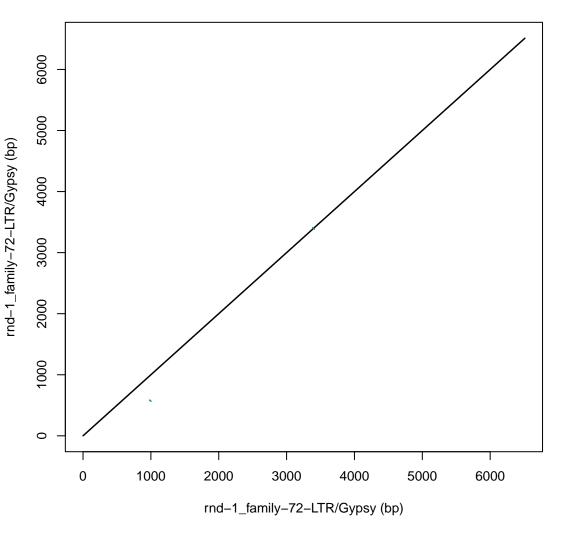
TE consensus (bp)

3000

4000

5000

6000



TE consensus structure and protein hits

