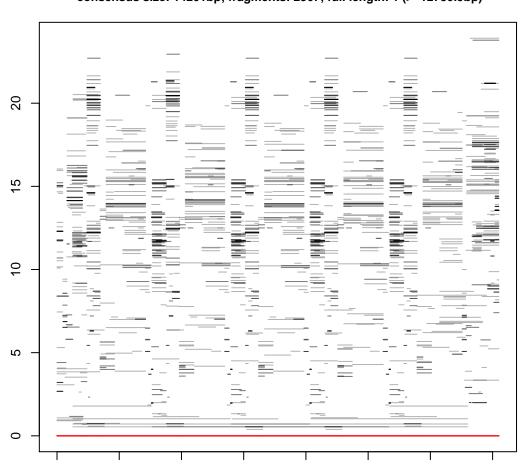
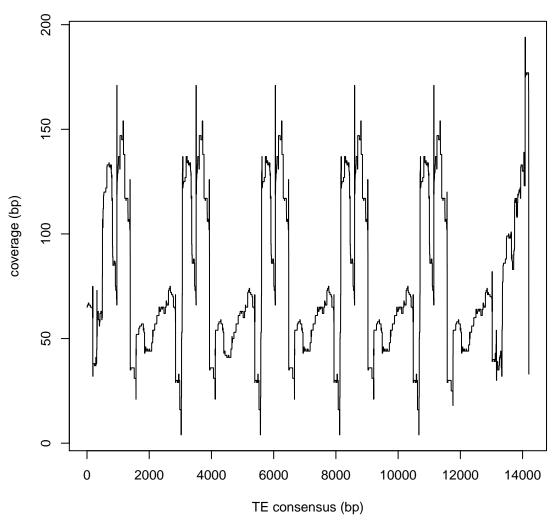
TE: rnd-1_family-15-LINE consensus size: 14201bp; fragments: 2567; full length: 1 (>=12780.9bp)



divergence to consensus (%)

TE consensus genomic coverage



TE consensus self dotplot (blastn)

TE consensus (bp)

8000

10000

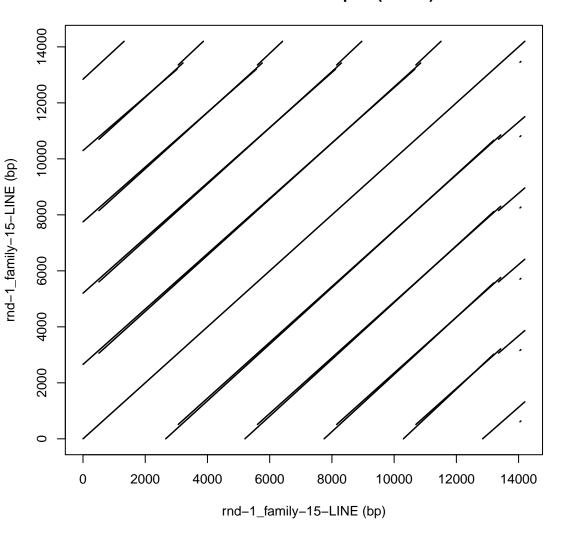
12000

14000

6000

2000

4000



TE consensus structure and protein hits

