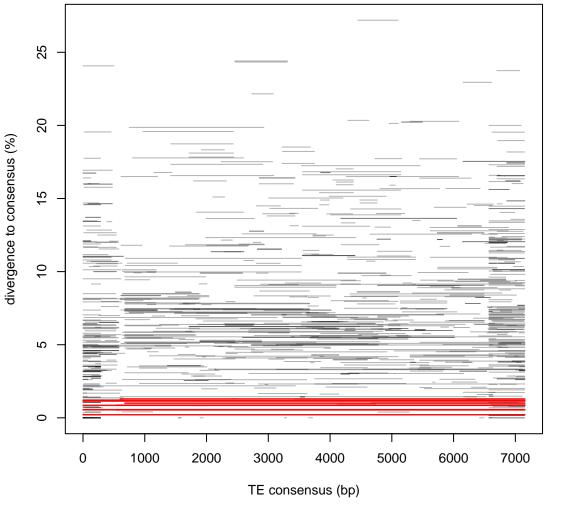
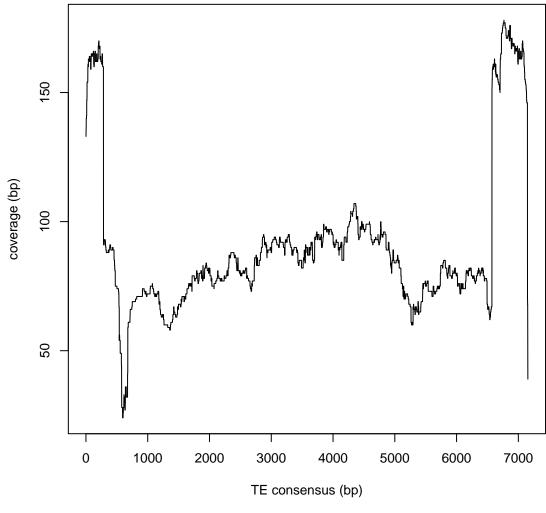
TE consensus genomic coverage





TE consensus self dotplot (blastn)

md-1_family-188-LTR/Gypsy (bp)

0 1000 2000 3000 4000 5000 6000 7000

3000

4000

rnd-1_family-188-LTR/Gypsy (bp)

5000

6000

7000

1000

2000

TE consensus structure and protein hits

