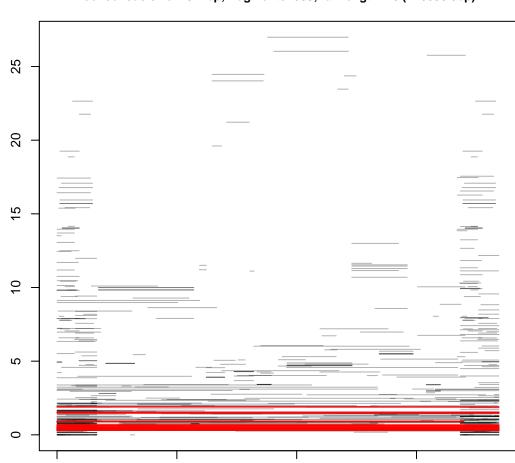
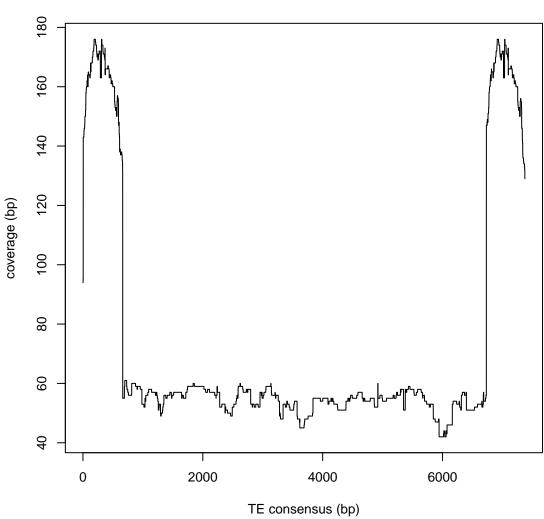
TE: rnd-5_family-1216-LTR/Gypsy consensus size: 7374bp; fragments: 583; full length: 16 (>=6636.6bp)



divergence to consensus (%)

0

TE consensus genomic coverage



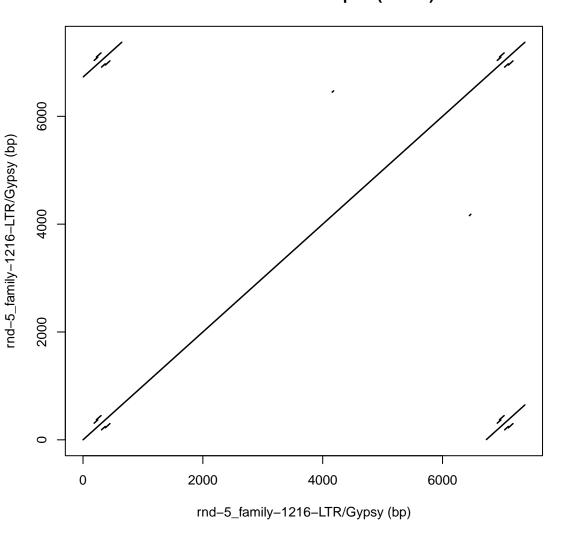
TE consensus self dotplot (blastn)

TE consensus (bp)

4000

6000

2000



TE consensus structure and protein hits

