TE: rnd-1\_family-470\_s\_1-LTR/Gypsy consensus size: 7382bp; fragments: 583; full length: 16 (>=6643.8bp)

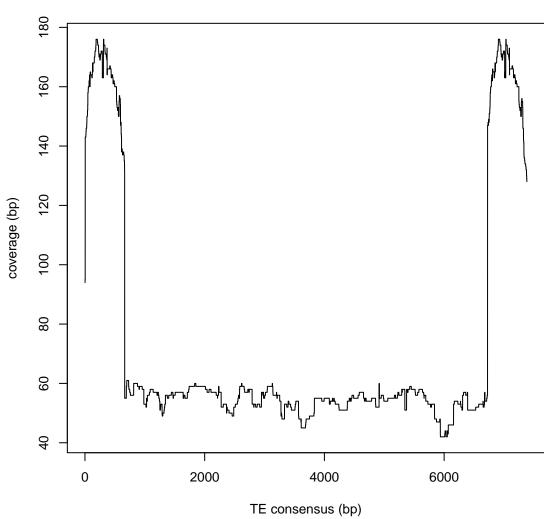
divergence to consensus (%)

0

0

2000

## TE consensus genomic coverage

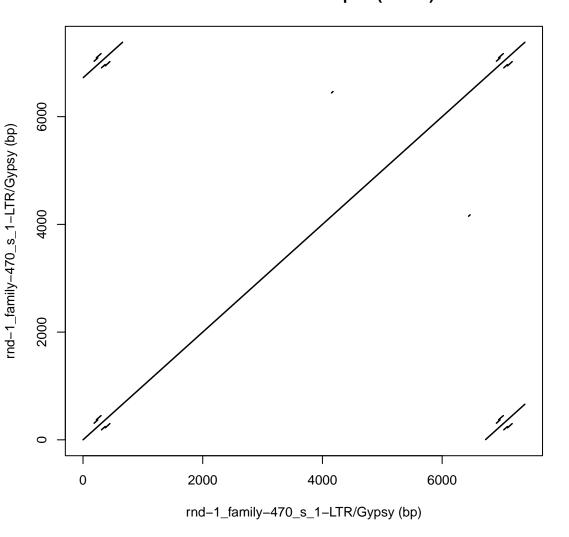


TE consensus self dotplot (blastn)

TE consensus (bp)

4000

6000



TE consensus structure and protein hits

