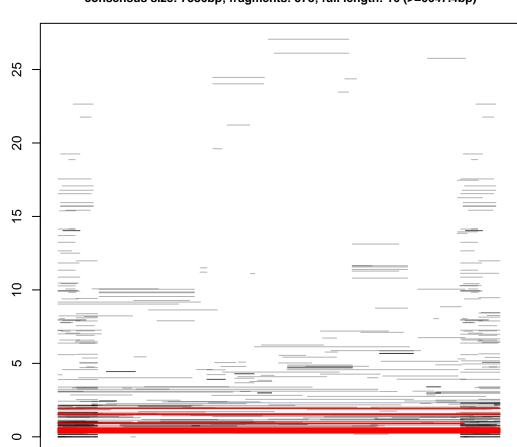
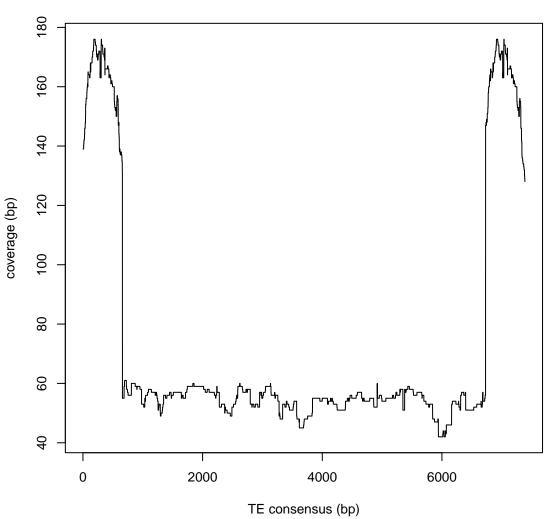
TE: rnd-1_family-512_s_1-LTR/Gypsy consensus size: 7386bp; fragments: 578; full length: 16 (>=6647.4bp)



divergence to consensus (%)

TE consensus genomic coverage



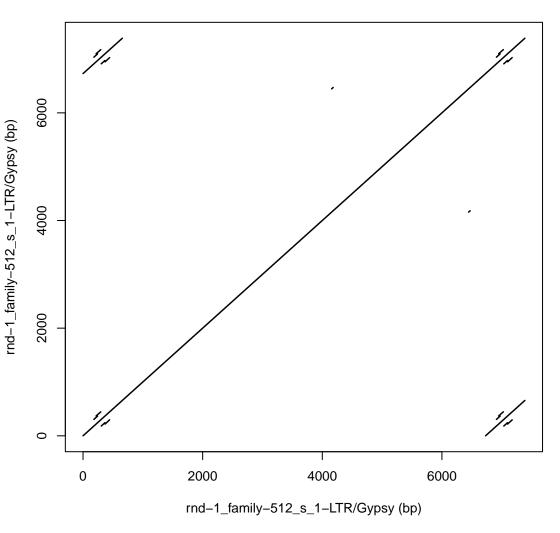
TE consensus self dotplot (blastn)

TE consensus (bp)

4000

6000

2000



TE consensus structure and protein hits

