**TE: rnd−5_family−884−LINE/I−Jockey**
**consensus size: 3554bp; fragments: 78; full length: 0 (>=3198.6bp)**

divergence to consensus (%)
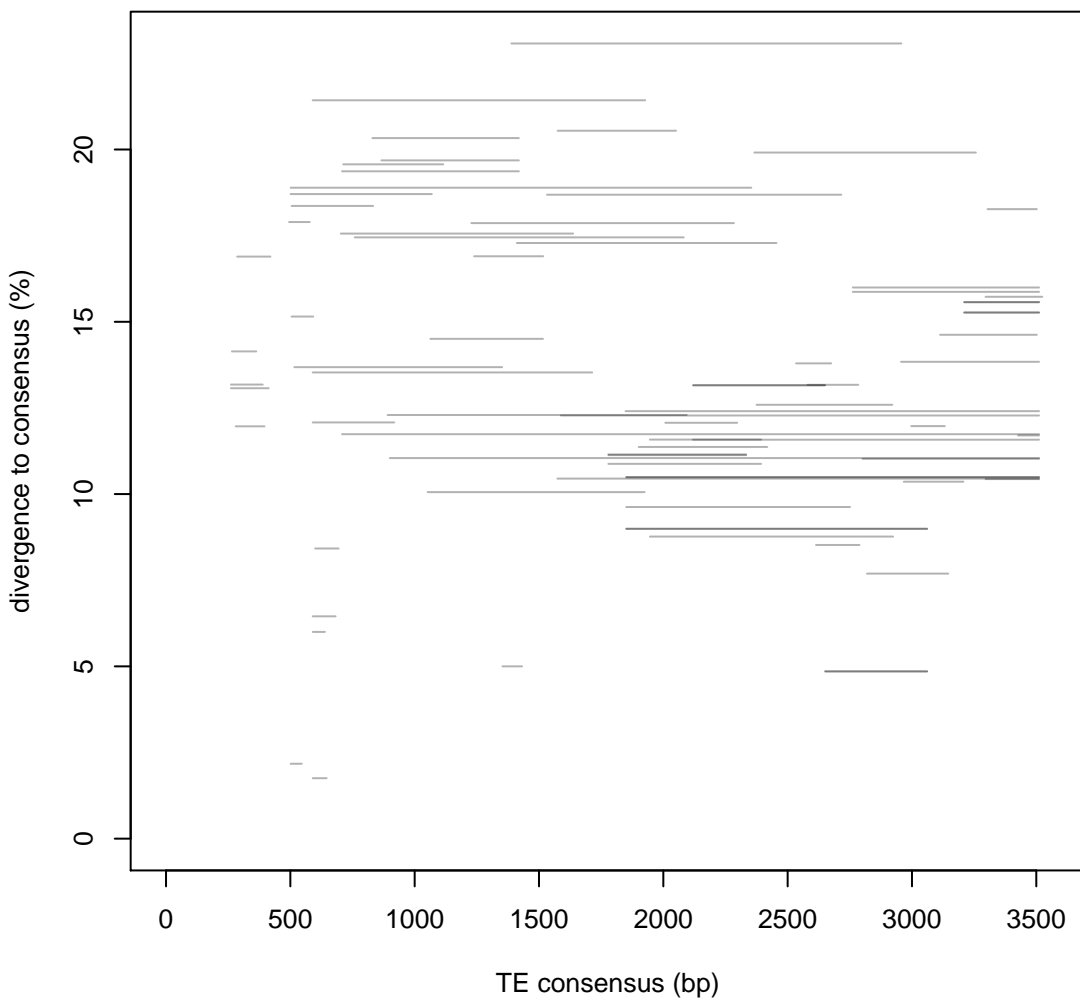
TE consensus (bp)

**TE consensus genomic coverage**

coverage (bp)
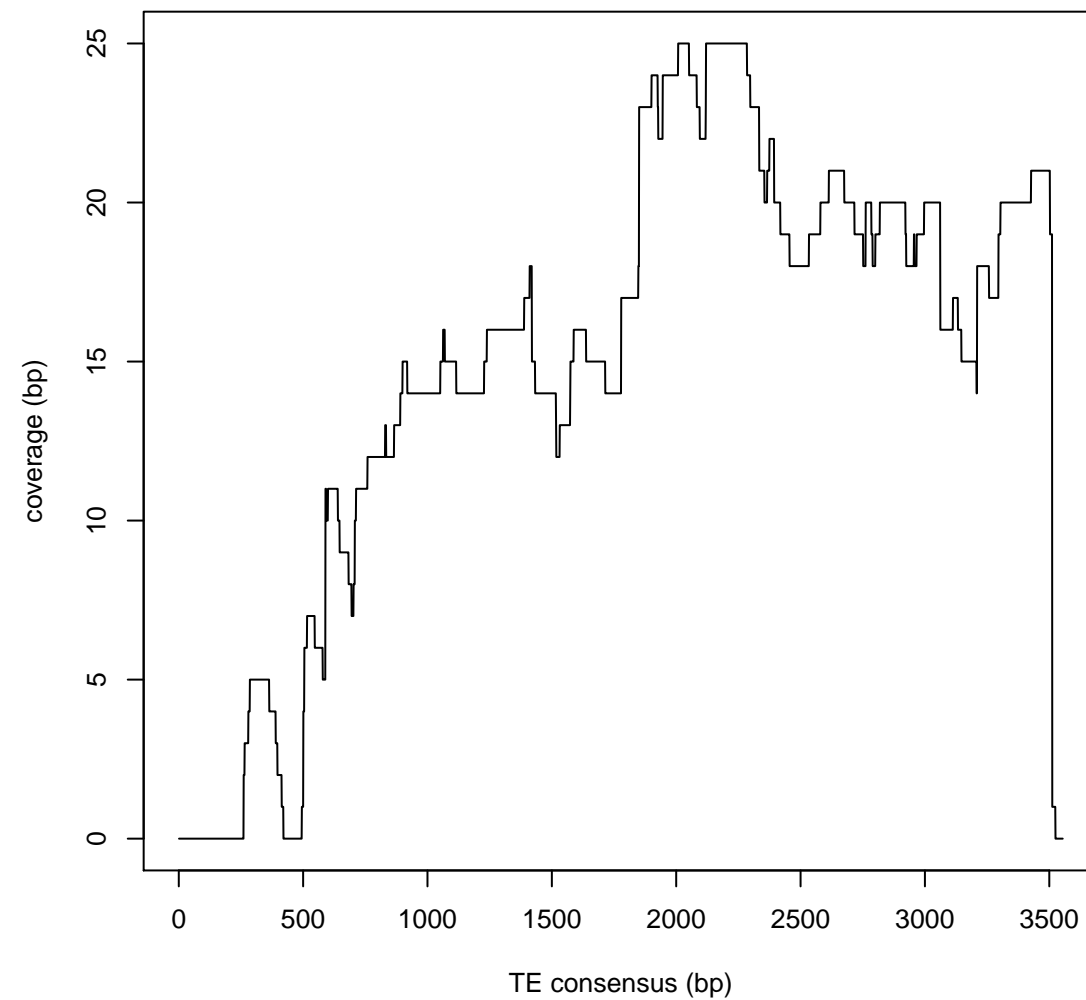
TE consensus (bp)

**TE consensus self dotplot (blastn)**

rnd−5_family−884−LINE/I−Jockey (bp)

rnd−5_family−884−LINE/I−Jockey (bp)
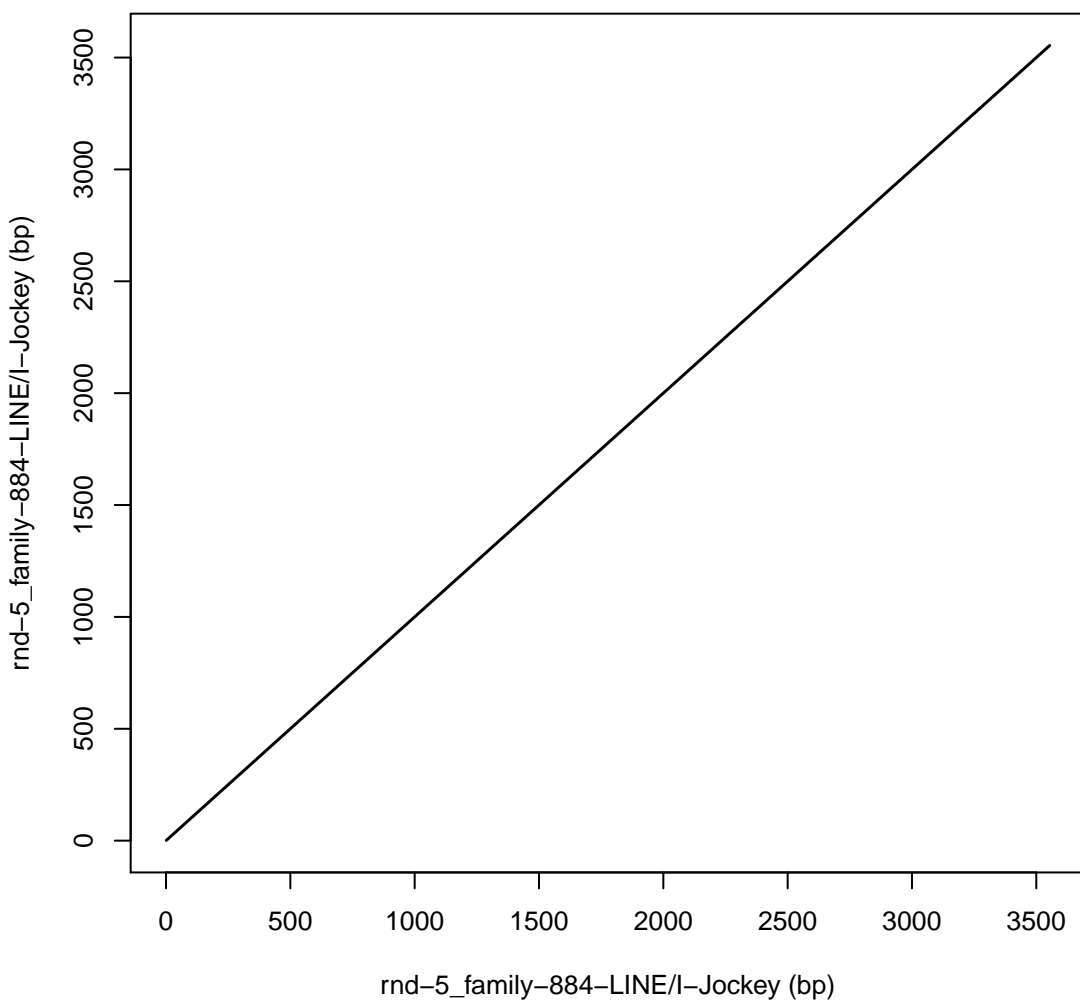
**TE consensus structure and protein hits**

rnd−5_family−884−LINE/I−Jockey (bp)