TE: rnd−1_family−542−LINE/I−Jockey
consensus size: 5009bp; fragments: 86; full length: 12 (>=4508.1bp)

divergence to consensus (%)
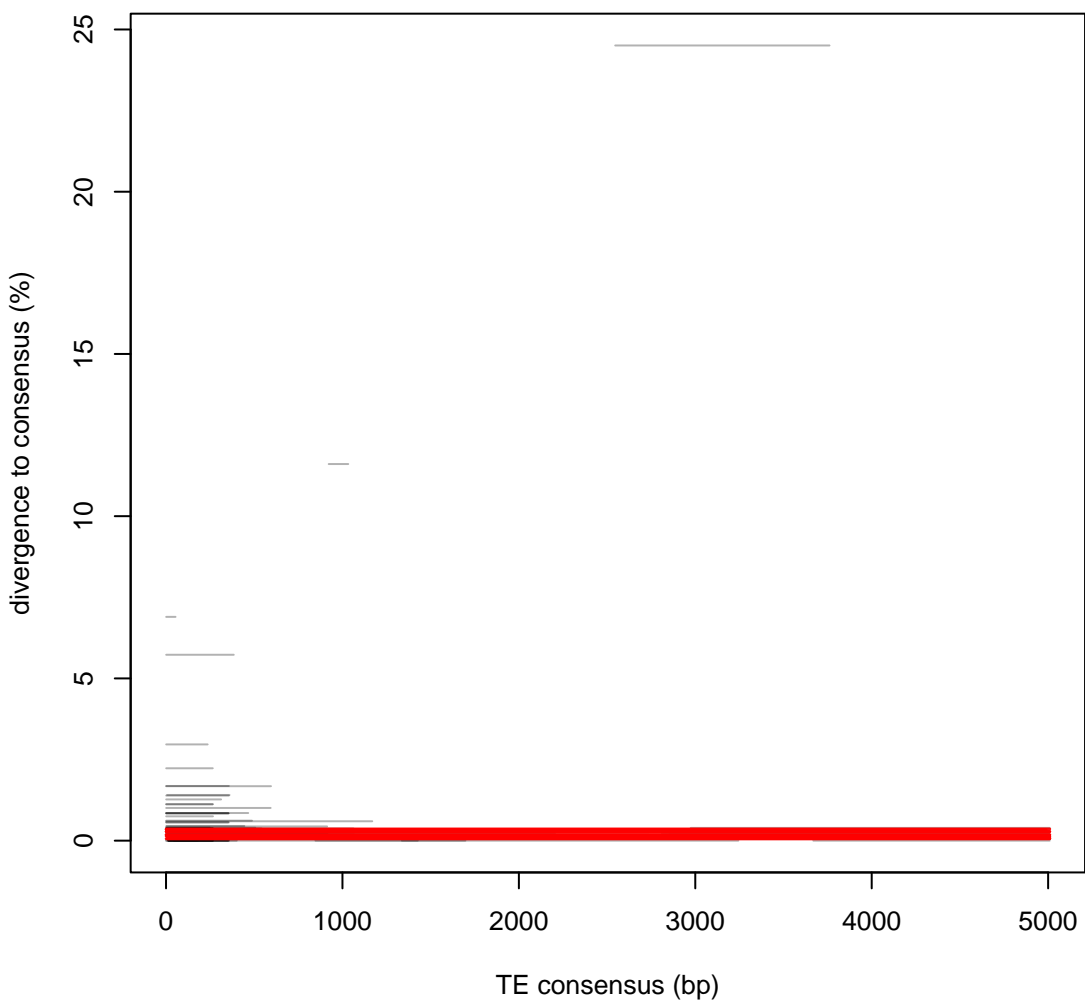
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
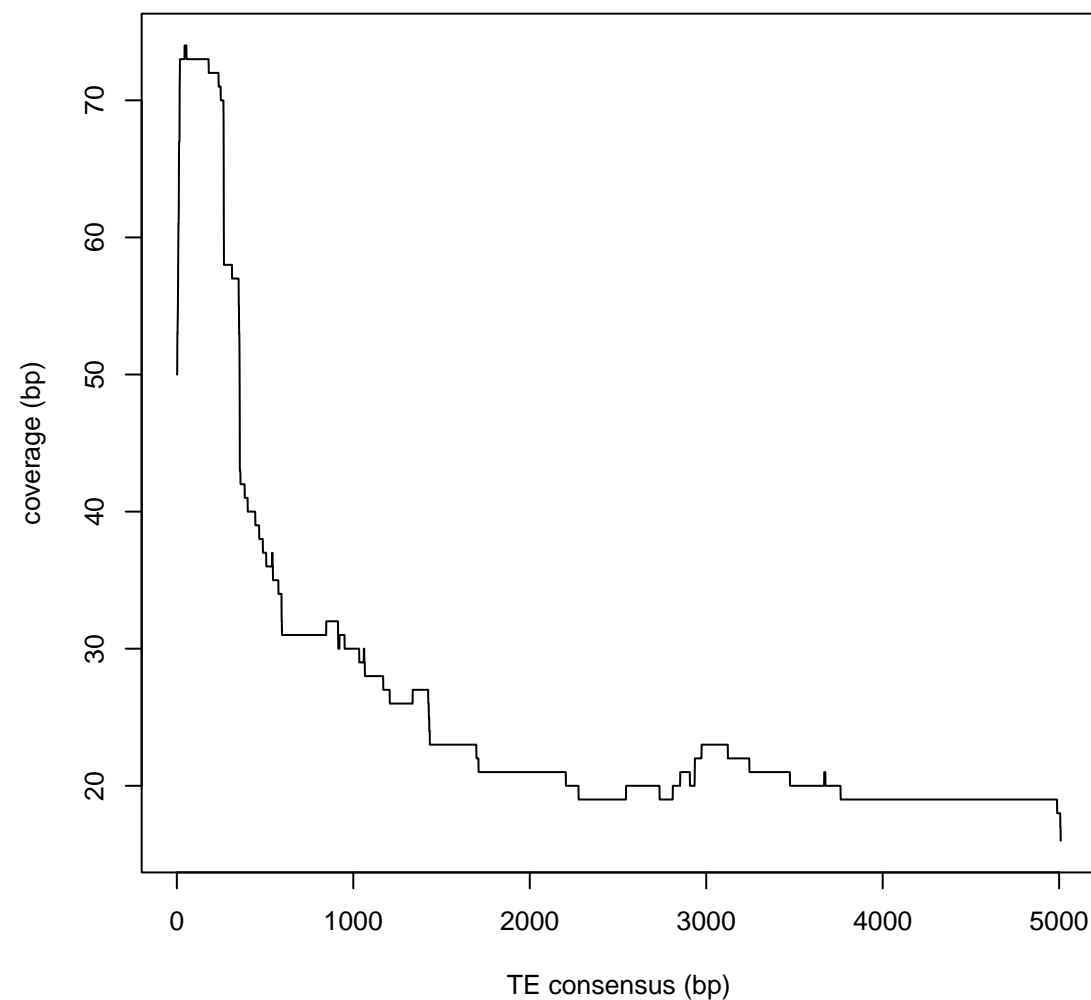
TE consensus (bp)

TE consensus self dotplot (blastn)

rnd−1_family−542−LINE/I−Jockey (bp)

rnd−1_family−542−LINE/I−Jockey (bp)
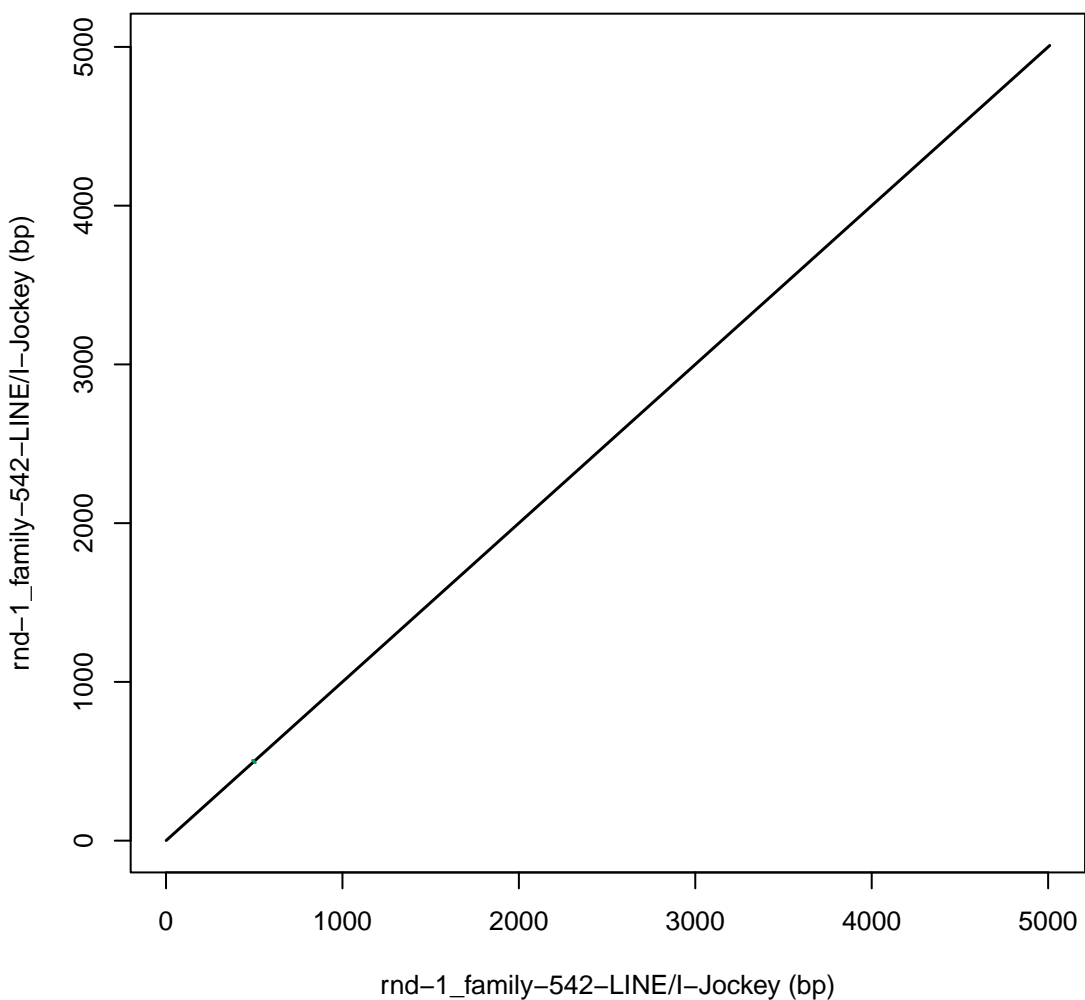
TE consensus structure and protein hits

LINEJ1_DM_pol LINE/I−Jockey

LINEJ1_DM_gag LINE/I−Jockey

rnd−1_family−542−LINE/I−Jockey (bp)