TE: rnd−1_family−462−LTR
consensus size: 2582bp; fragments: 186; full length: 1 (>=2323.8bp)

divergence to consensus (%)
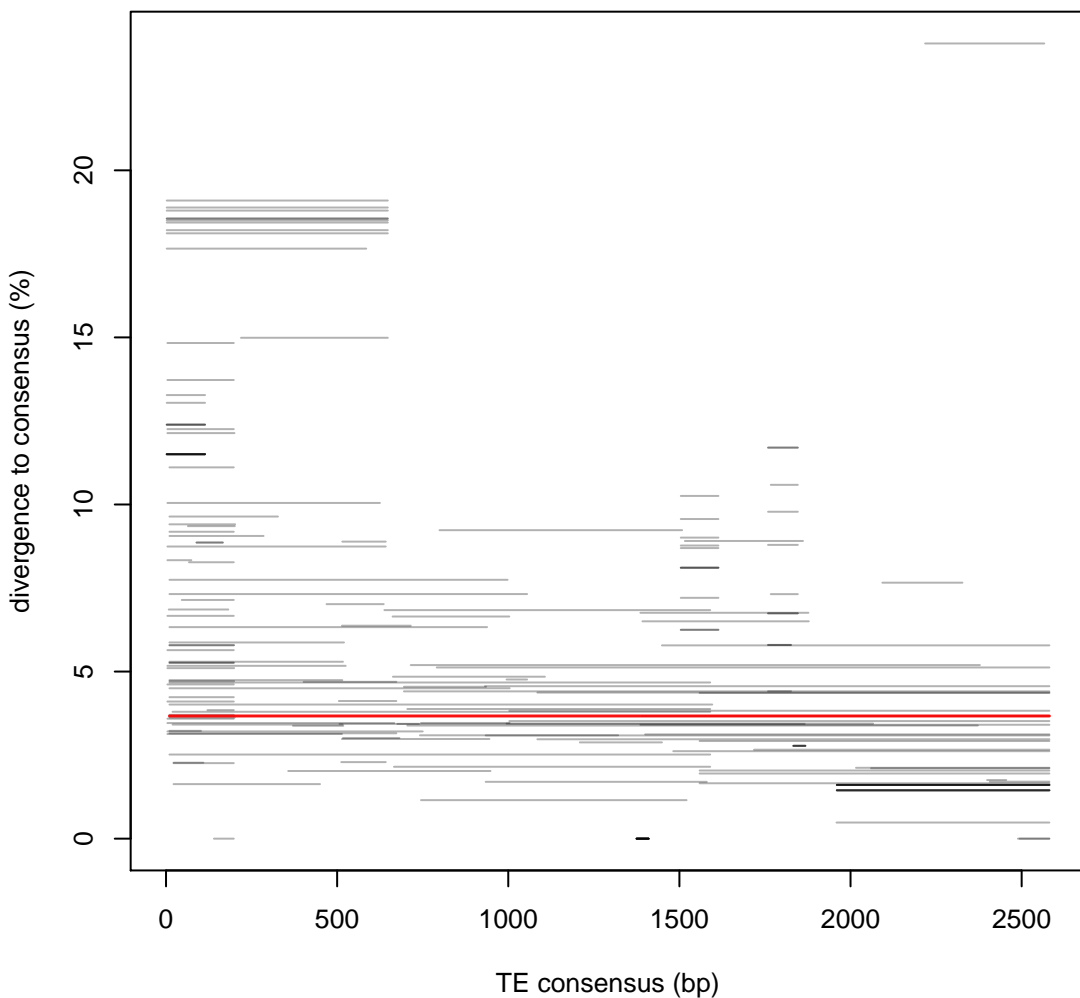
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
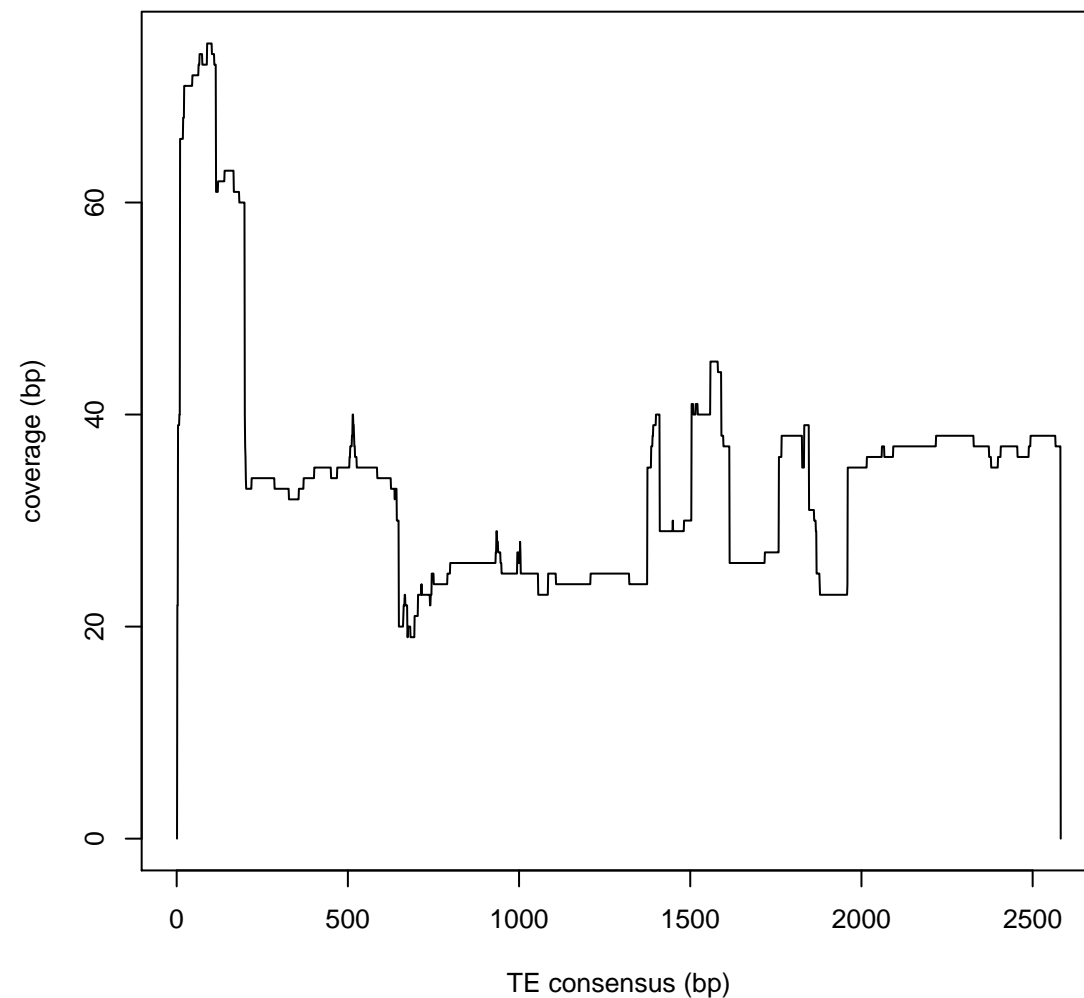
TE consensus (bp)

TE consensus self dotplot (blastn)

rnd−1_family−462−LTR (bp)

rnd−1_family−462−LTR (bp)
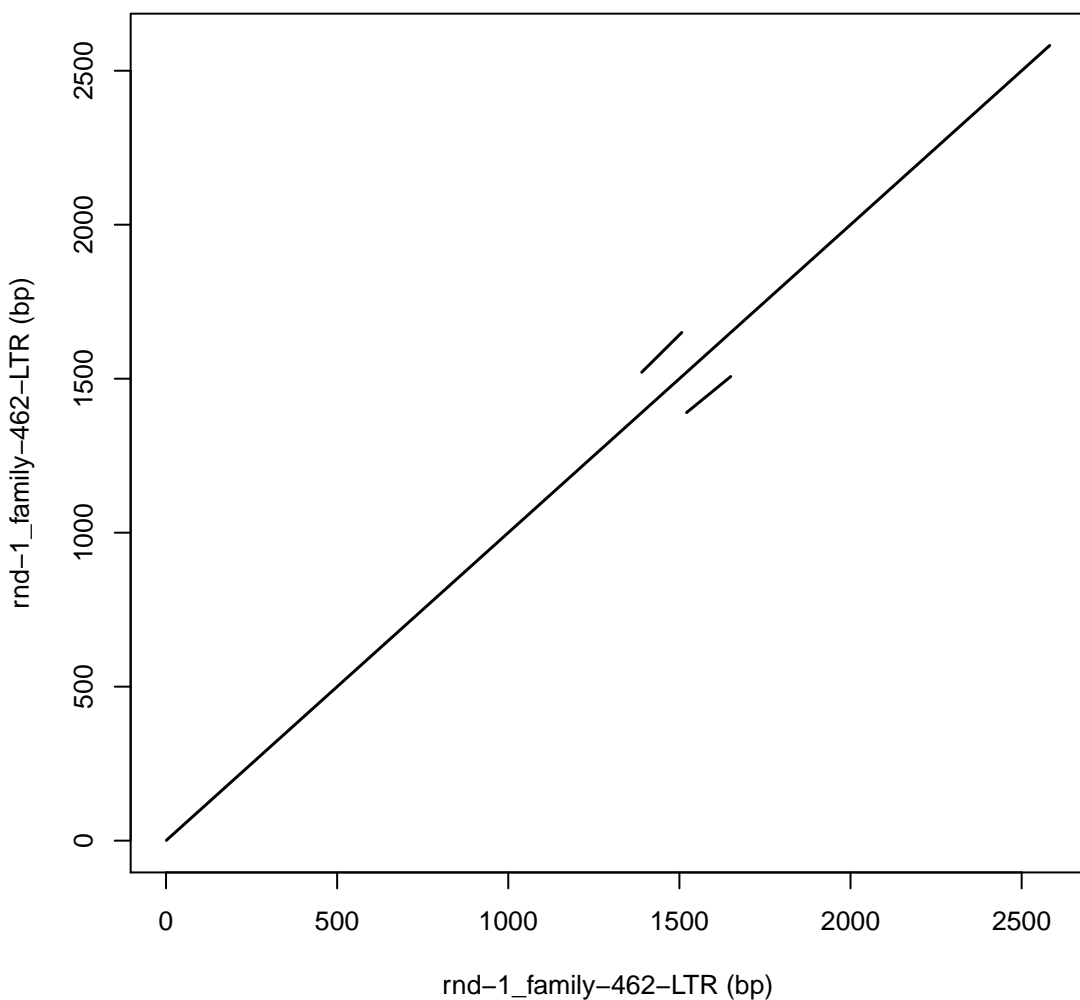
TE consensus structure and protein hits

Copia36−PTR LTR/Copia

BEL−22_DTa_pol LTR/Pa

rnd−1_family−462−LTR (bp)