TE: rnd−6_family−188−TIR
consensus size: 2259bp; fragments: 24; full length: 0 (>=2033.1bp)

TE consensus genomic coverage
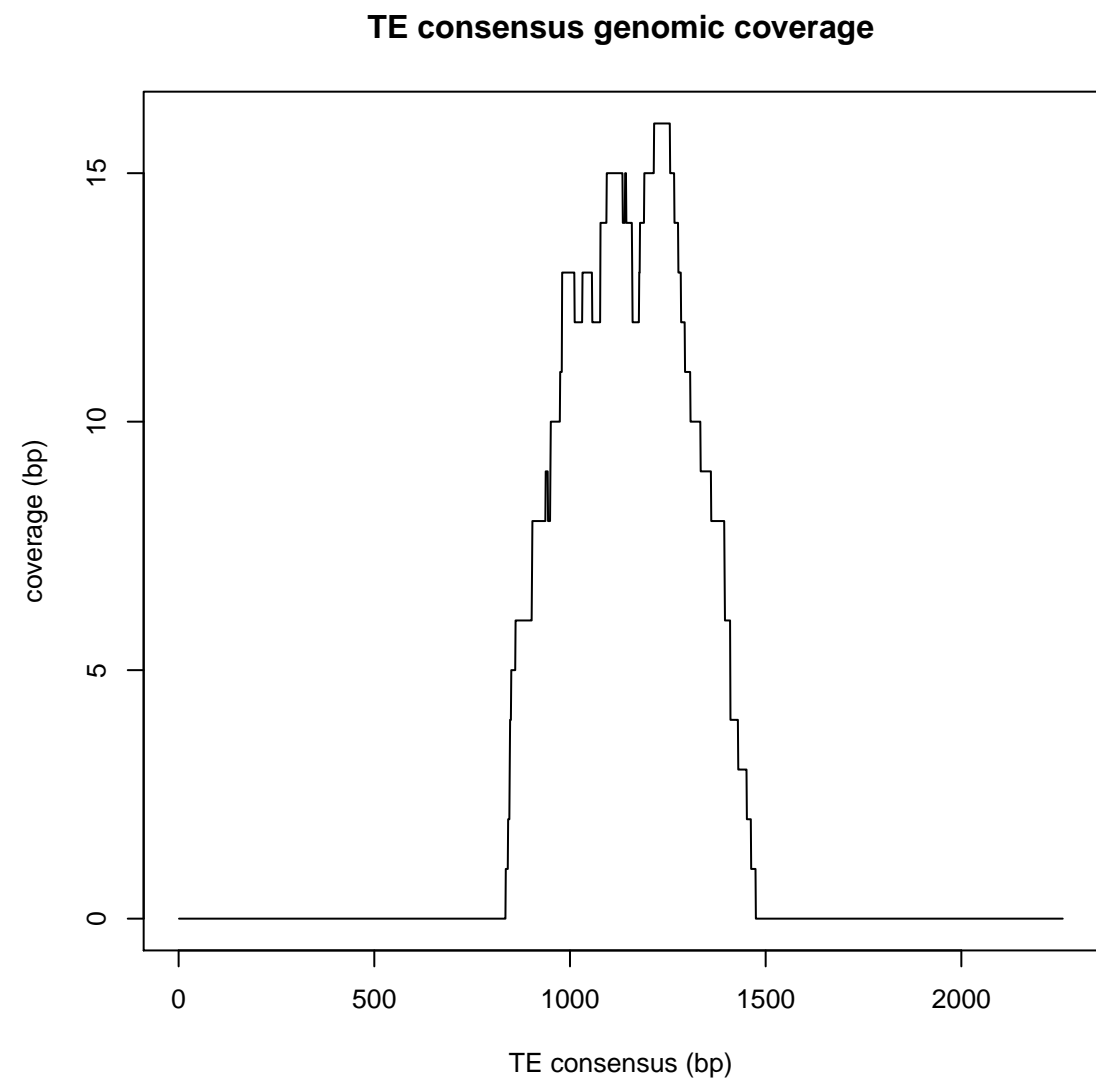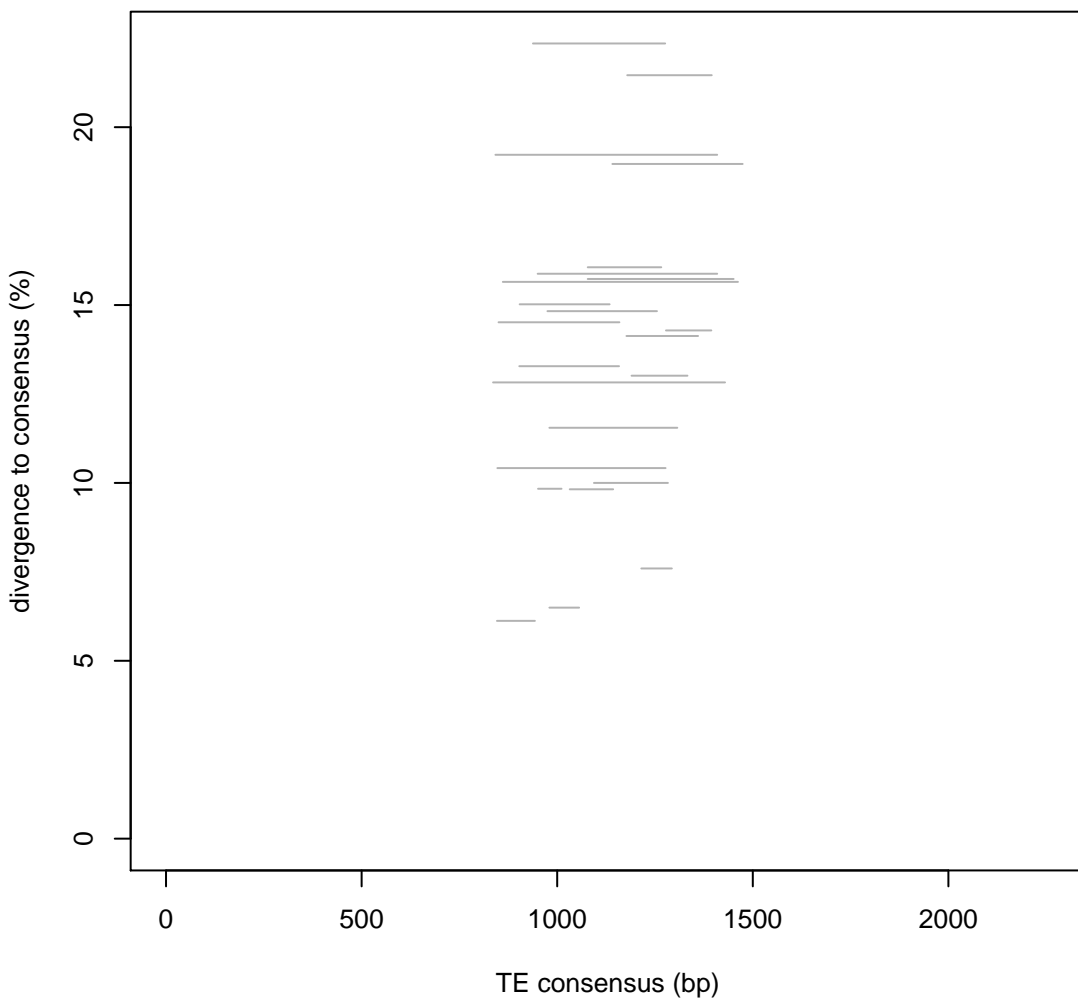
TE consensus self dotplot (blastn)
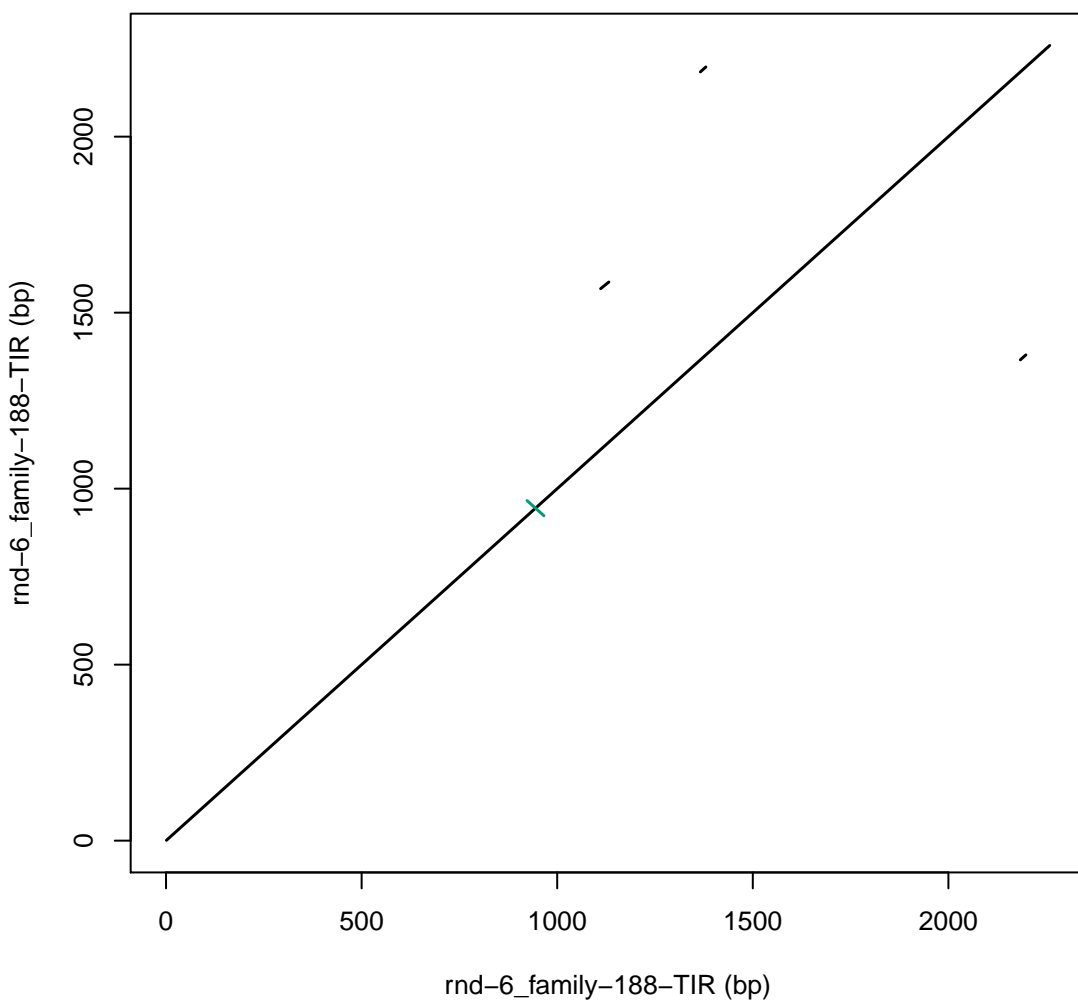
TE consensus structure and protein hits

no orf >400 bp detected