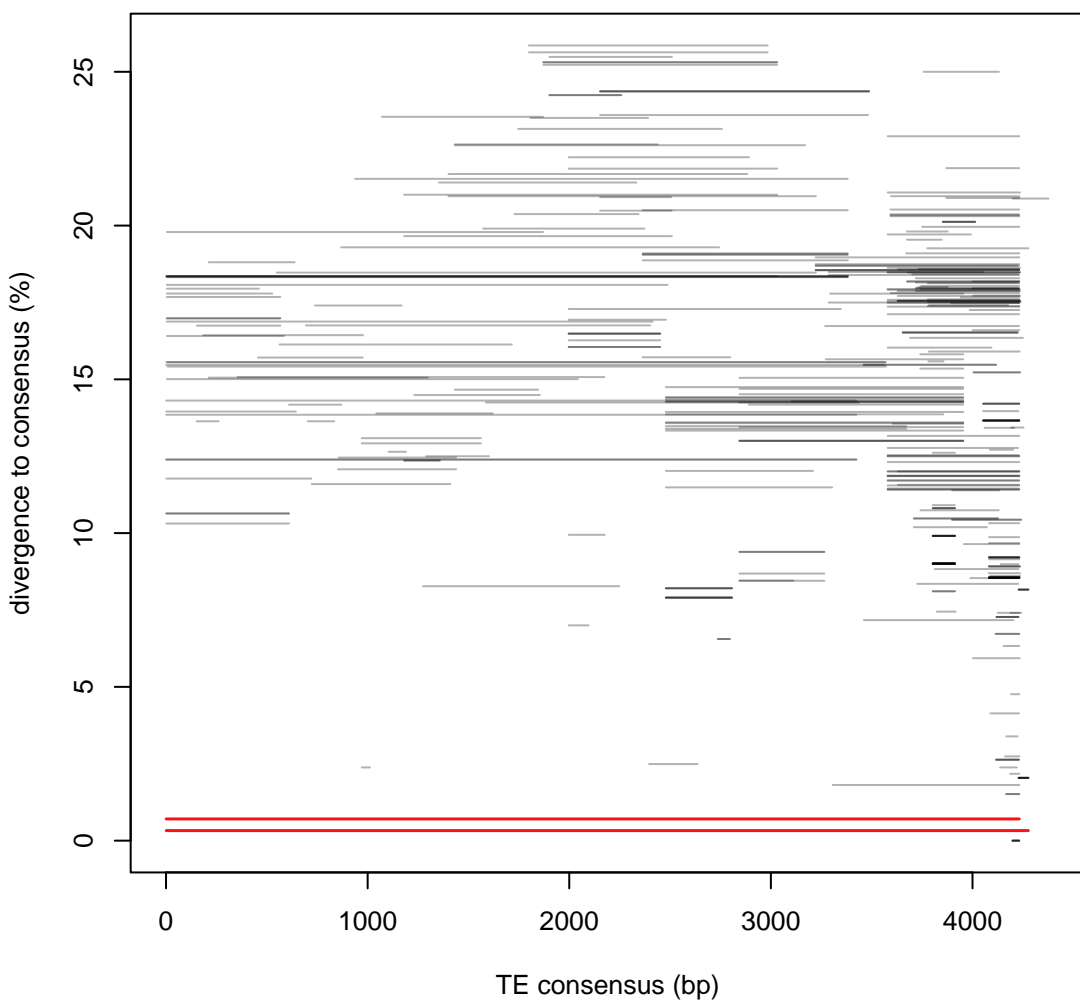
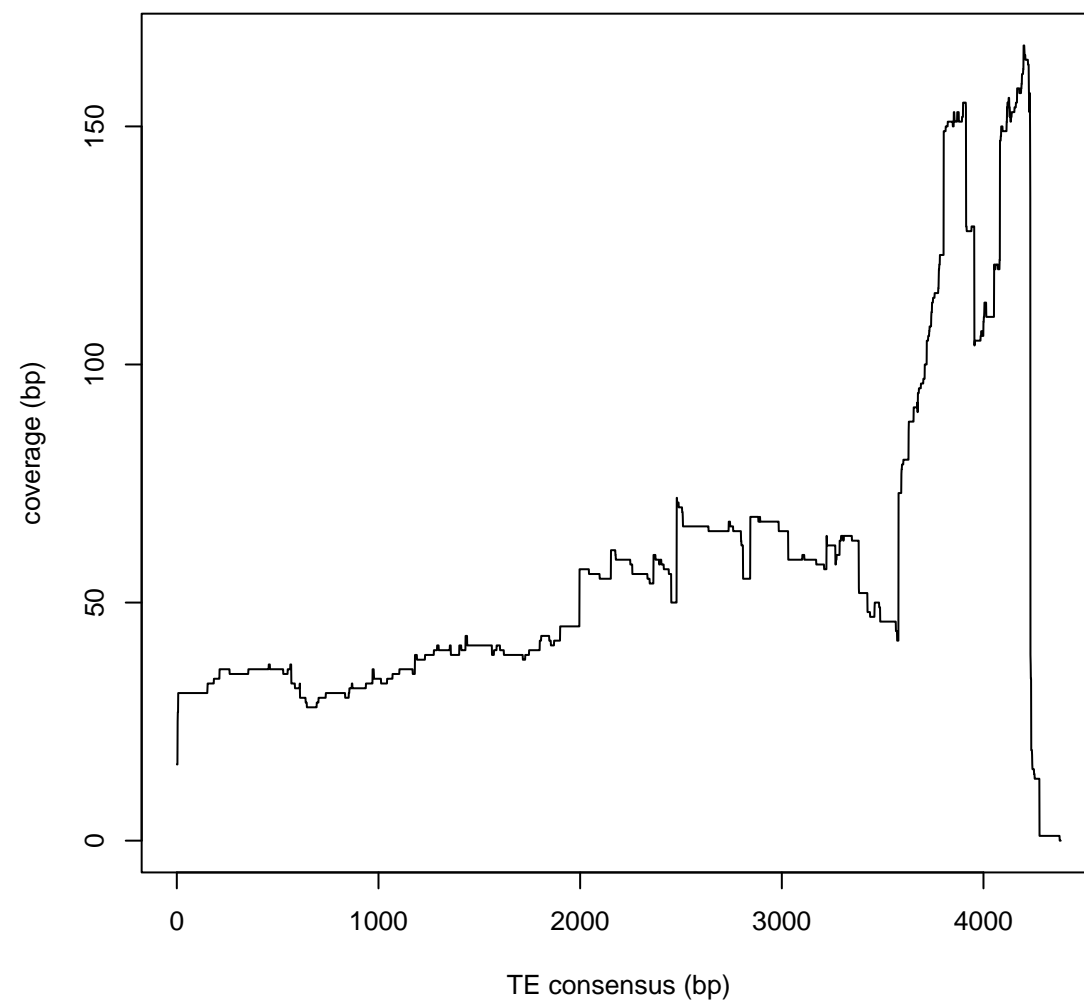


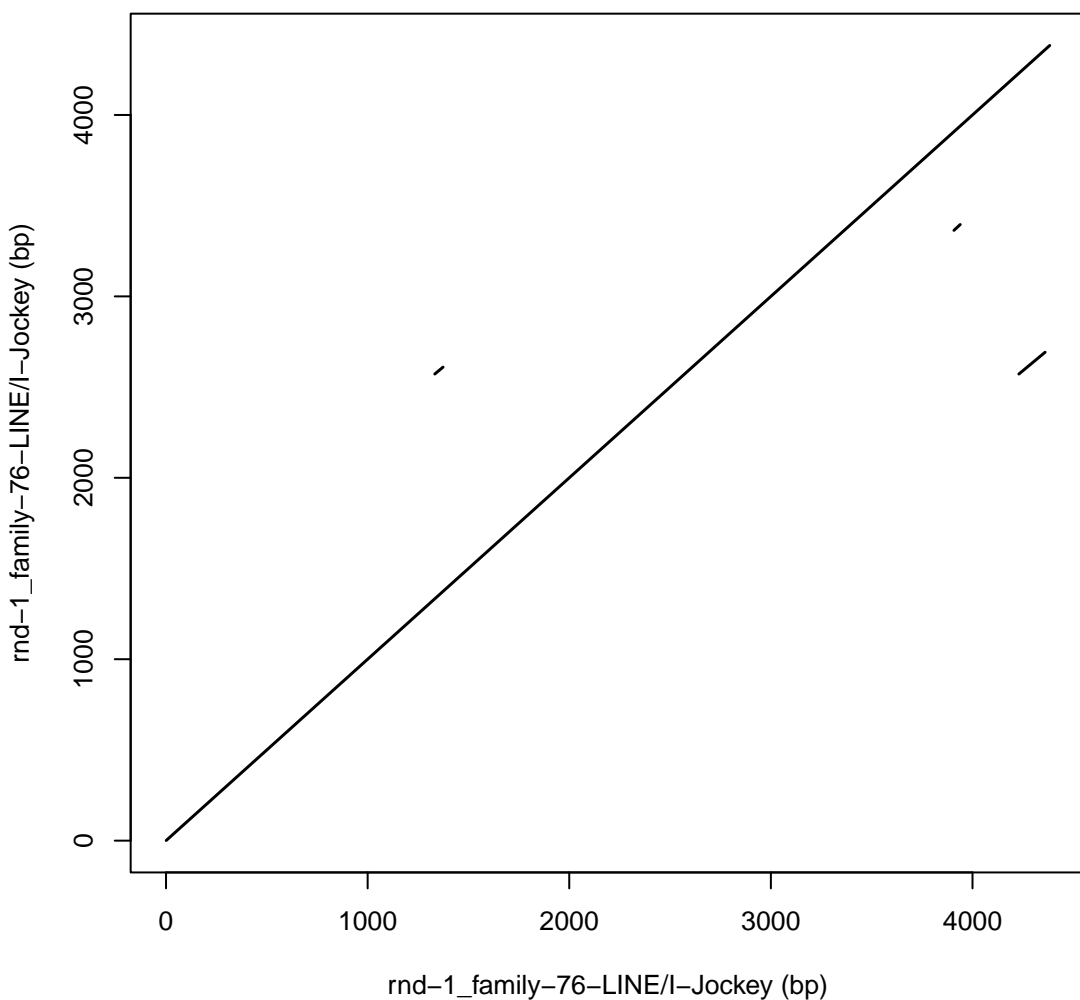
TE: rnd-1_family-76-LINE/I-Jockey
consensus size: 4383bp; fragments: 371; full length: 2 (≥ 3944.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

