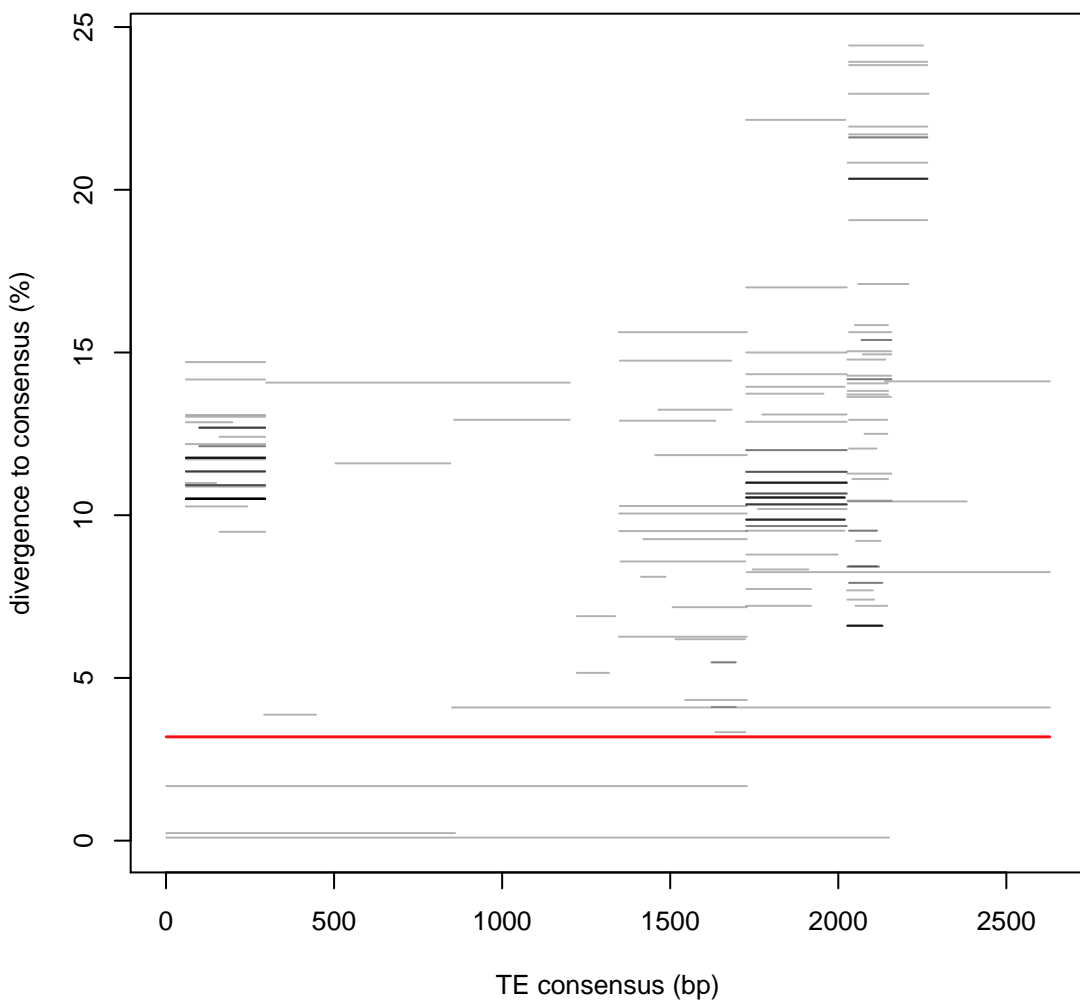
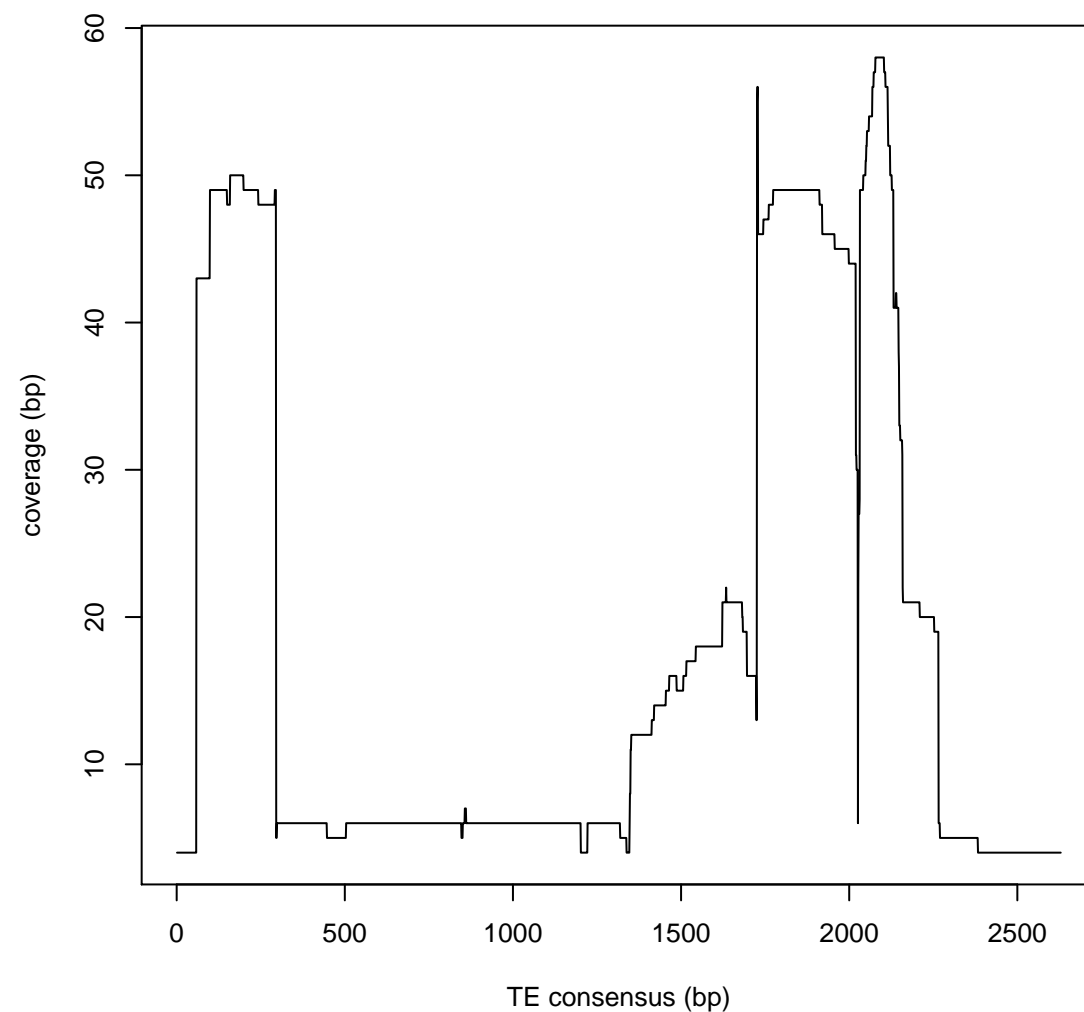


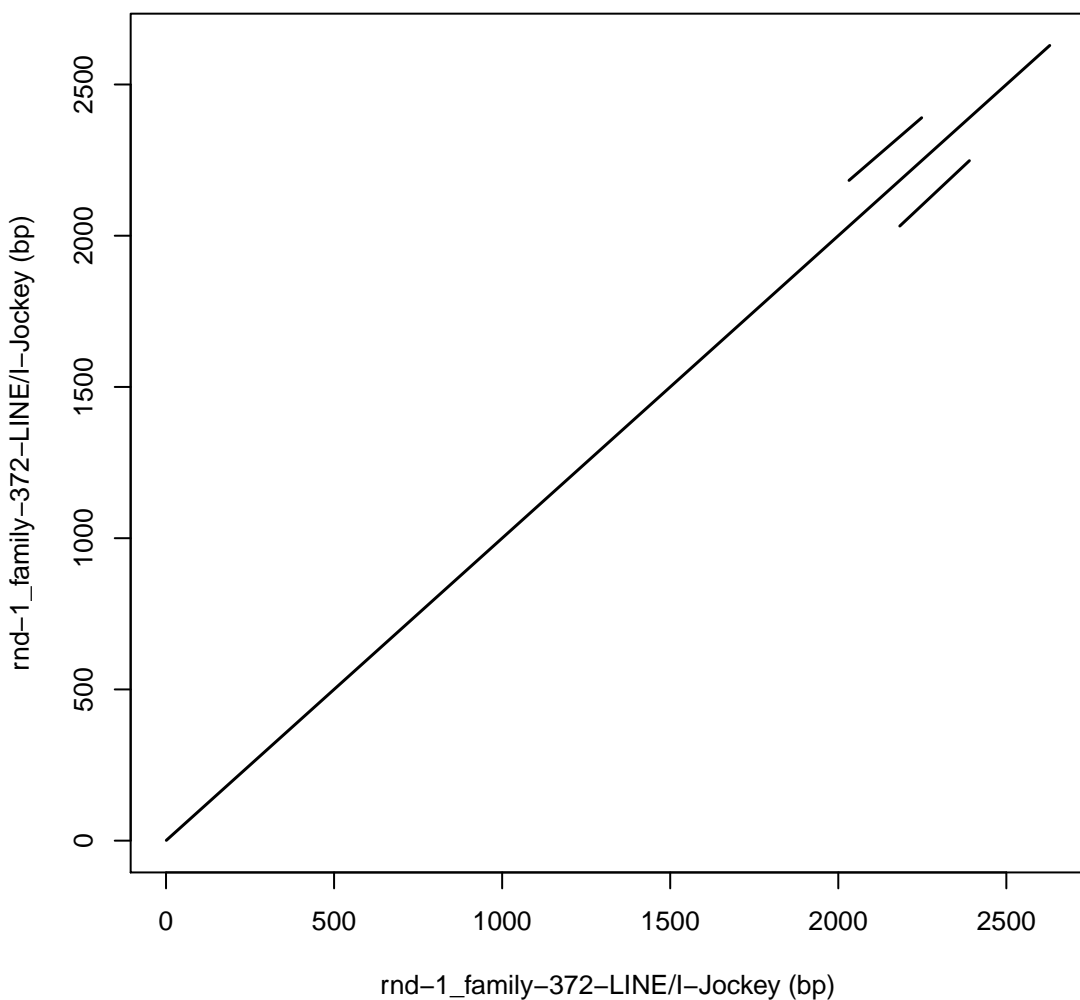
TE: rnd-1_family-372-LINE/I-Jockey
consensus size: 2629bp; fragments: 178; full length: 1 (>=2366.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

