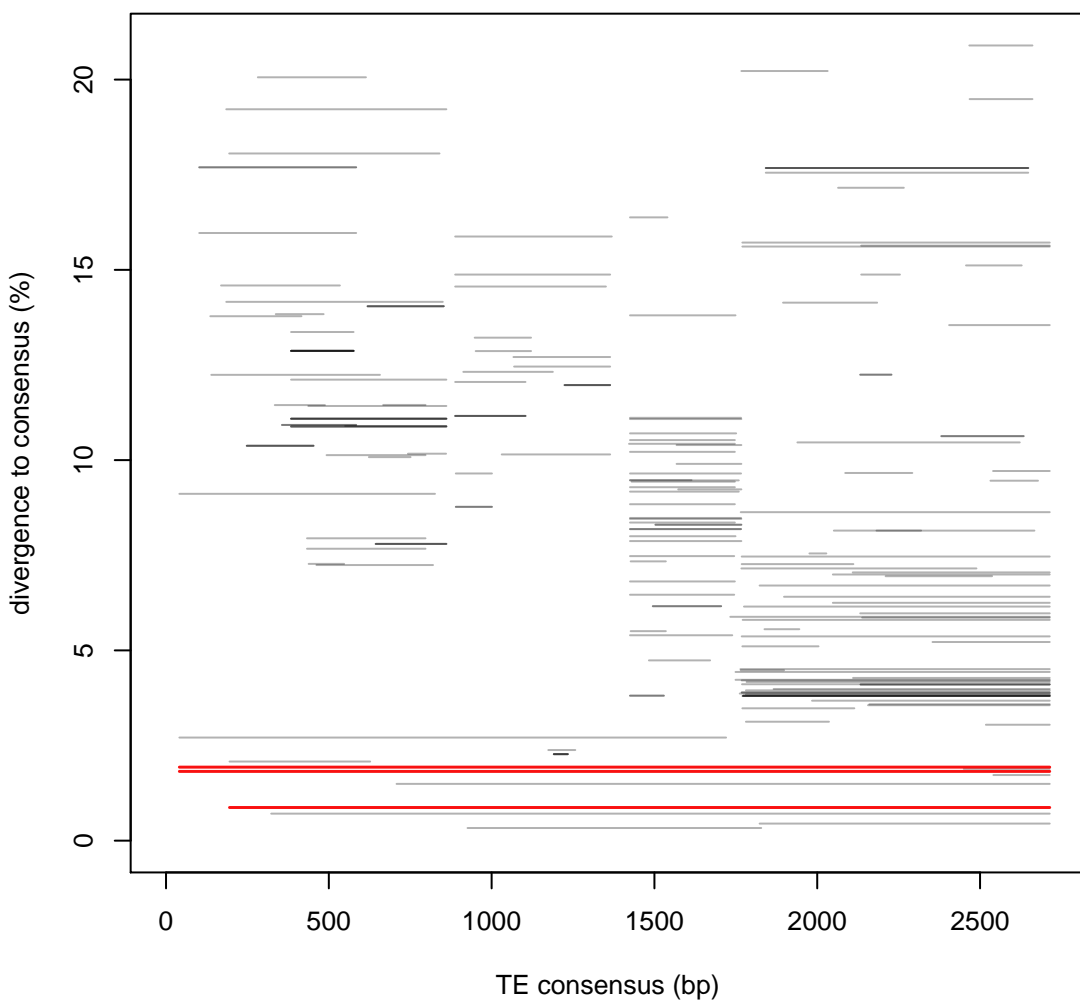
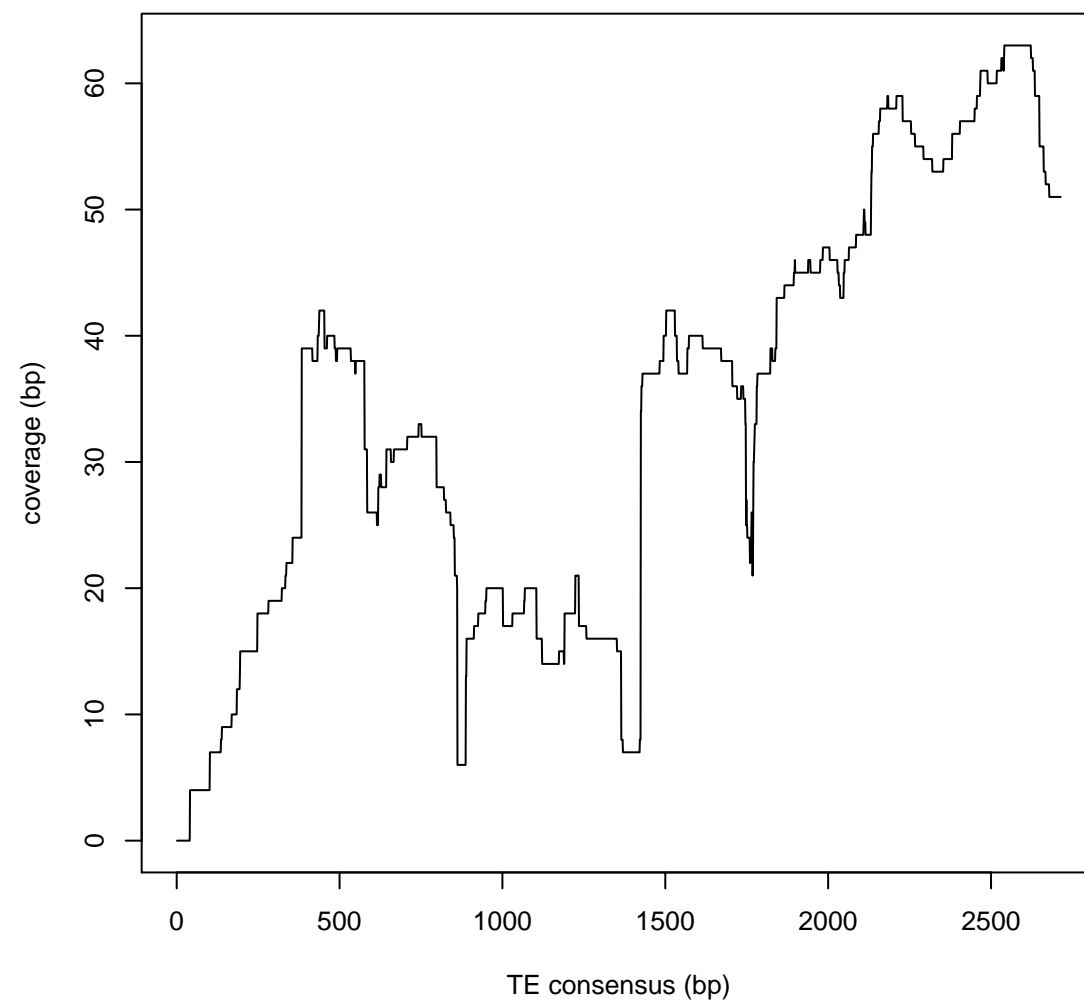


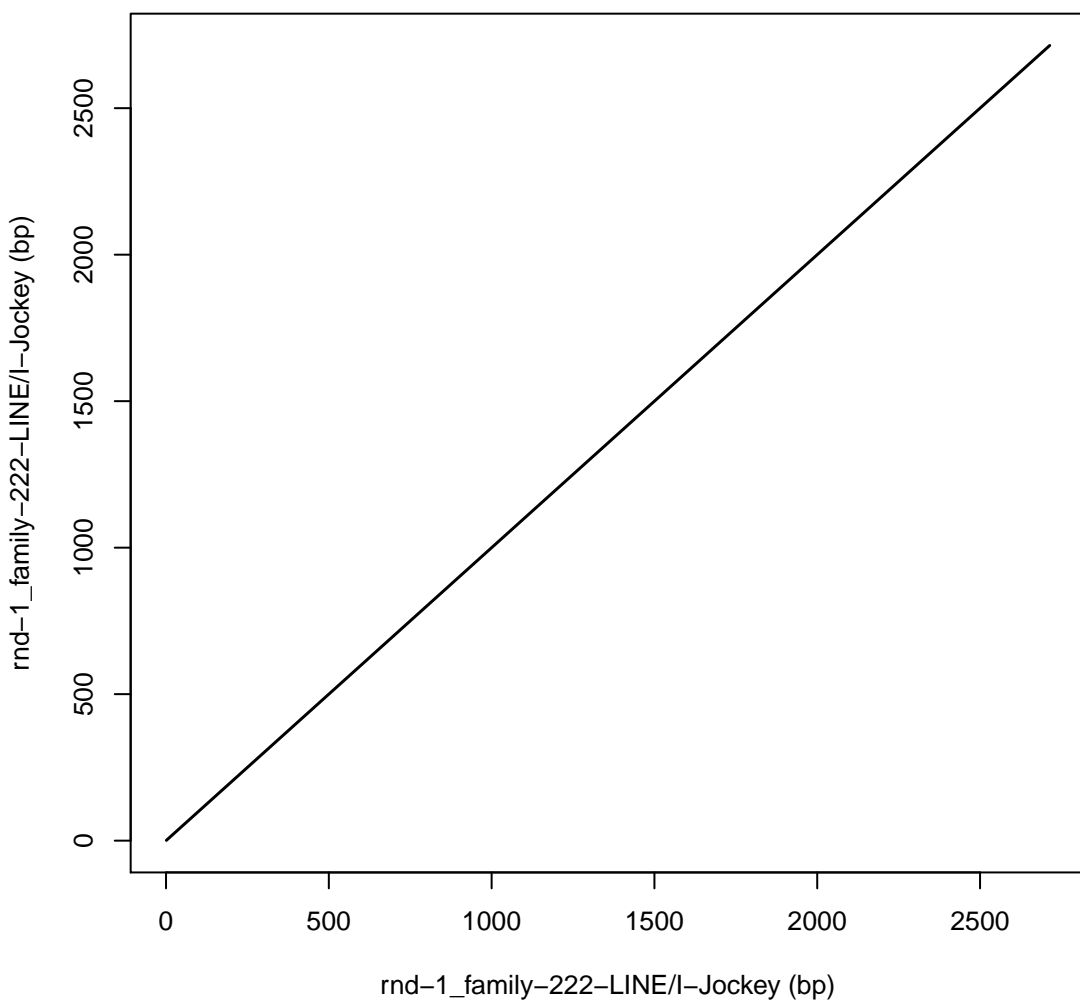
TE: rnd-1_family-222-LINE/I-Jockey
consensus size: 2714bp; fragments: 194; full length: 3 (>=2442.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

