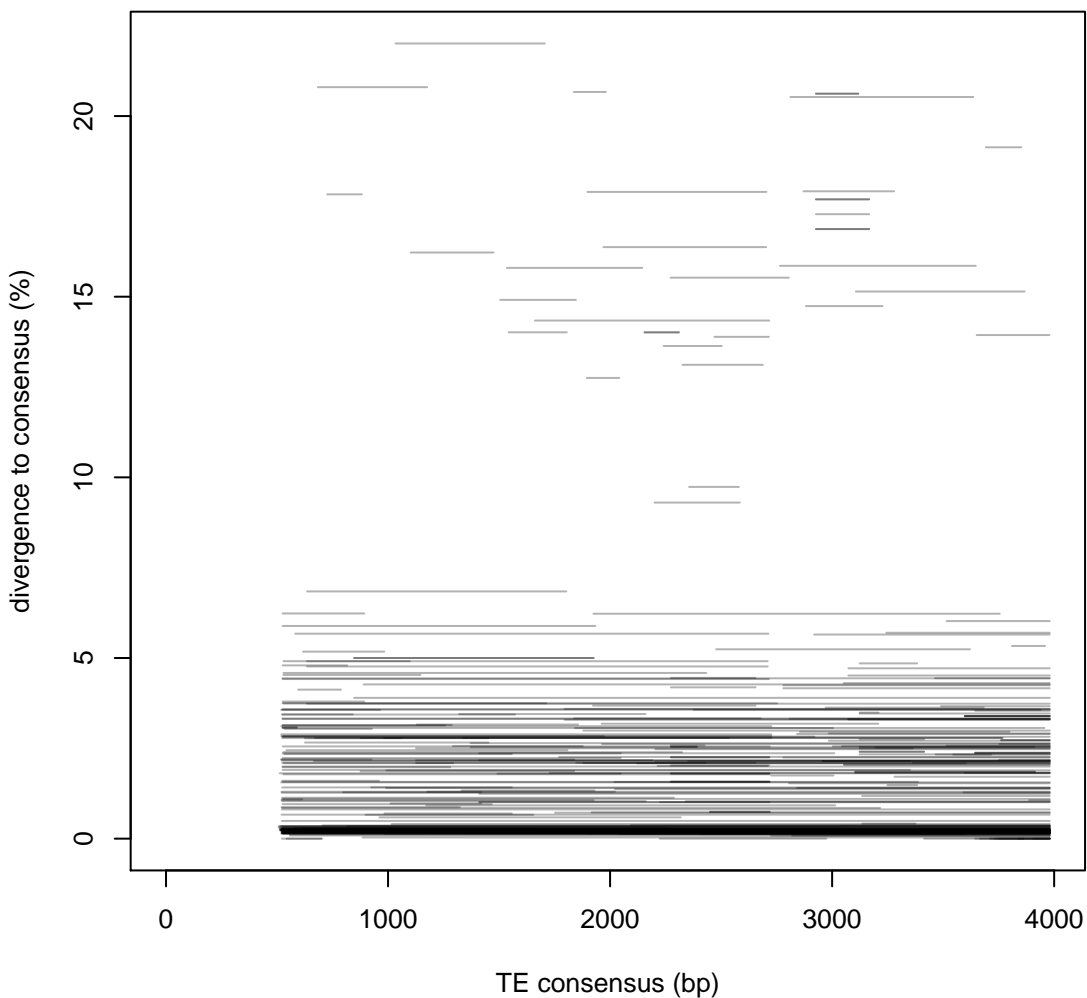
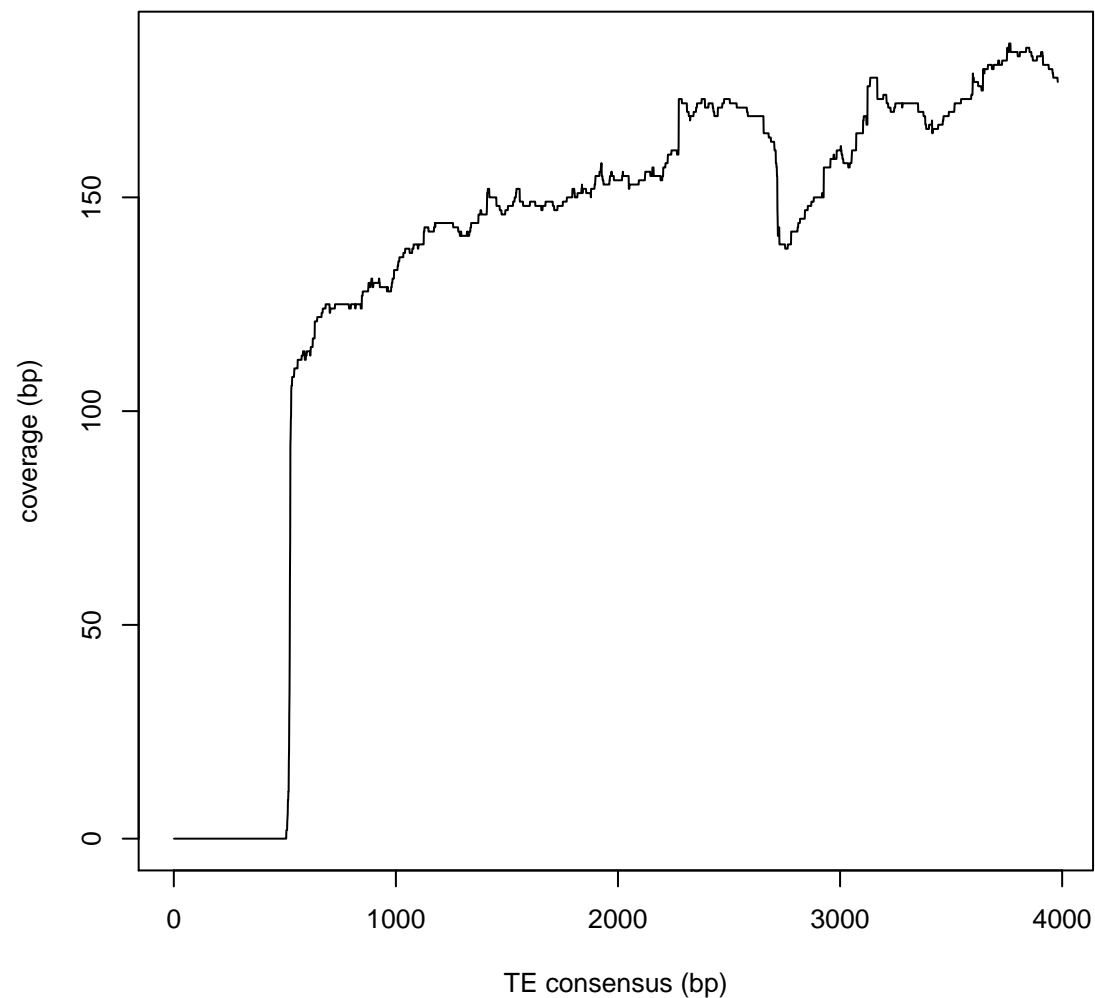


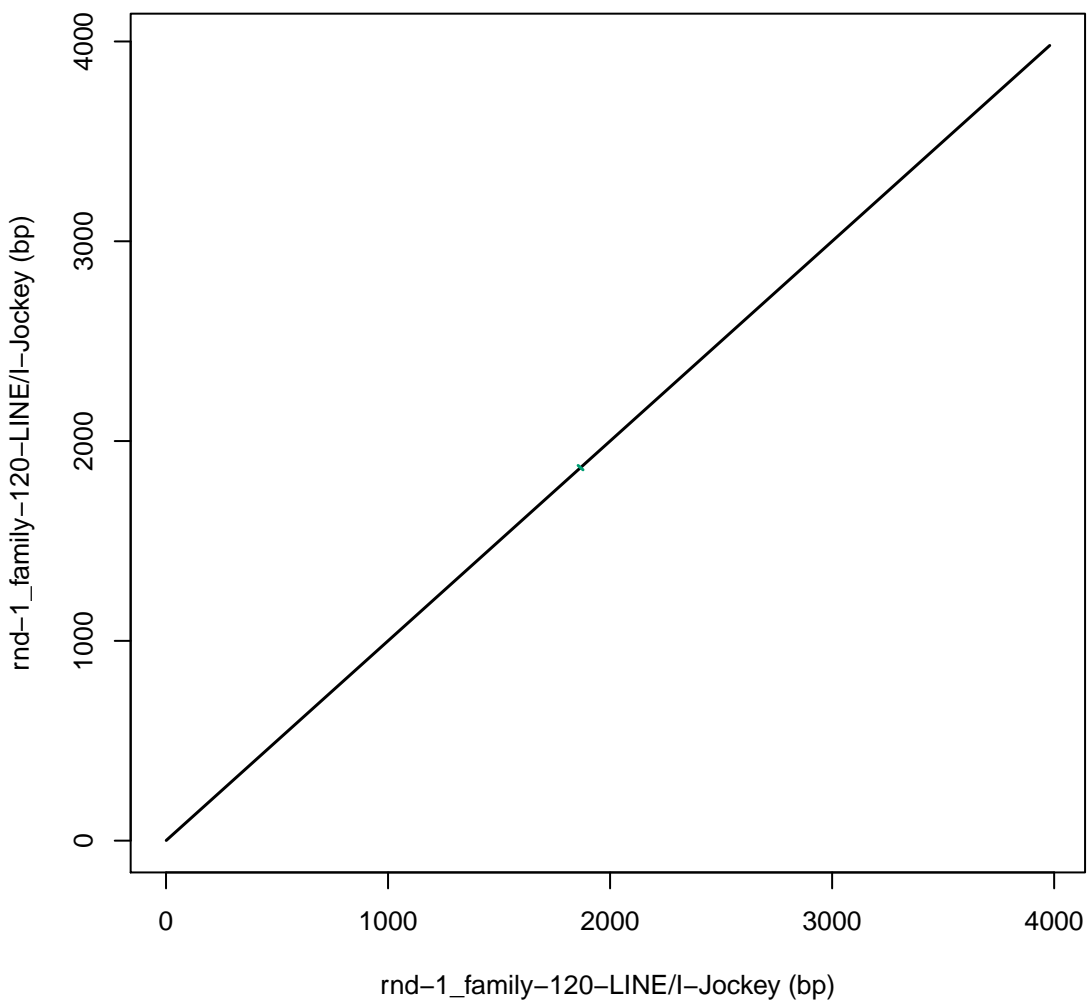
TE: rnd-1_family-120-LINE/I-Jockey
consensus size: 3980bp; fragments: 340; full length: 0 (>=3582bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

