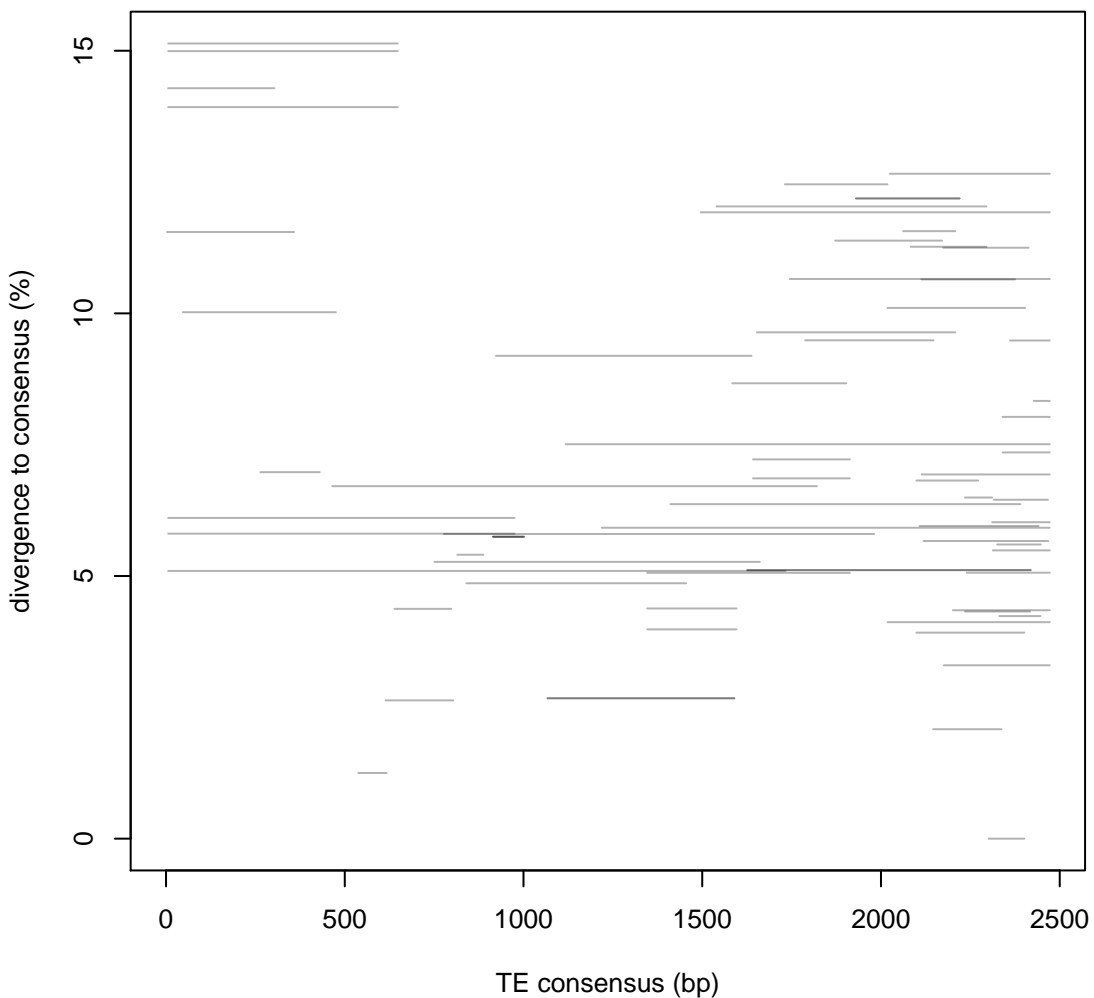
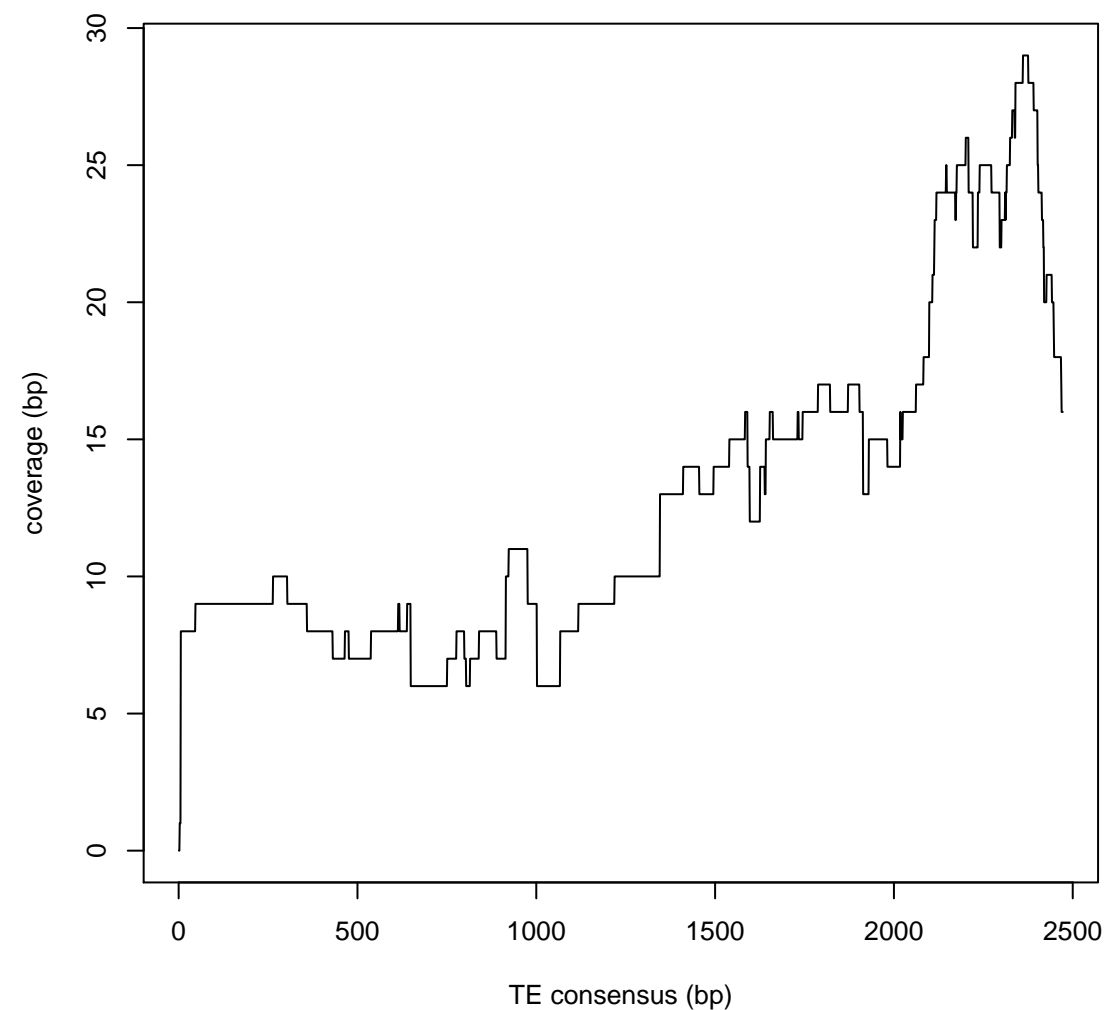


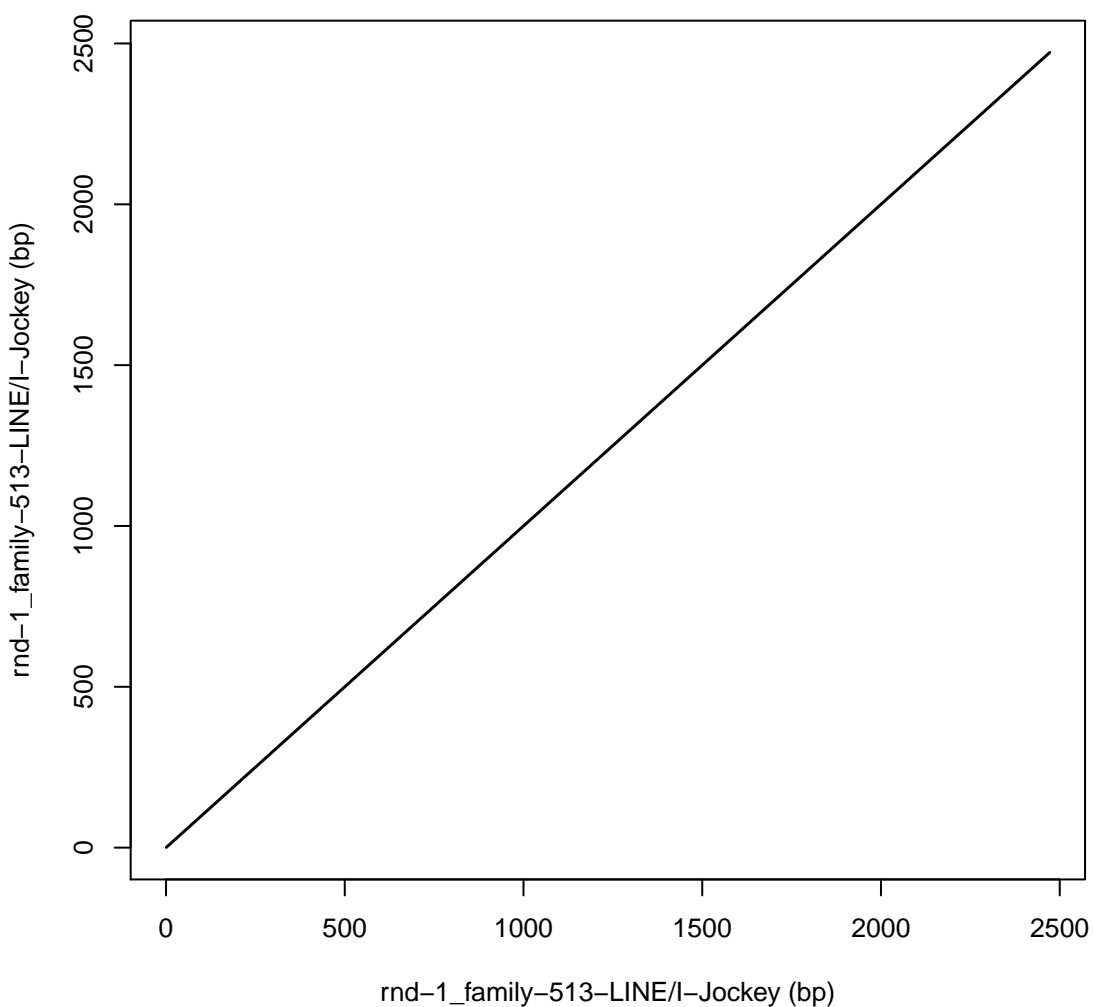
TE: rnd-1_family-513-LINE/I-Jockey
consensus size: 2472bp; fragments: 72; full length: 0 (>=2224.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

