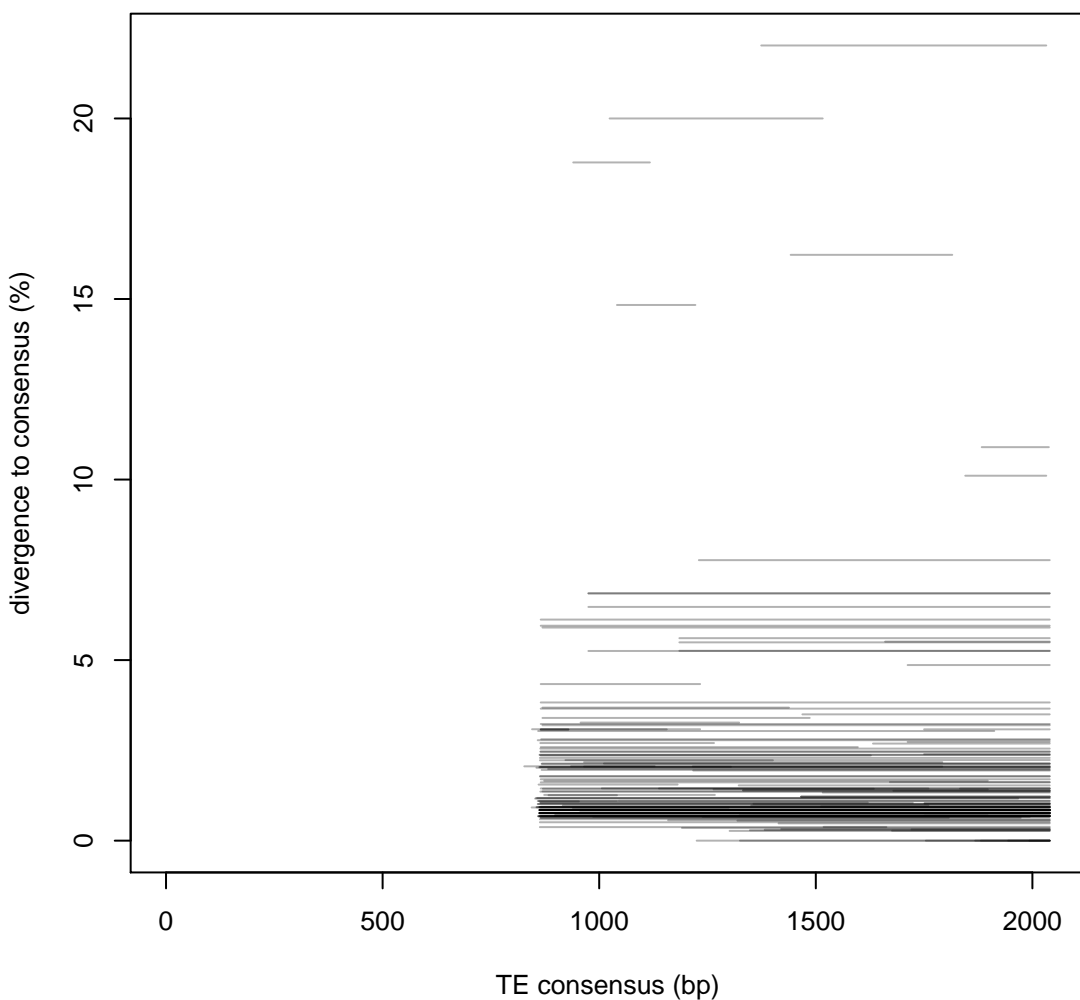
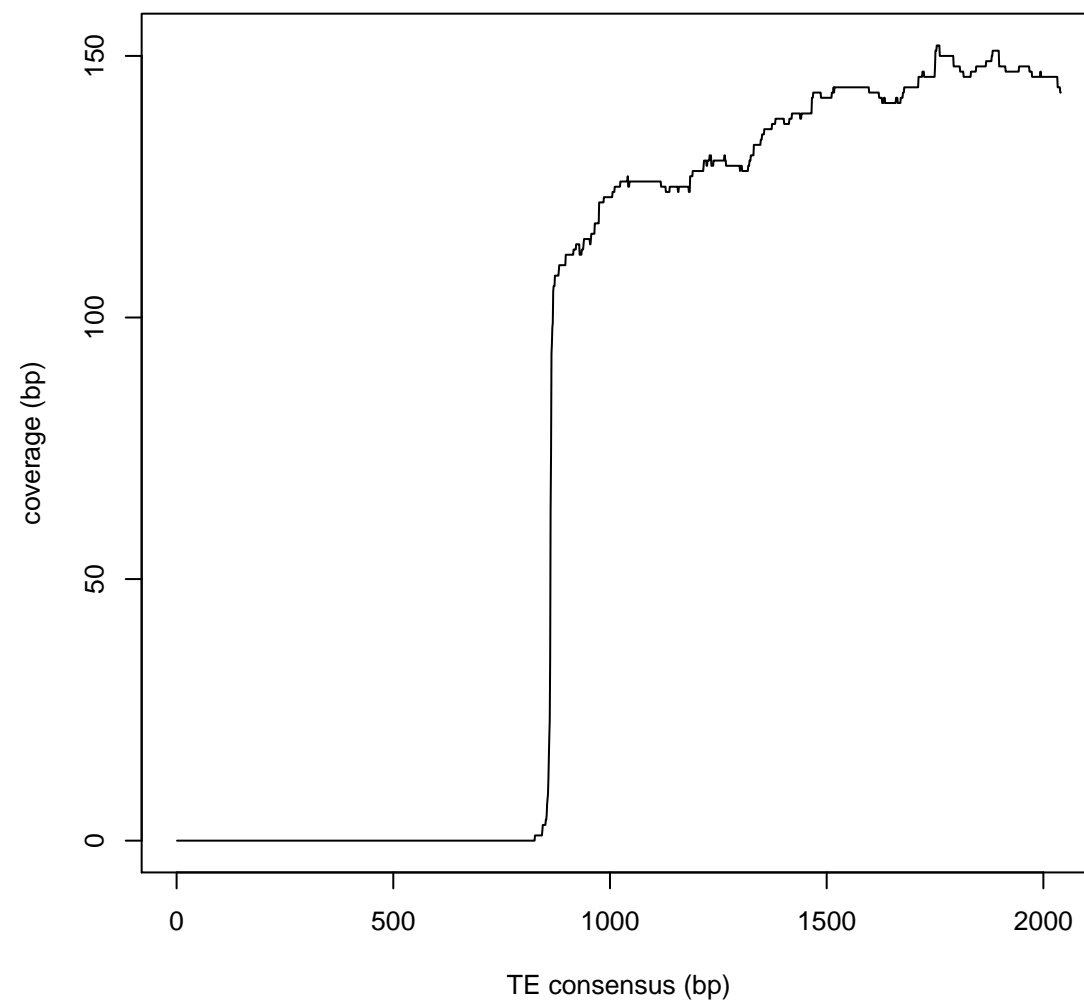


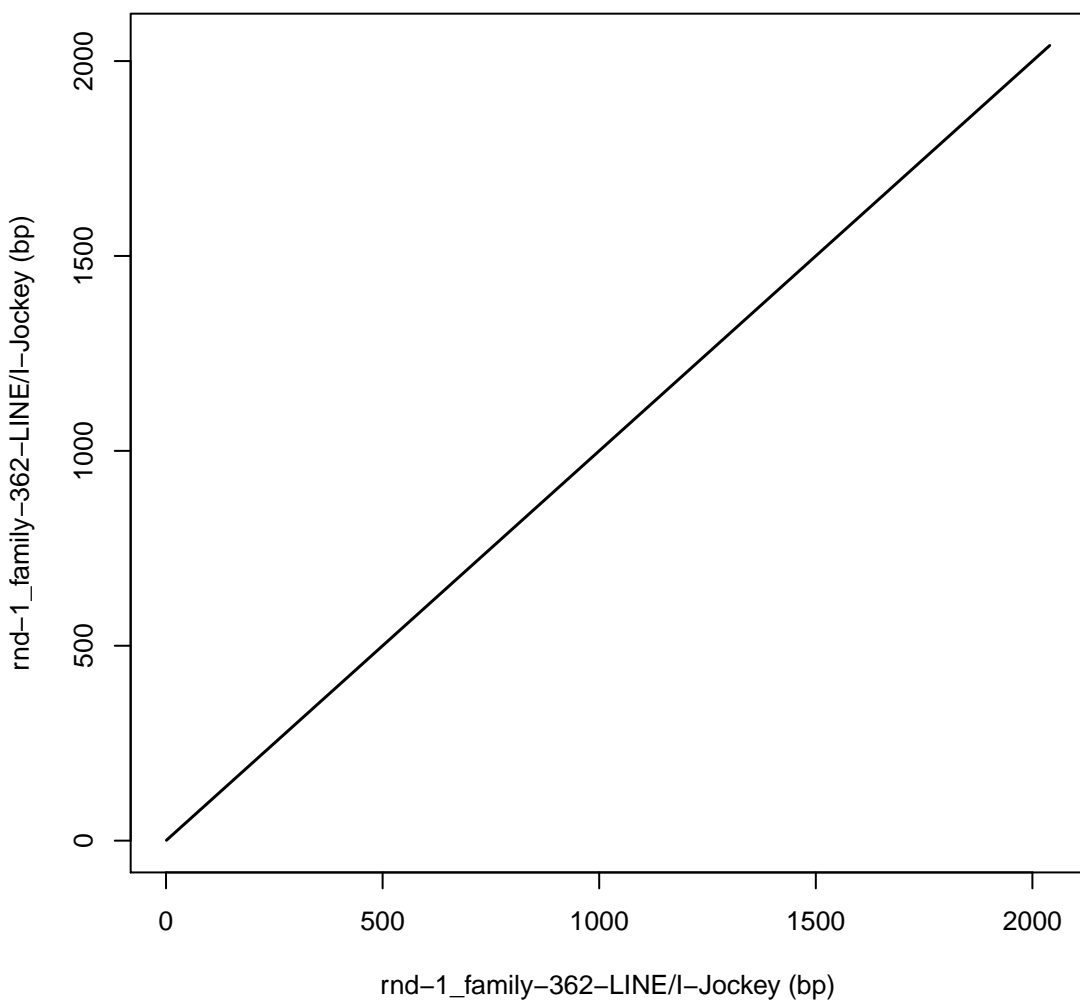
TE: rnd-1_family-362-LINE/I-Jockey
consensus size: 2040bp; fragments: 186; full length: 0 (>=1836bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

