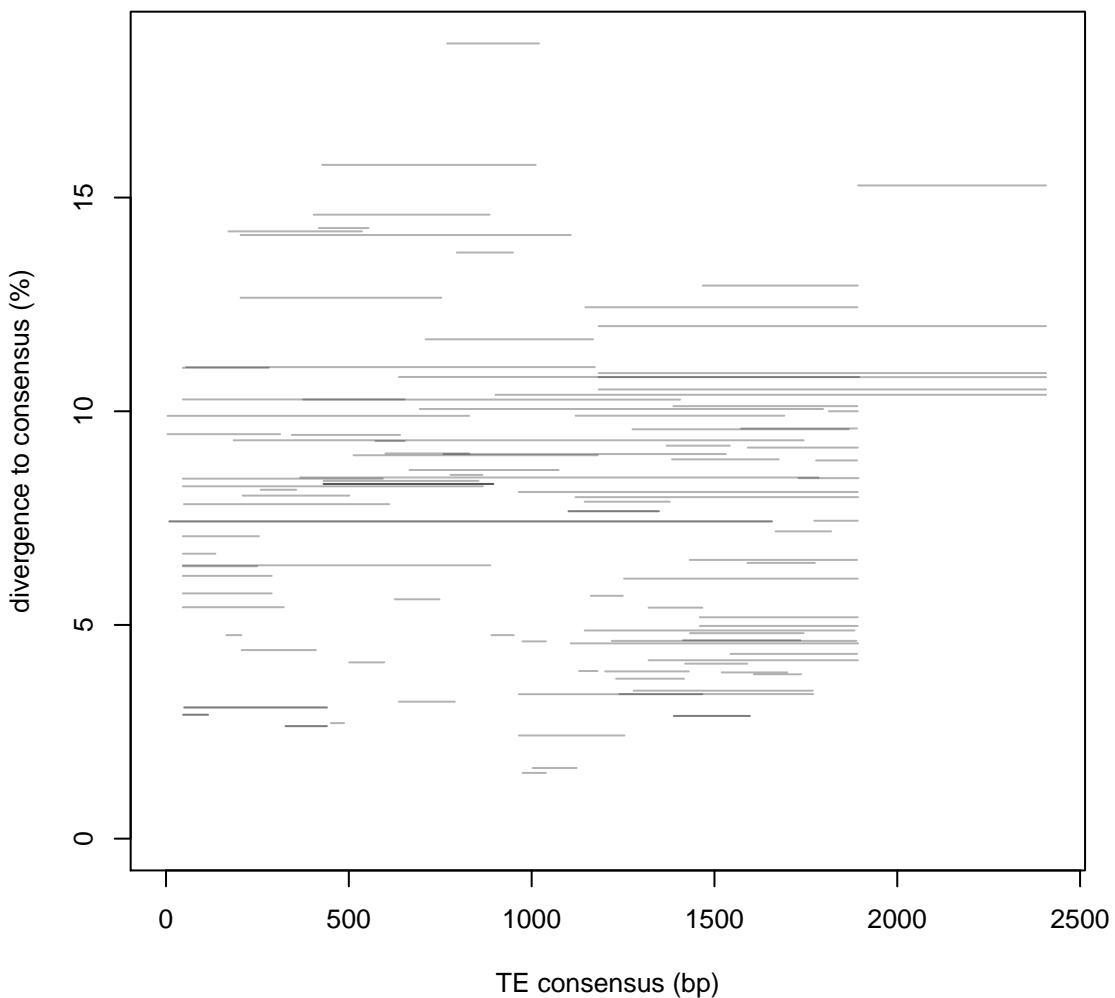
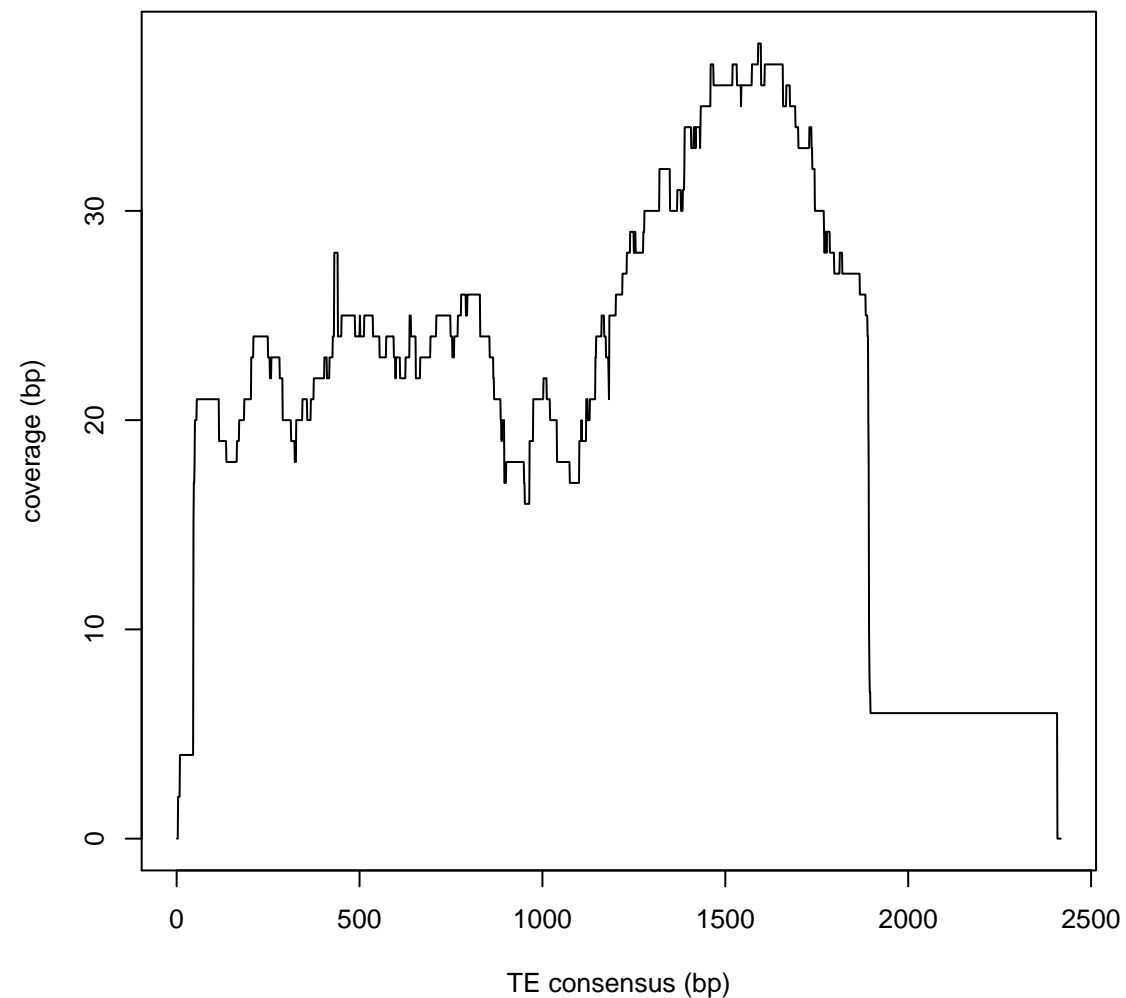


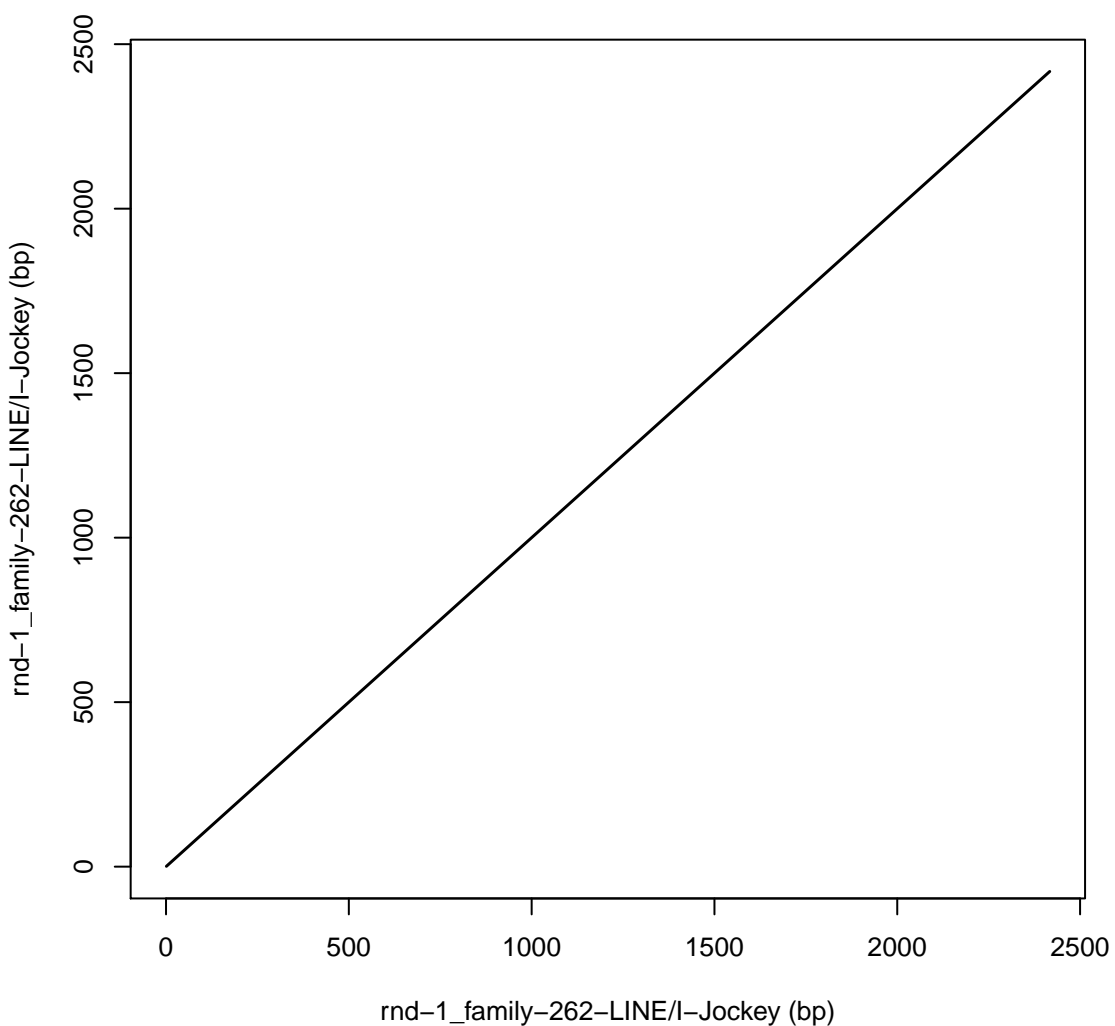
TE: rnd-1_family-262-LINE/I-Jockey
consensus size: 2417bp; fragments: 111; full length: 0 (>=2175.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

