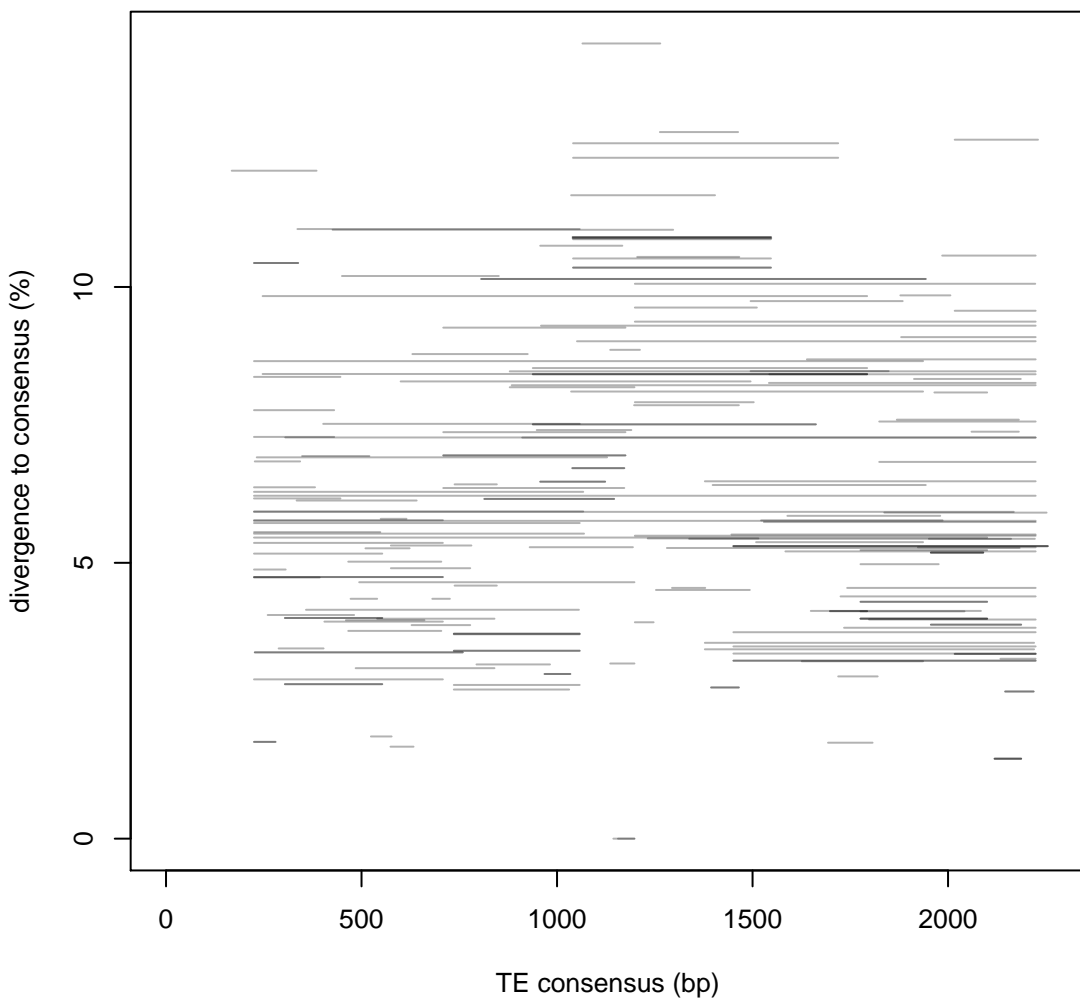
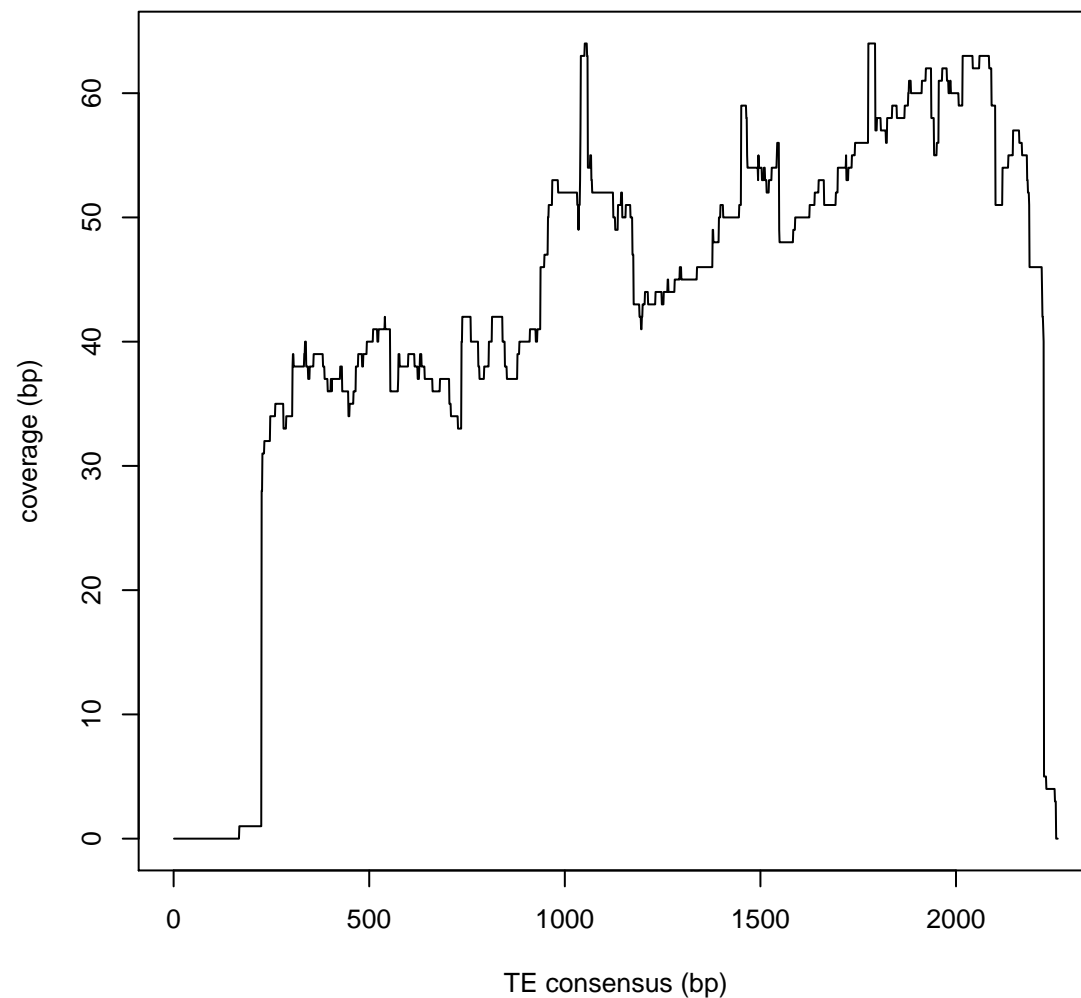


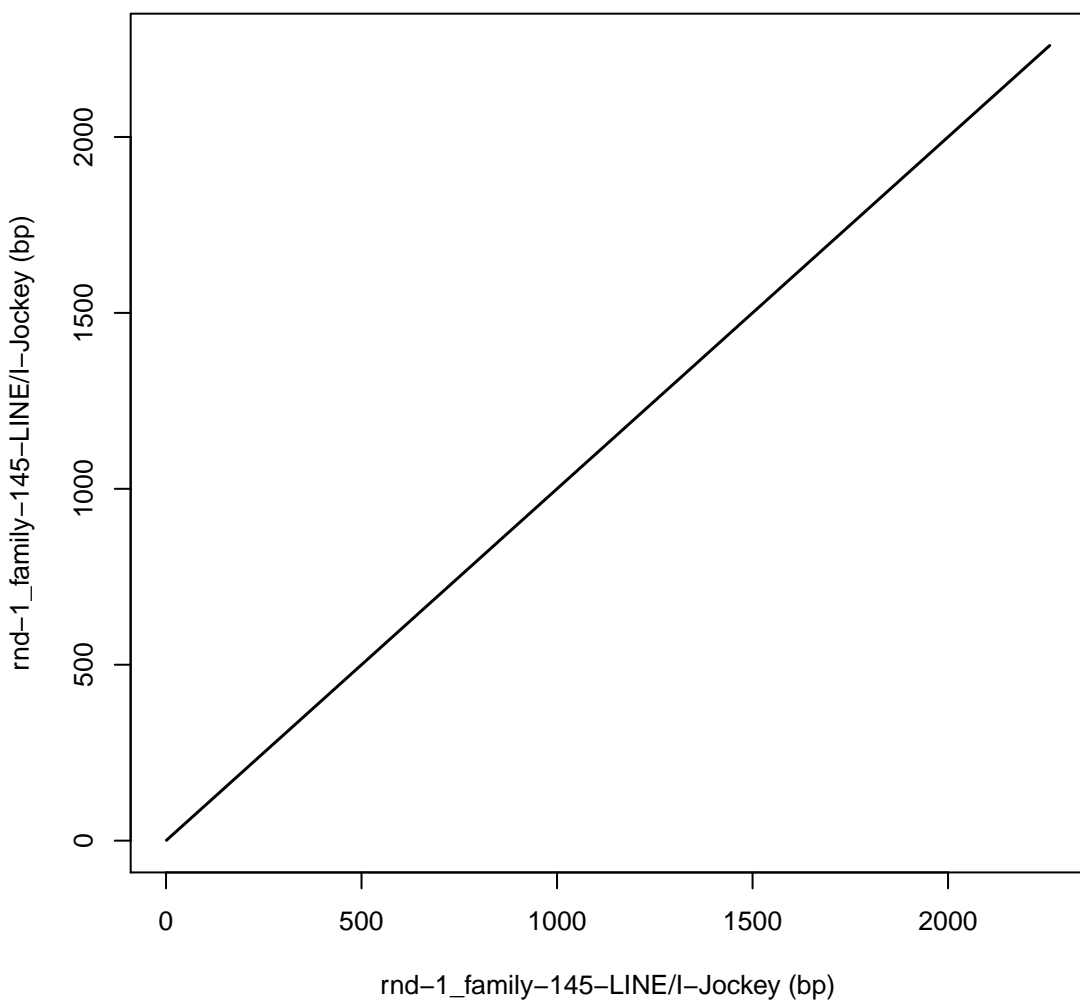
TE: rnd-1_family-145-LINE/I-Jockey
consensus size: 2260bp; fragments: 206; full length: 0 (>=2034bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

