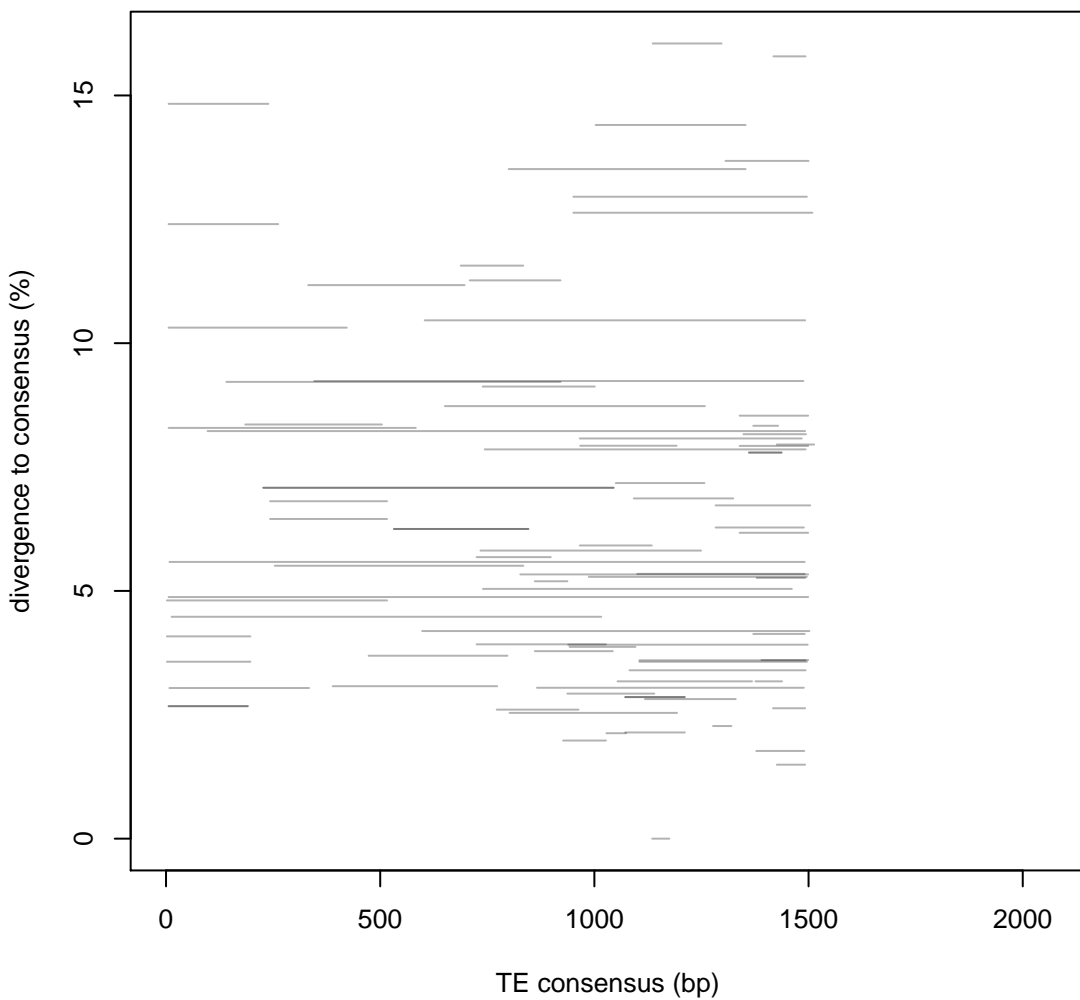
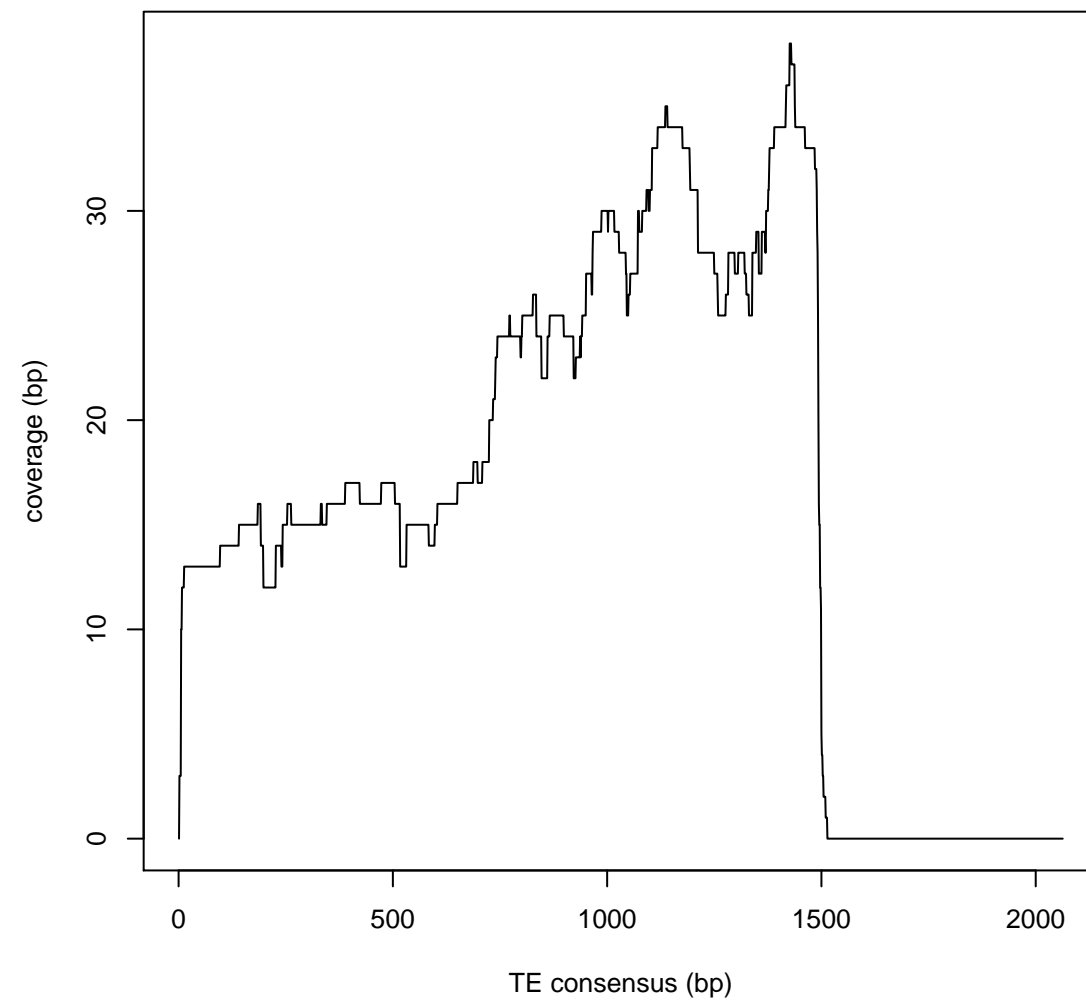


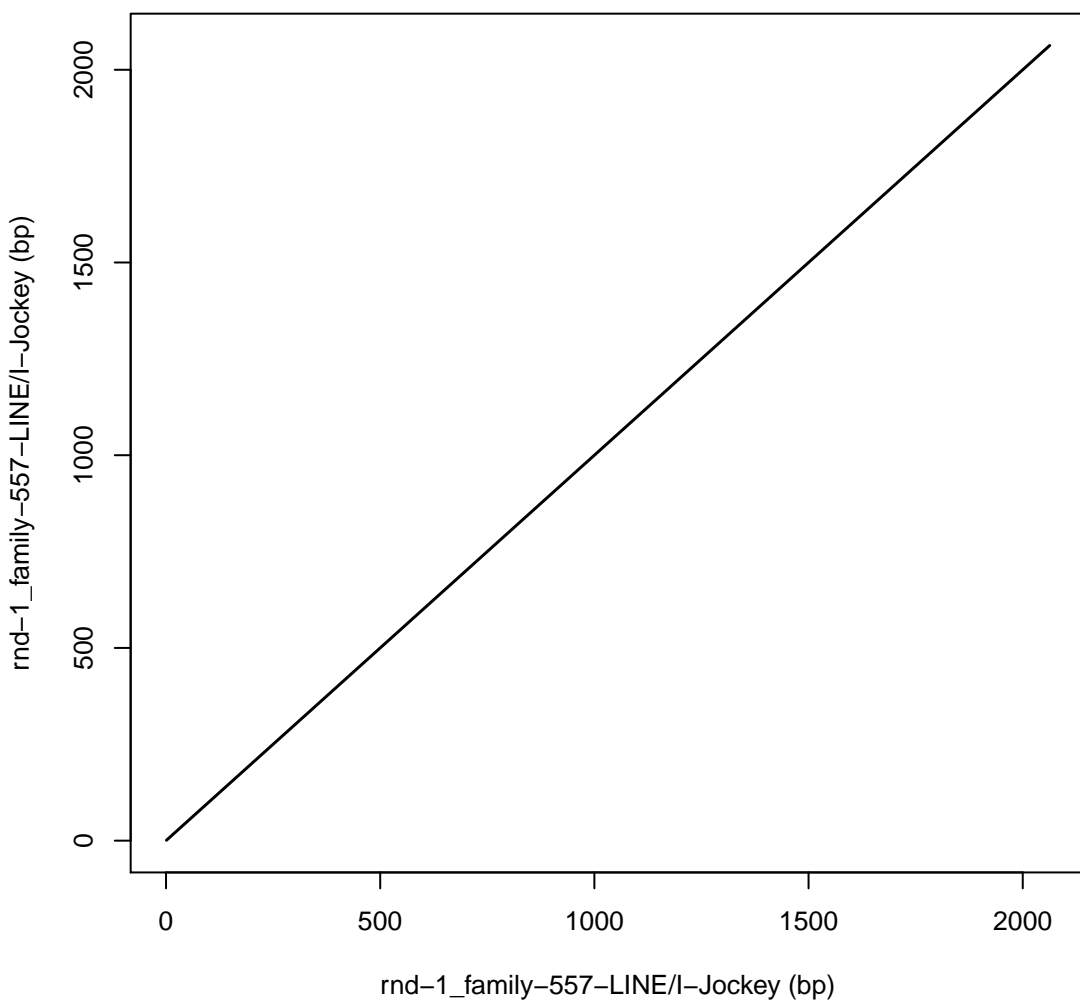
TE: rnd-1_family-557-LINE/I-Jockey
consensus size: 2063bp; fragments: 90; full length: 0 (≥ 1856.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

