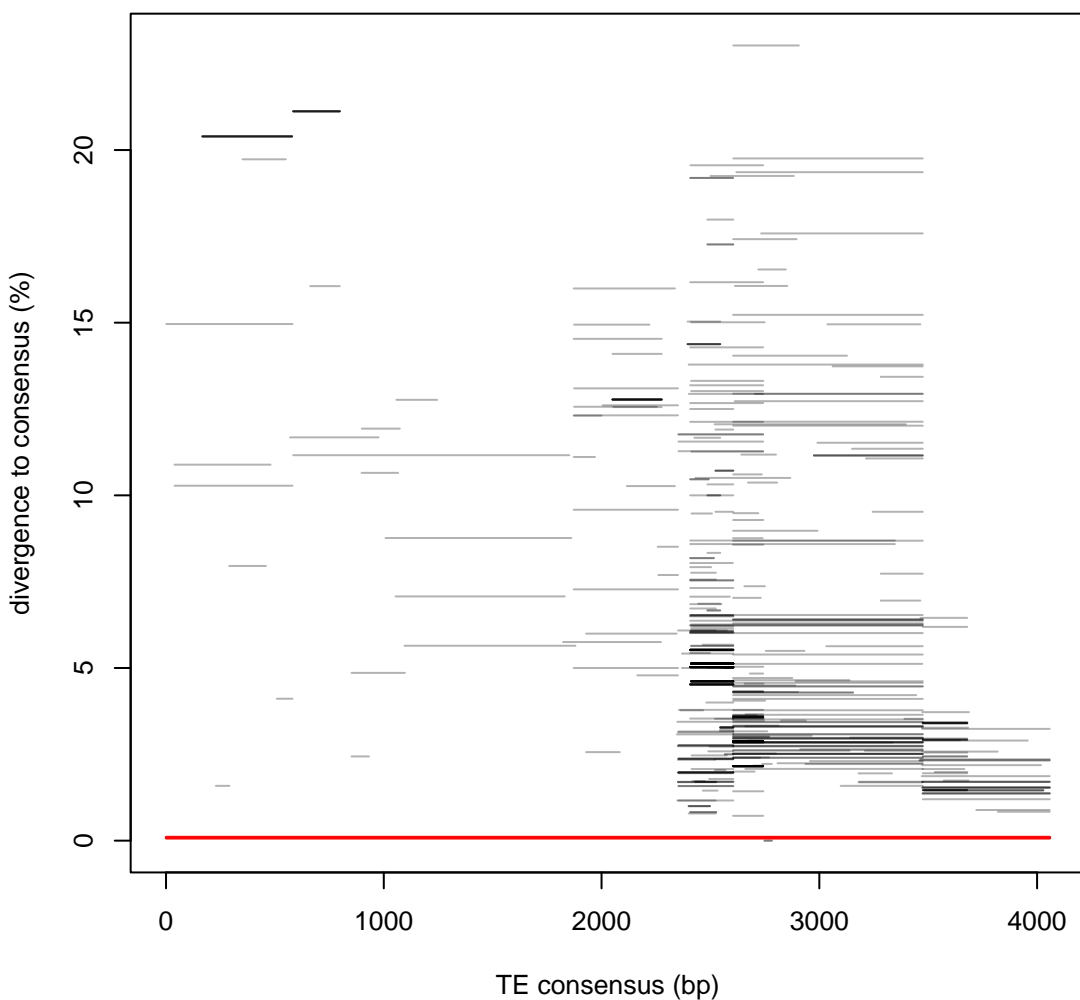
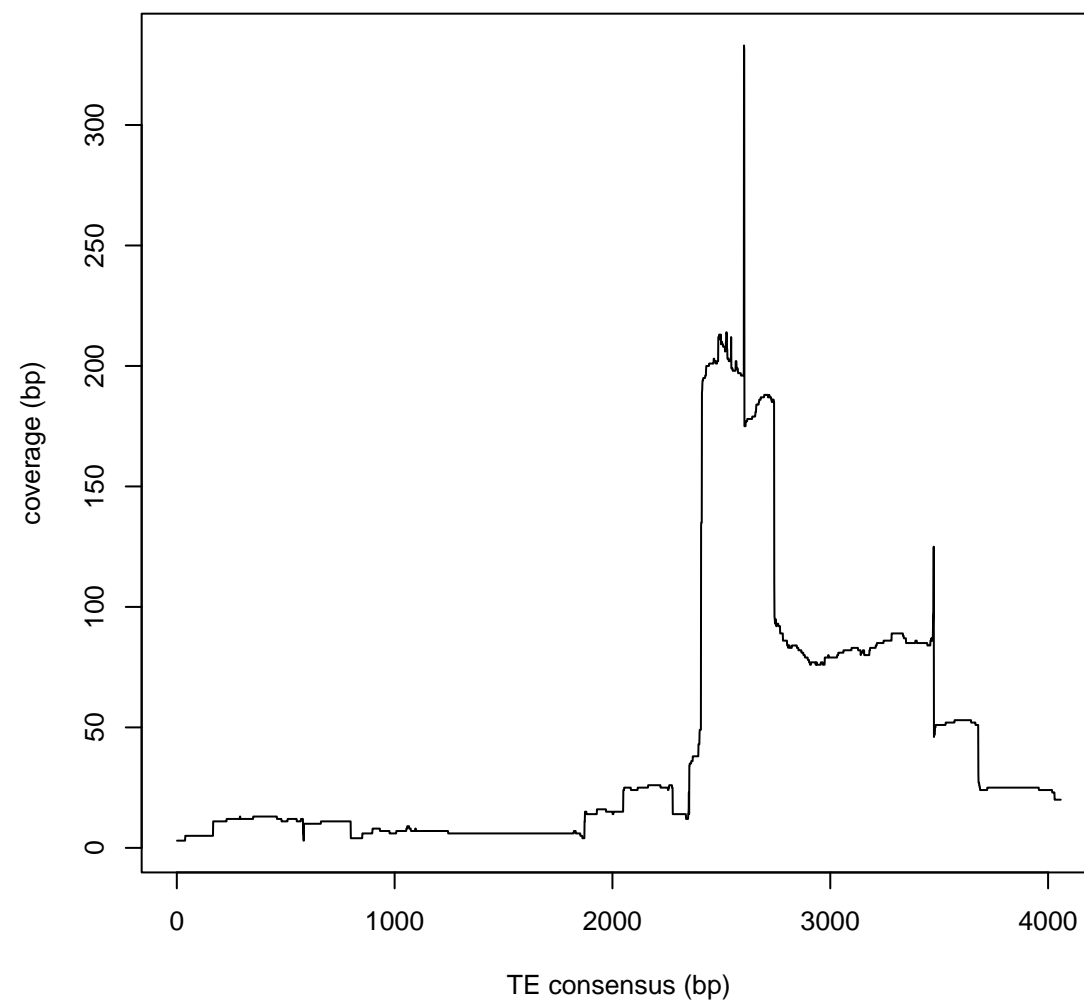


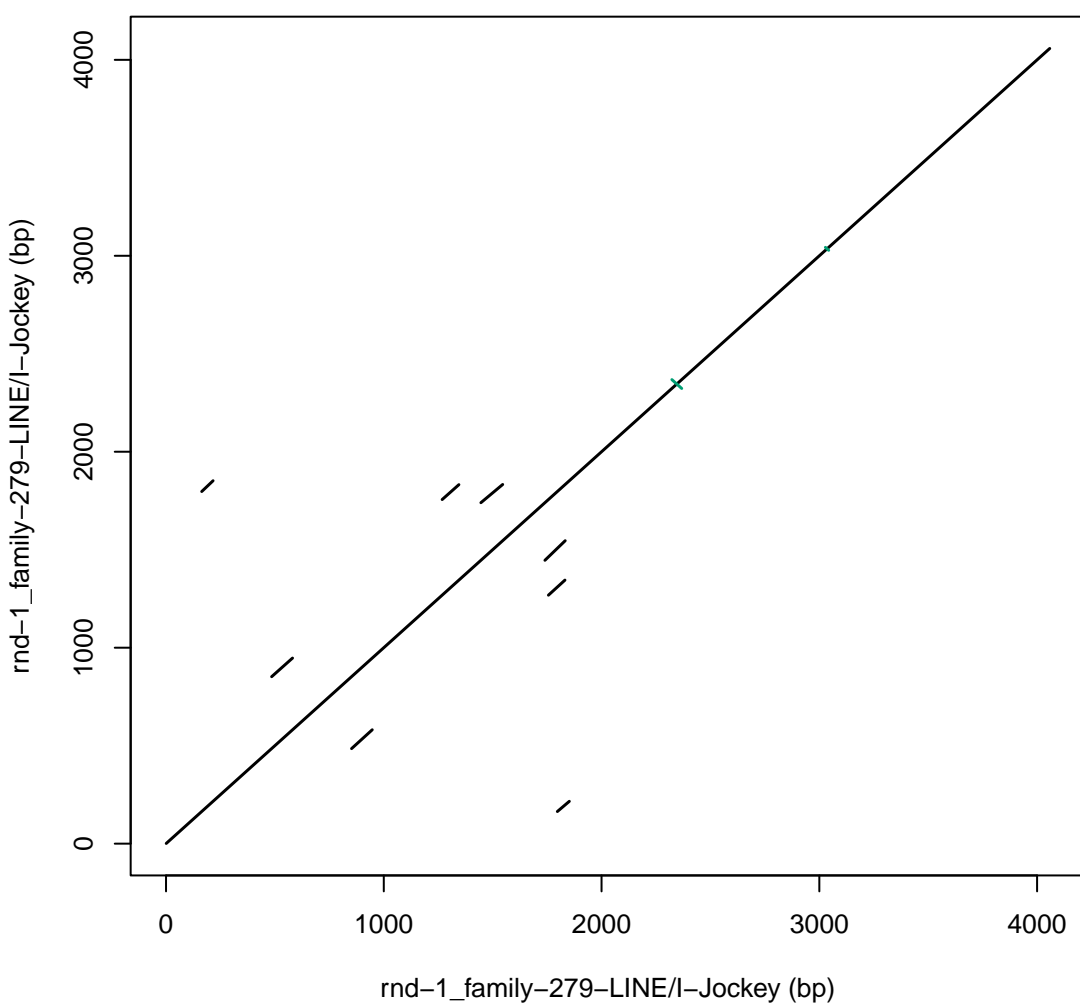
TE: rnd-1_family-279-LINE/I-Jockey
consensus size: 4058bp; fragments: 543; full length: 2 (≥ 3652.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

