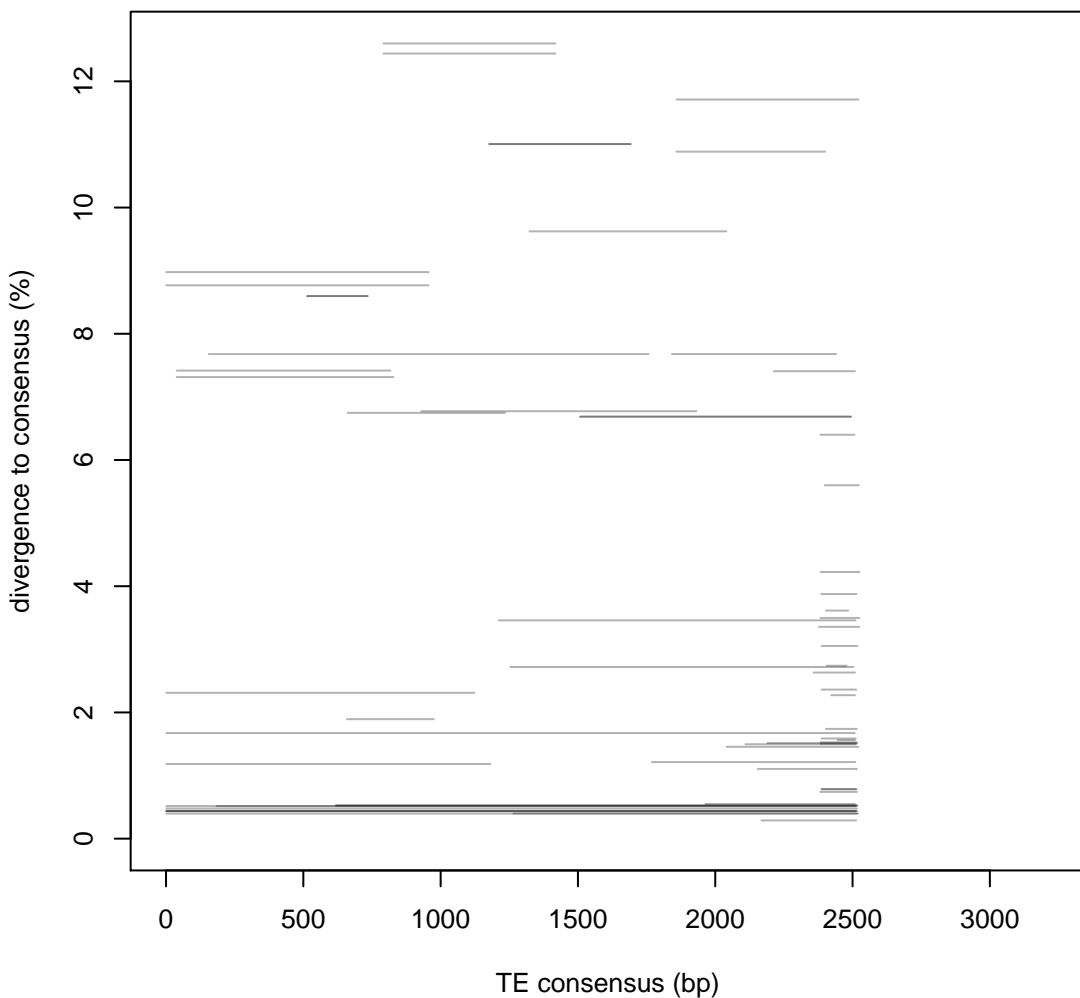
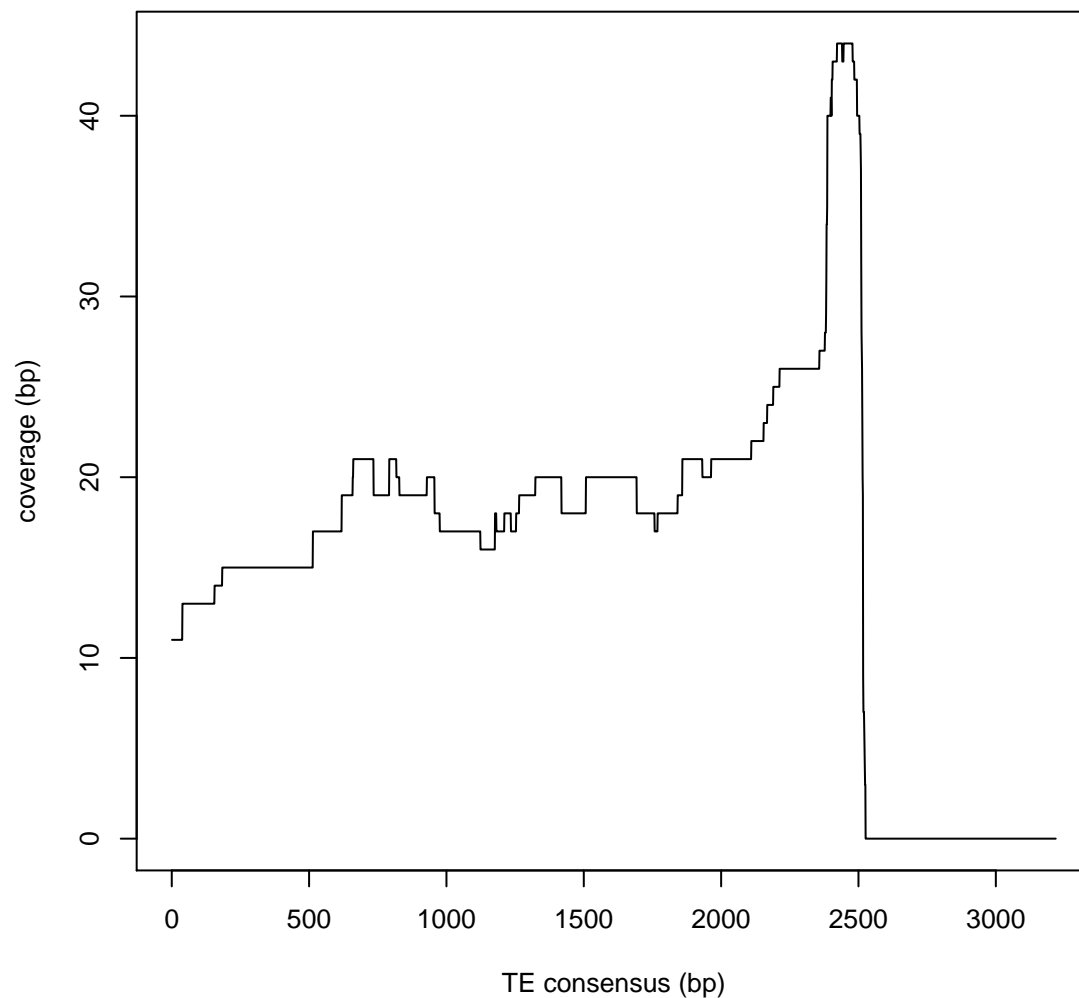


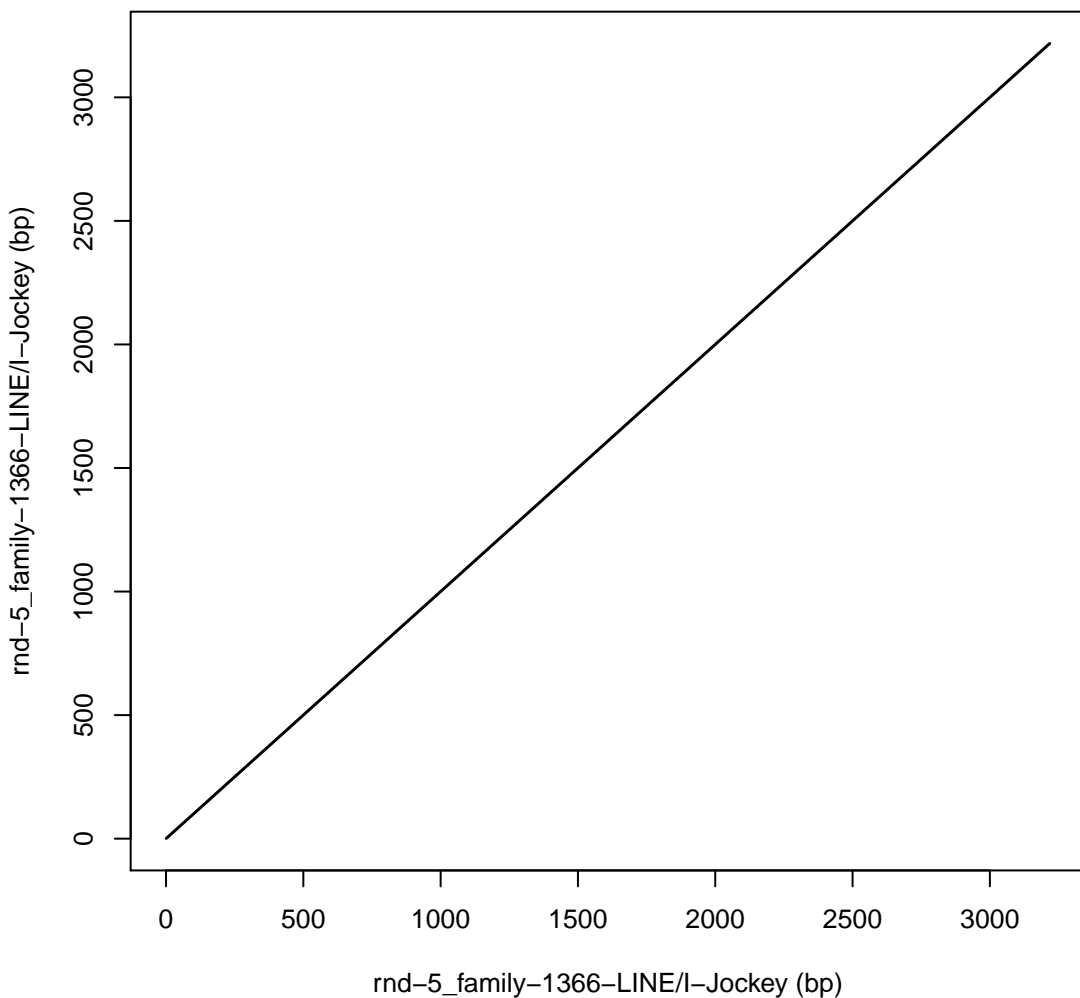
TE: rnd-5_family-1366-LINE/I-Jockey
consensus size: 3218bp; fragments: 63; full length: 0 (≥ 2896.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

