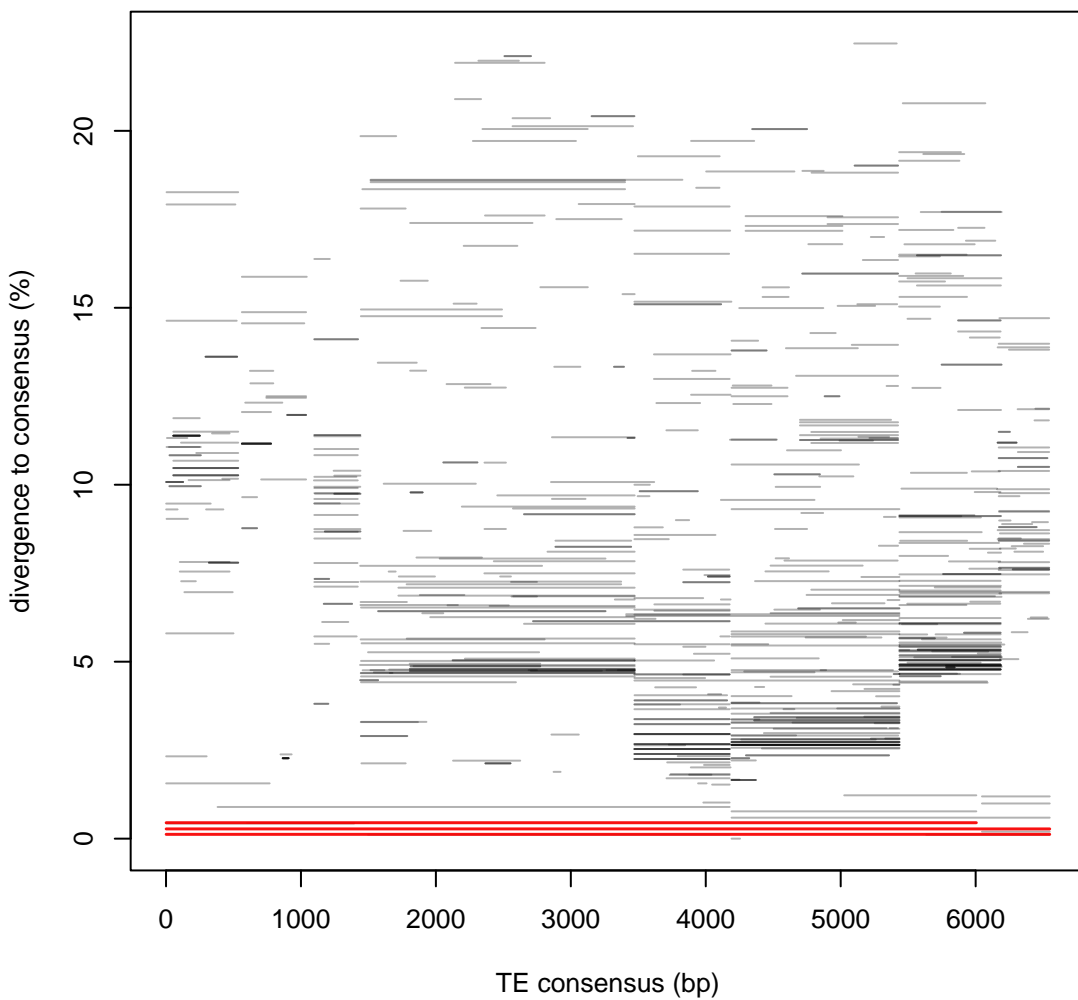
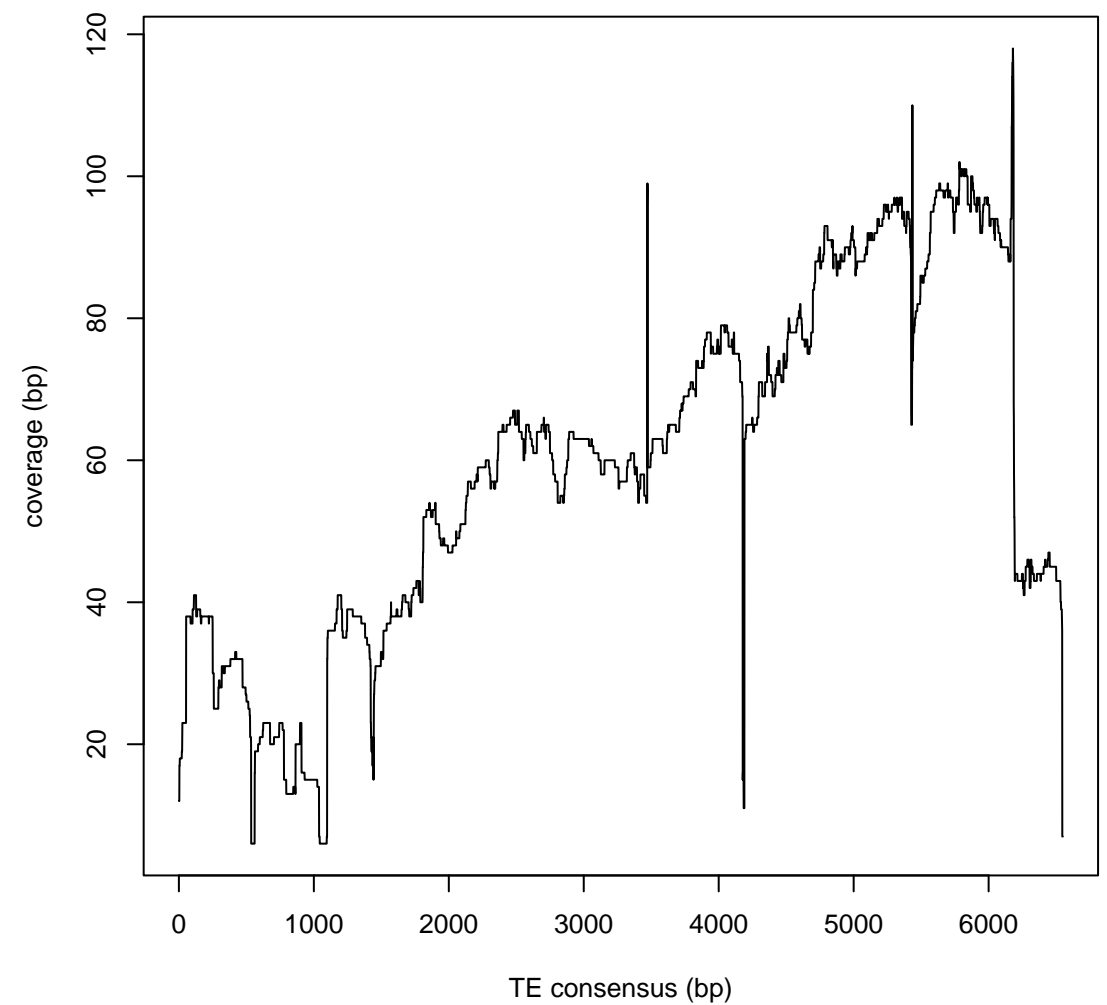


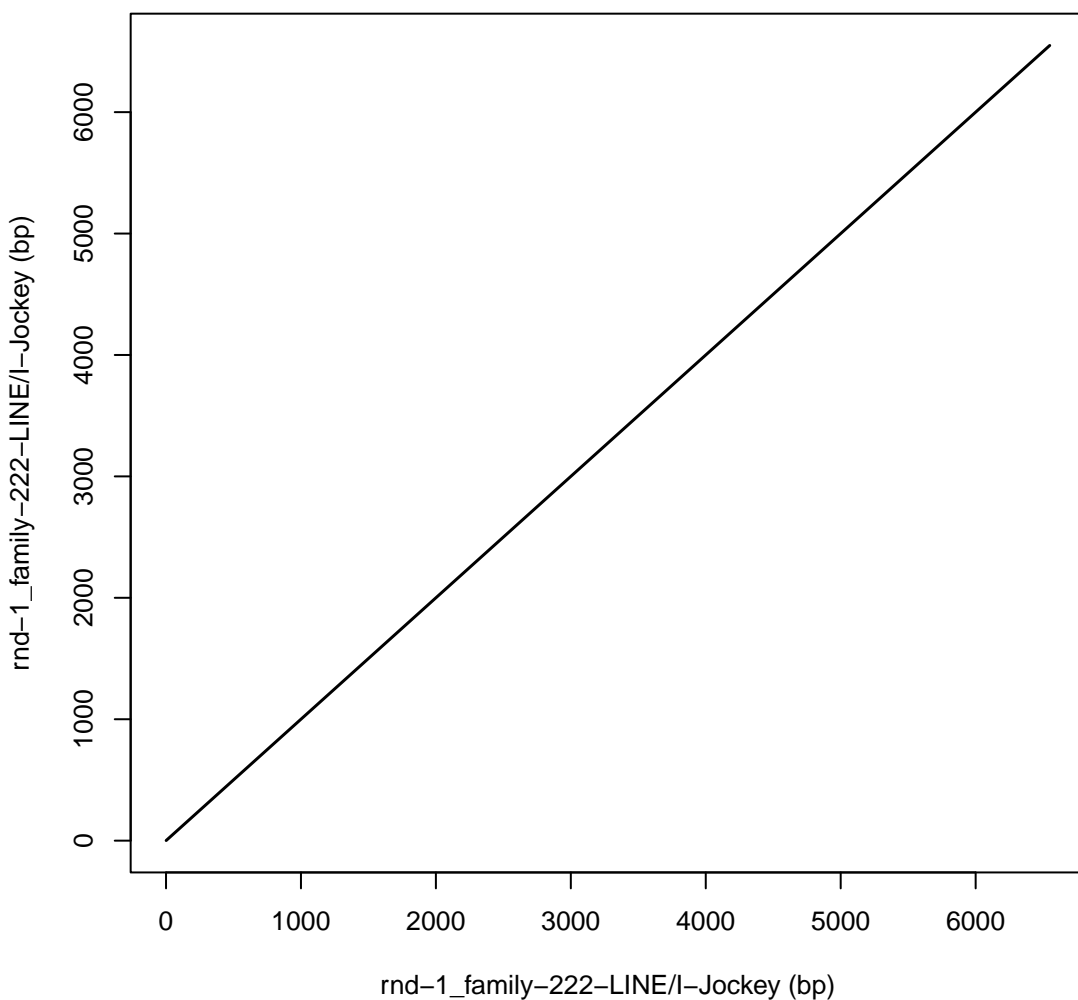
TE: rnd-1_family-222-LINE/I-Jockey
consensus size: 6549bp; fragments: 687; full length: 3 (≥ 5894.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

