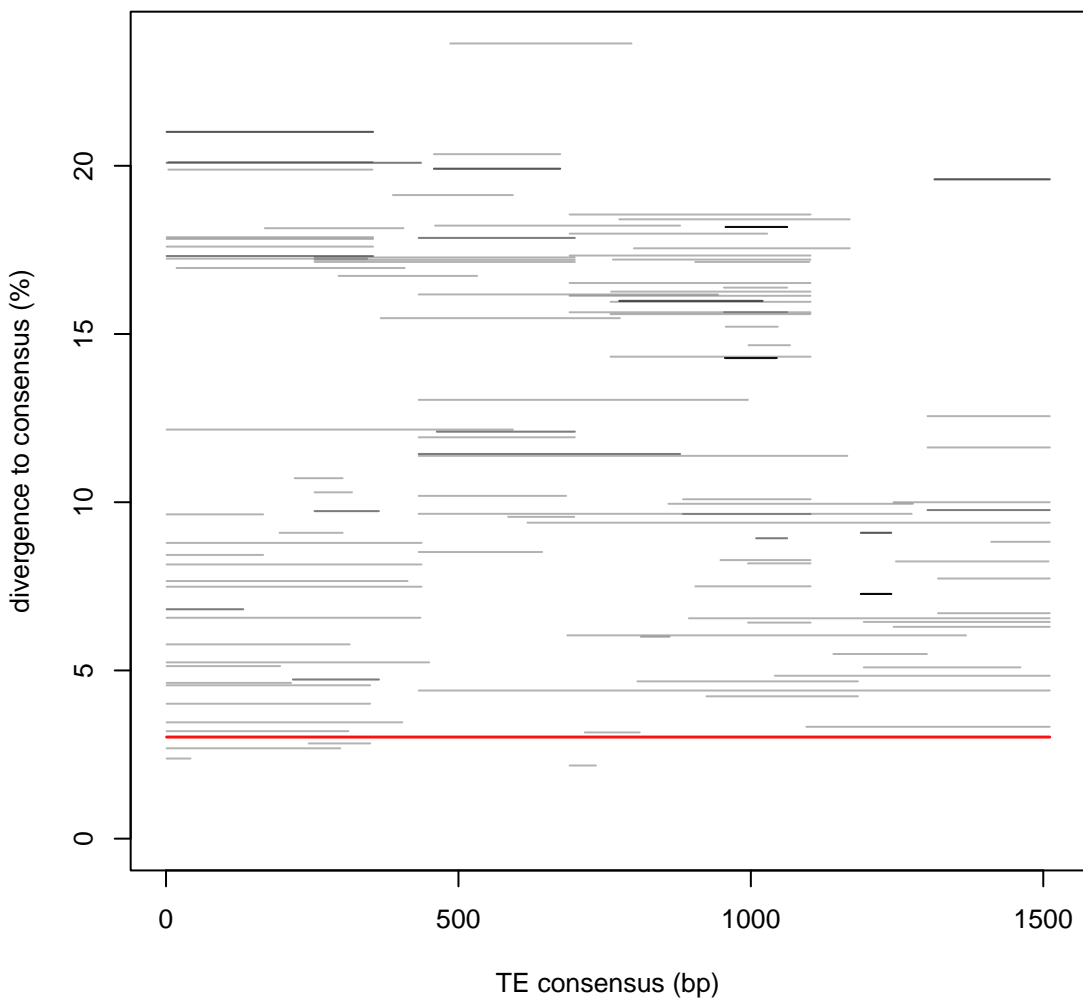
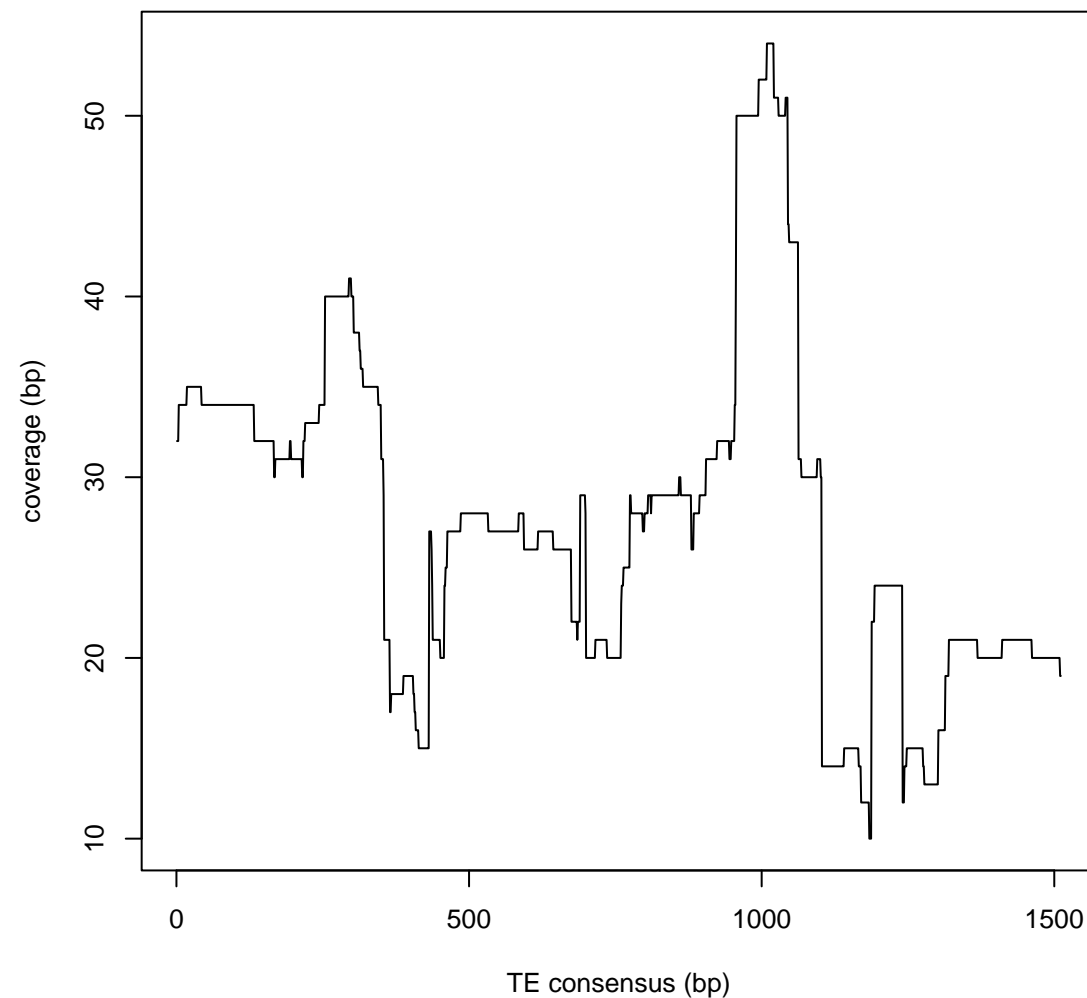


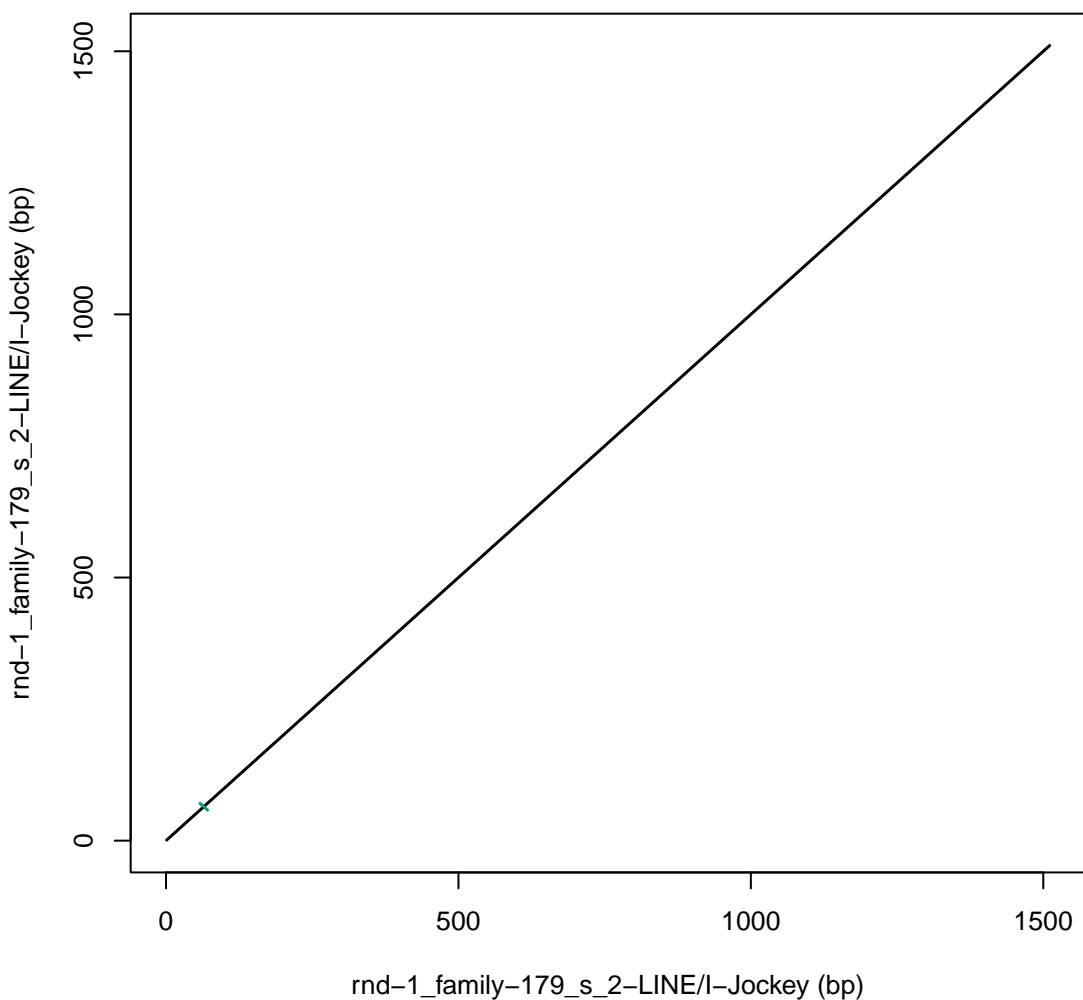
TE: rnd-1_family-179_s_2-LINE/l-Jockey
consensus size: 1511bp; fragments: 154; full length: 1 (≥ 1359.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

