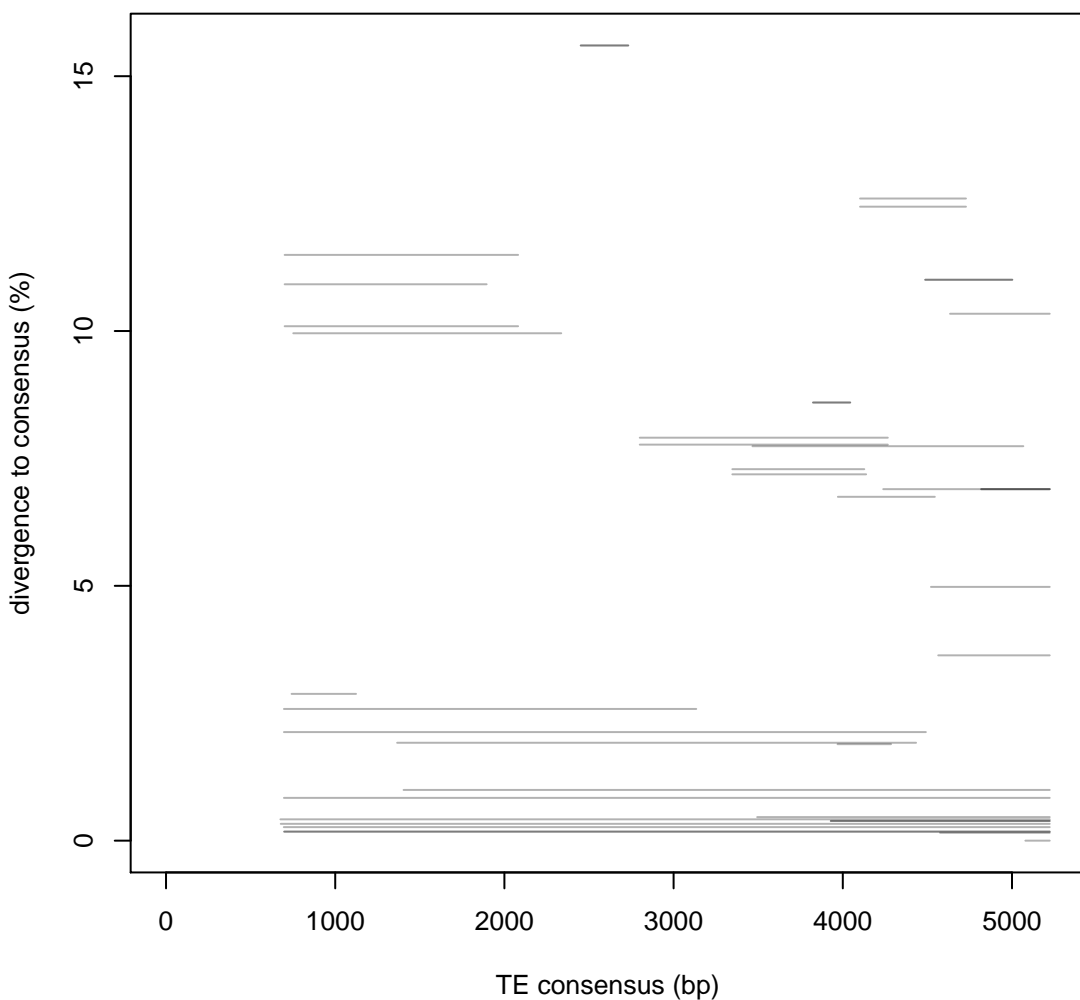
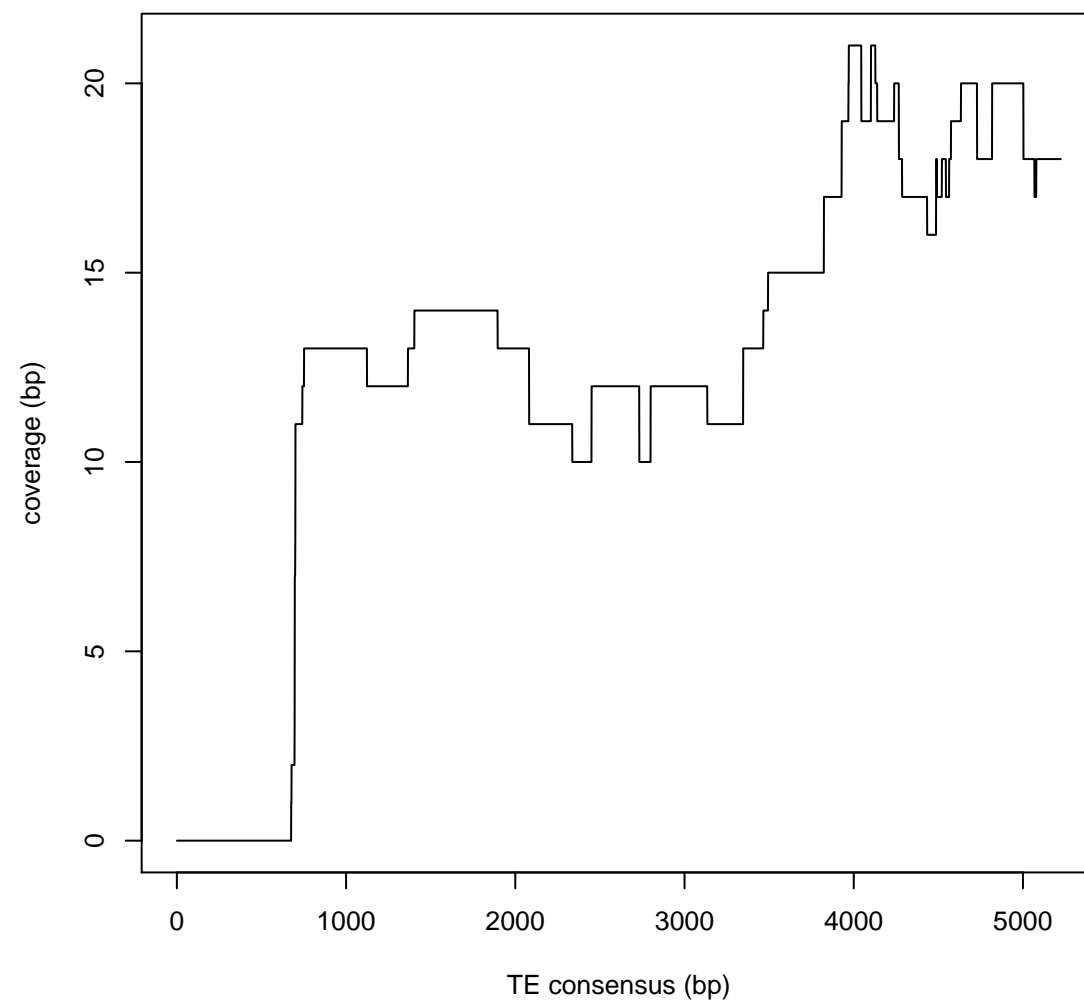


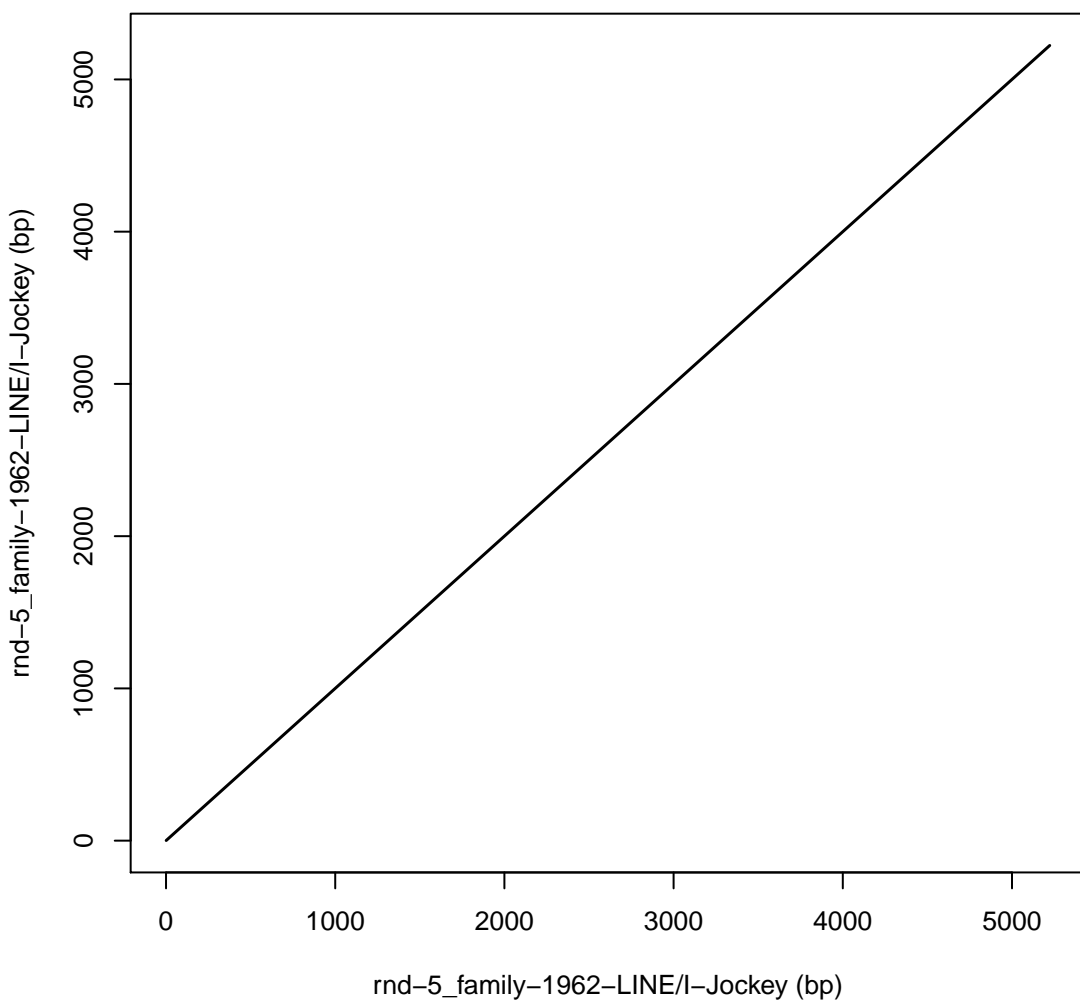
TE: rnd-5_family-1962-LINE/I-Jockey
consensus size: 5223bp; fragments: 41; full length: 0 (≥ 4700.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

