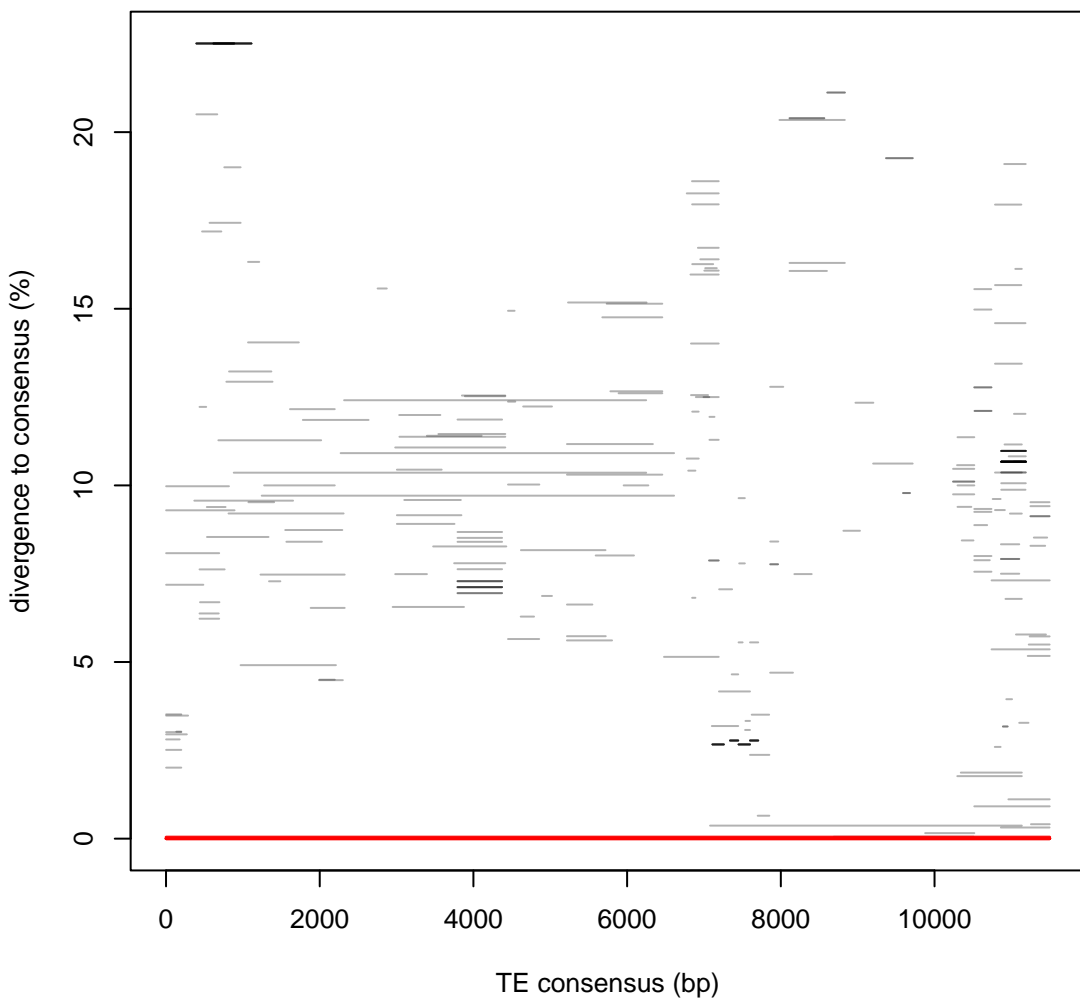
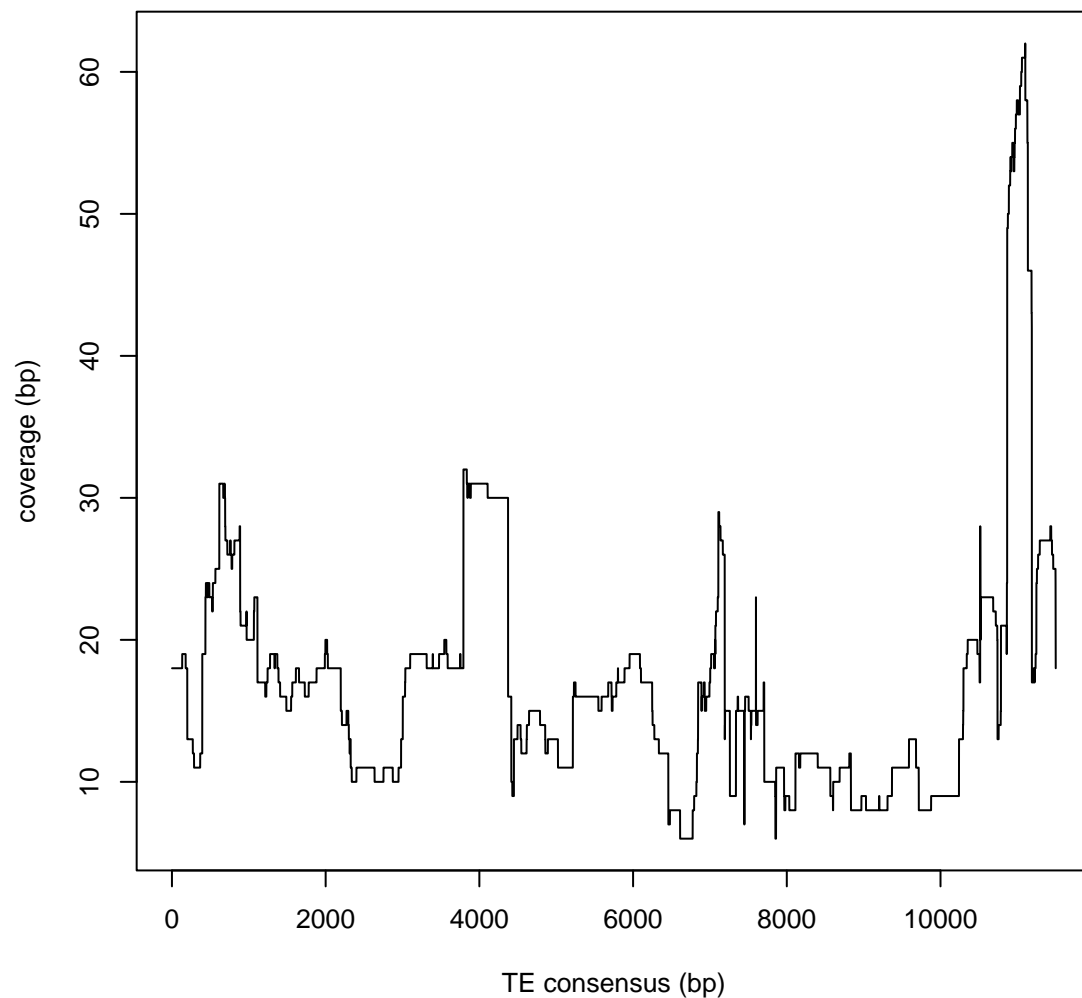


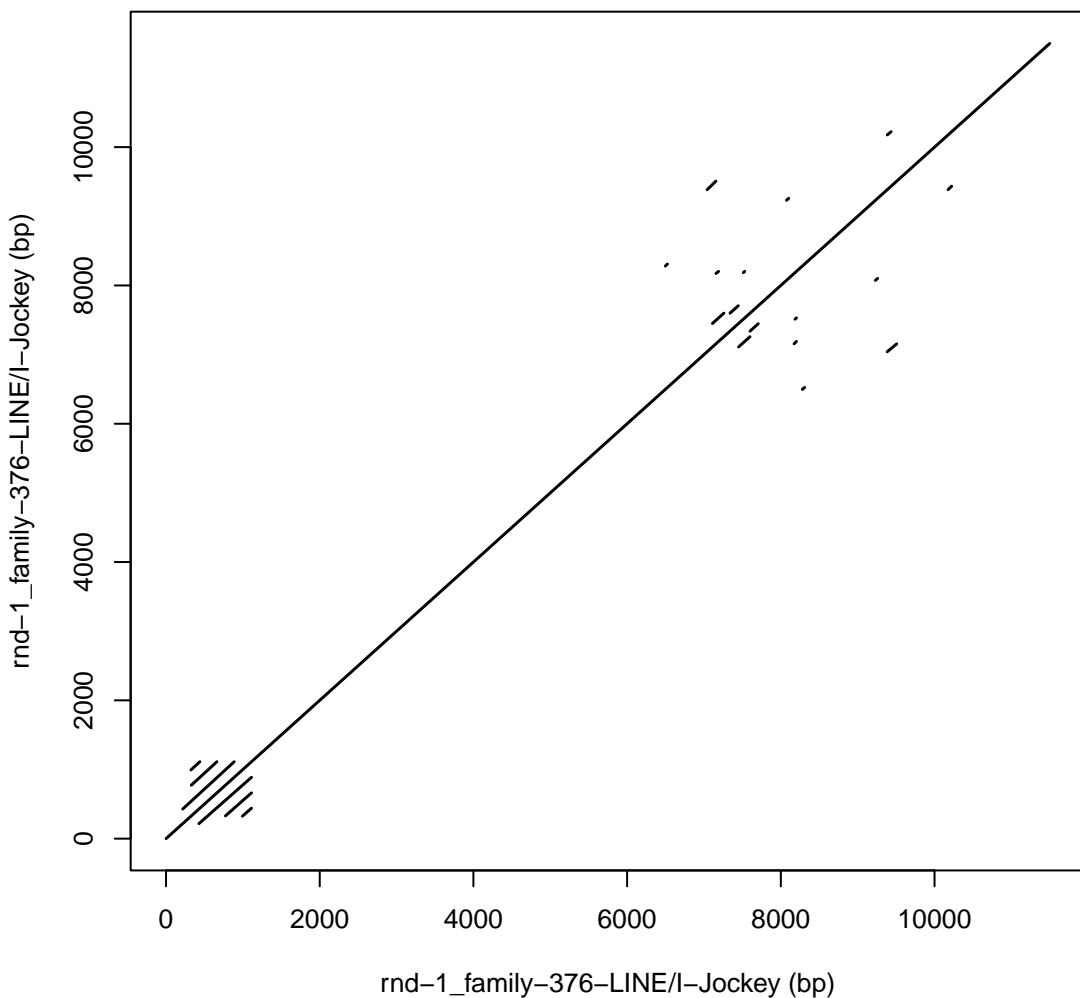
TE: rnd-1_family-376-LINE/I-Jockey
consensus size: 11499bp; fragments: 296; full length: 5 (≥ 10349.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

