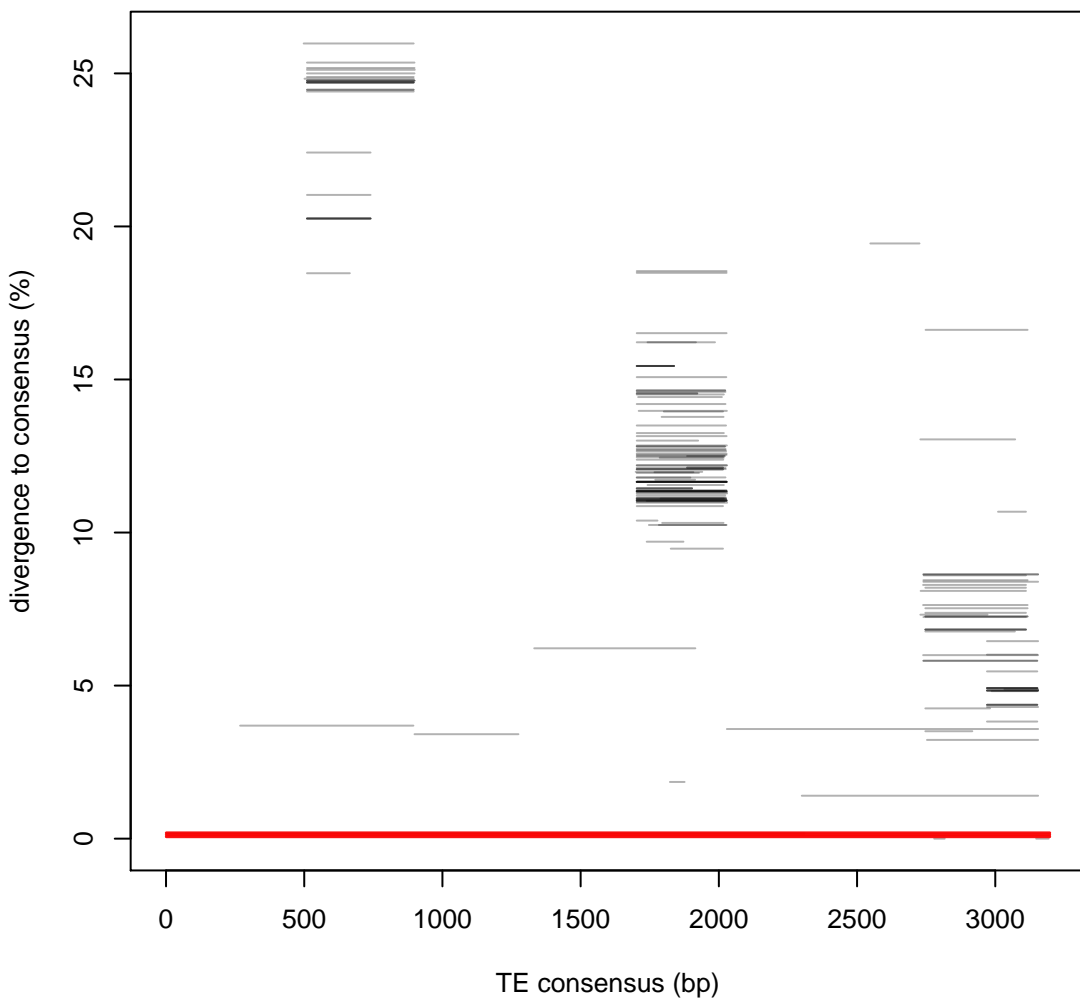
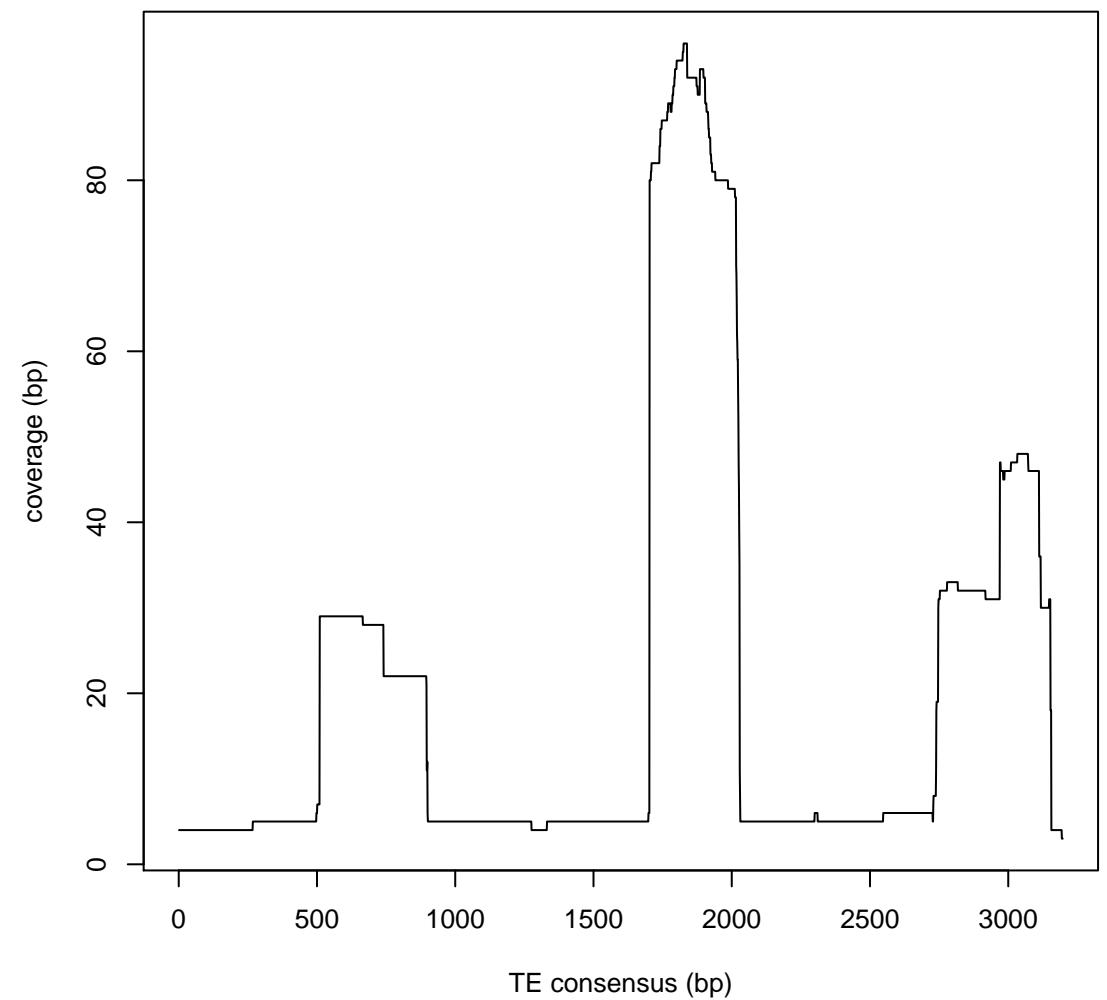


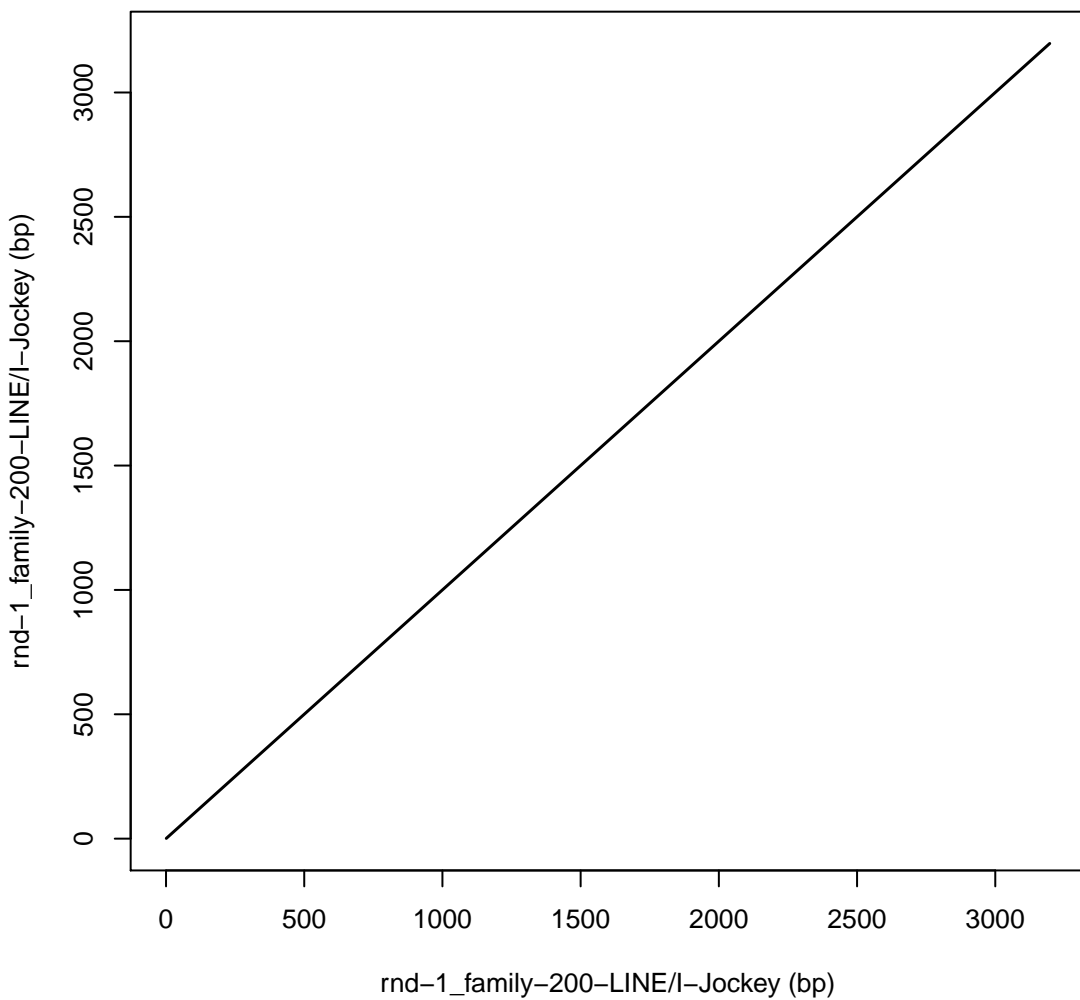
TE: rnd-1_family-200-LINE/l-Jockey
consensus size: 3197bp; fragments: 177; full length: 3 (≥ 2877.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

