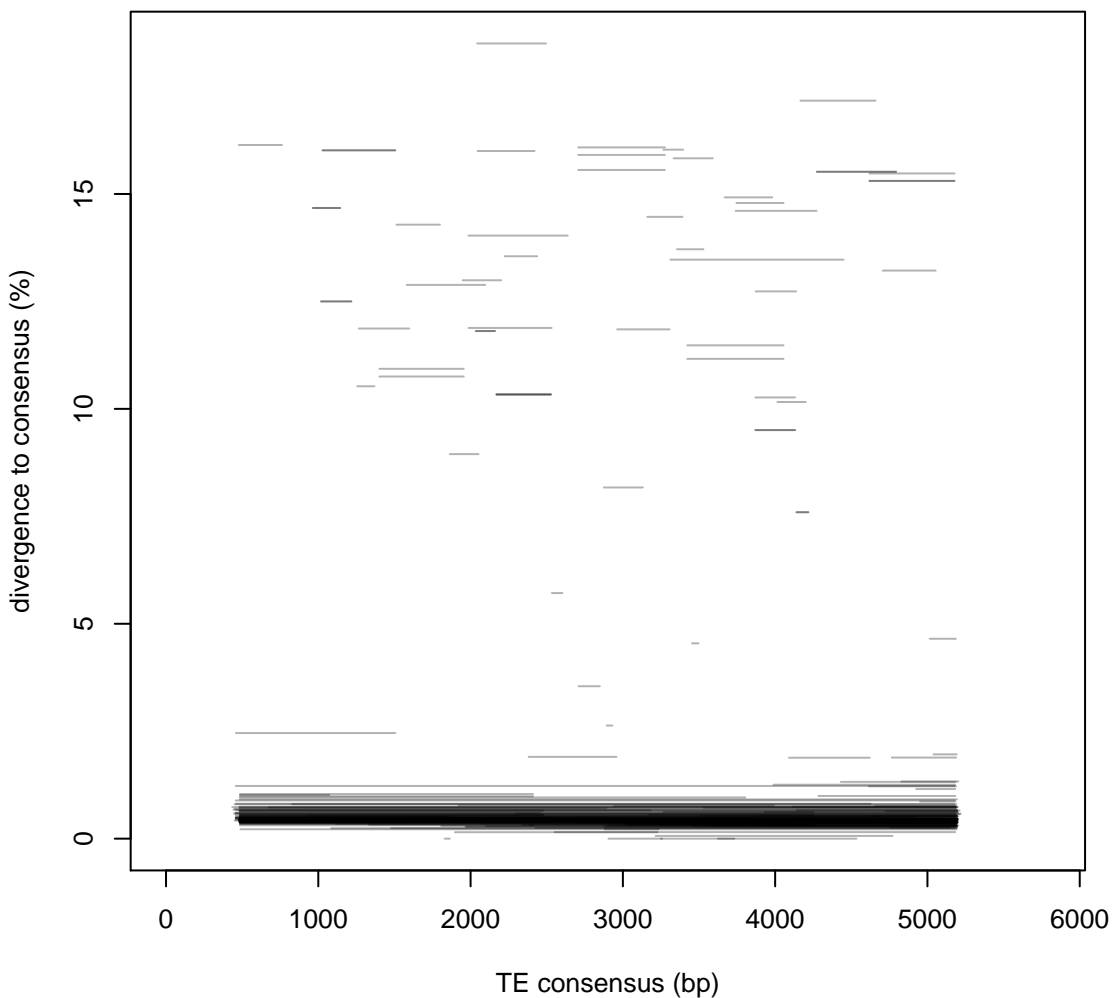
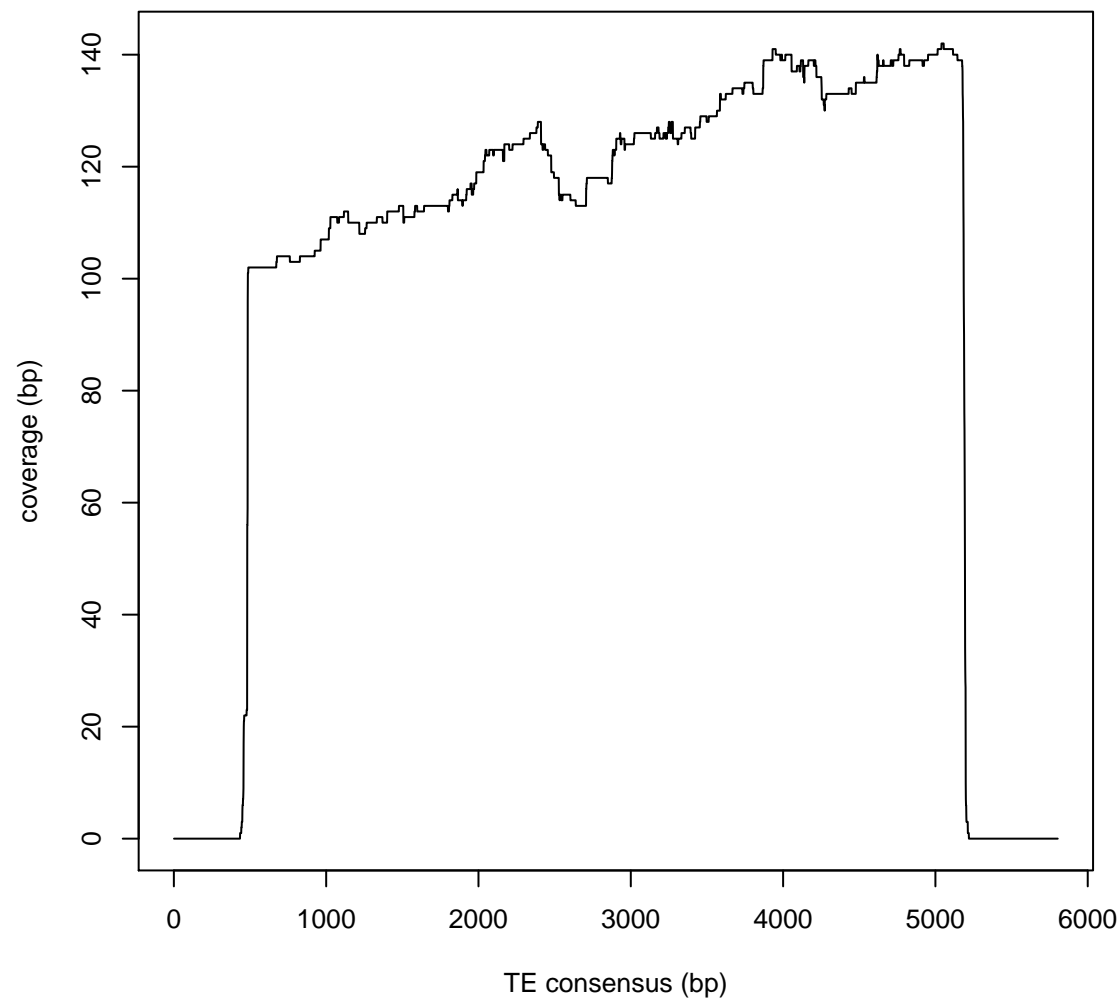


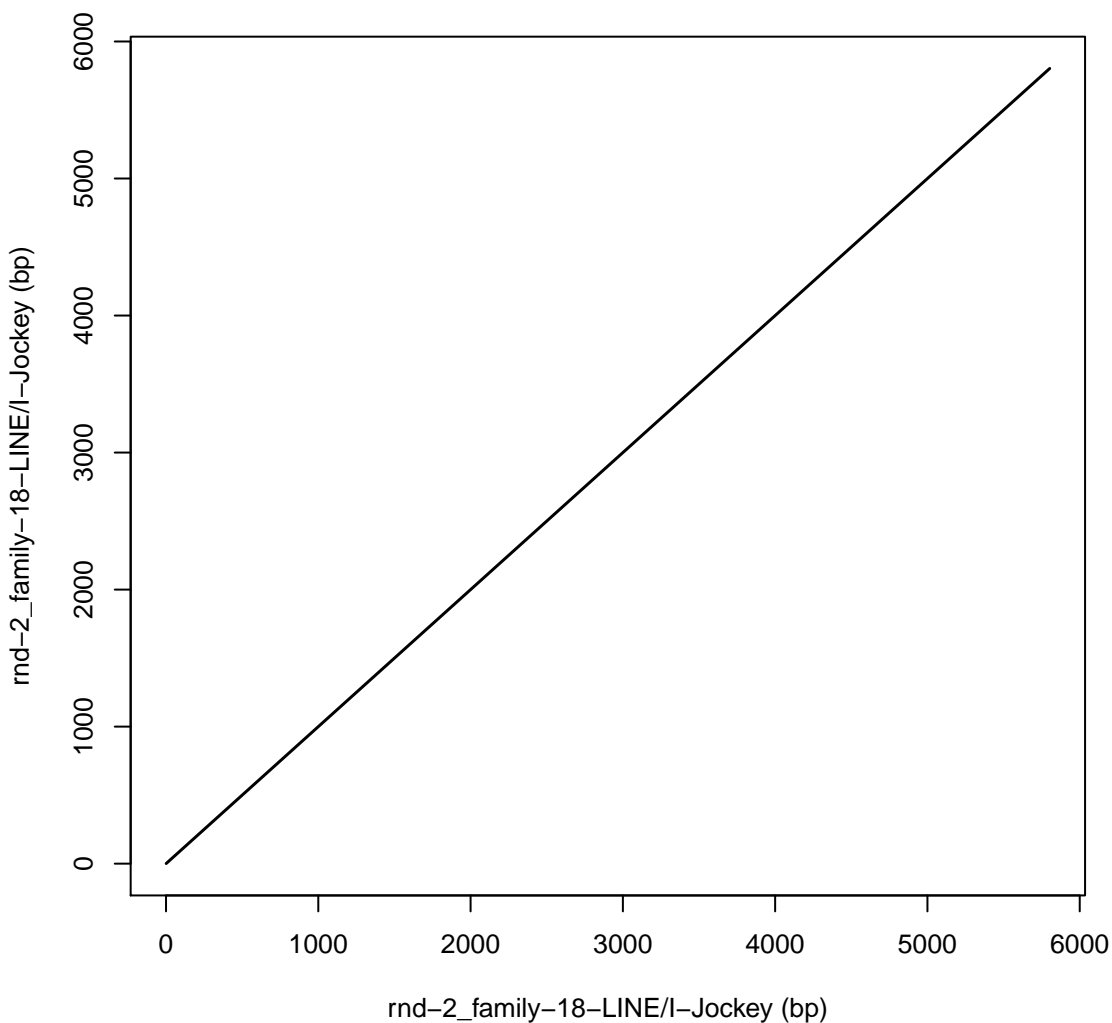
TE: rnd-2_family-18-LINE/I-Jockey
consensus size: 5803bp; fragments: 238; full length: 0 (≥ 5222.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

