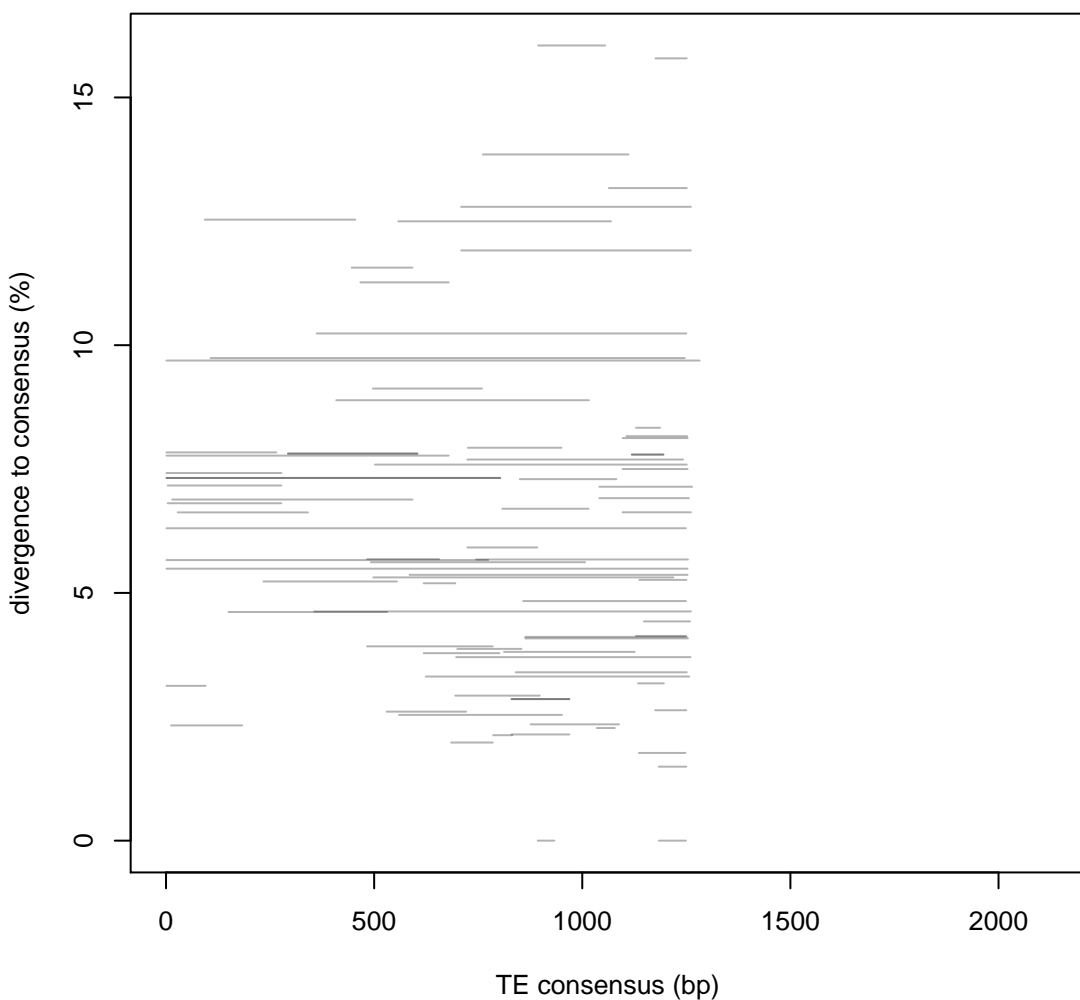
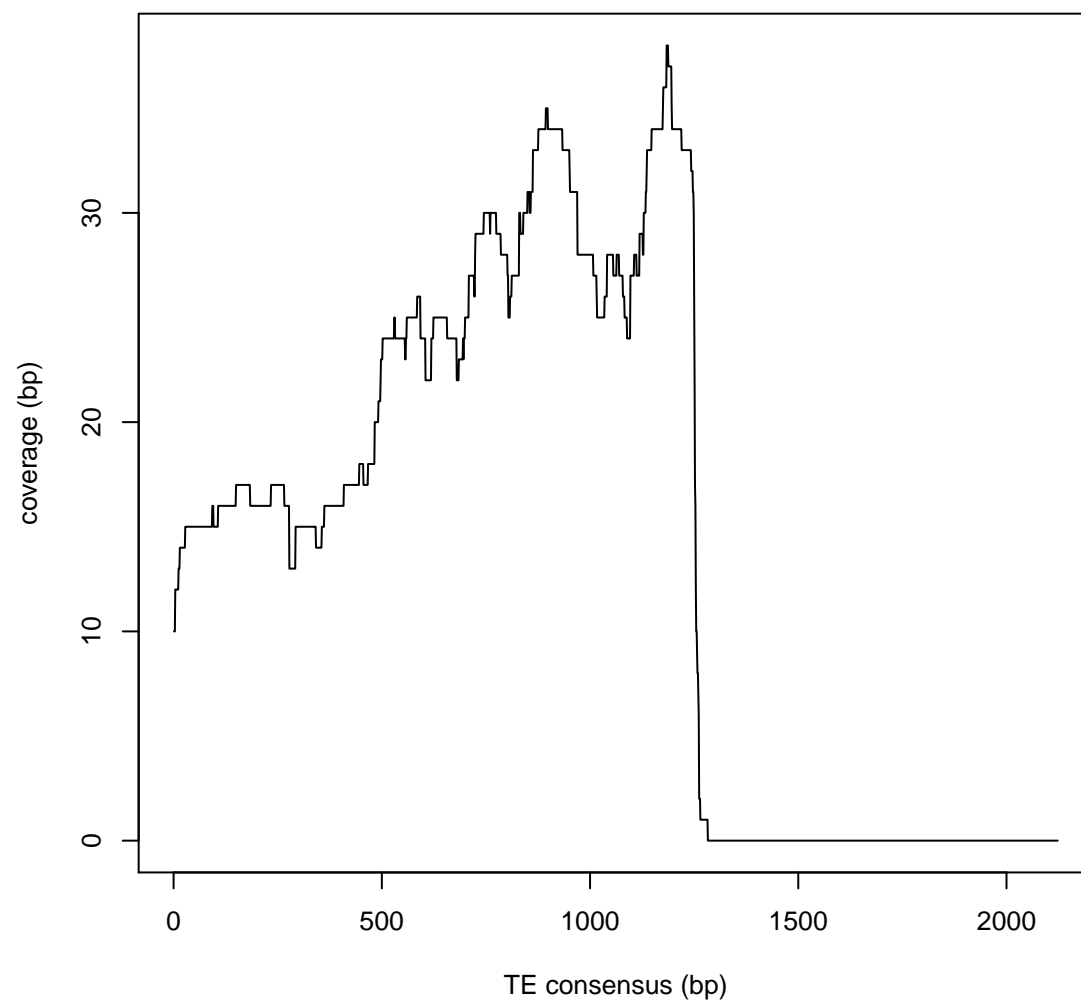


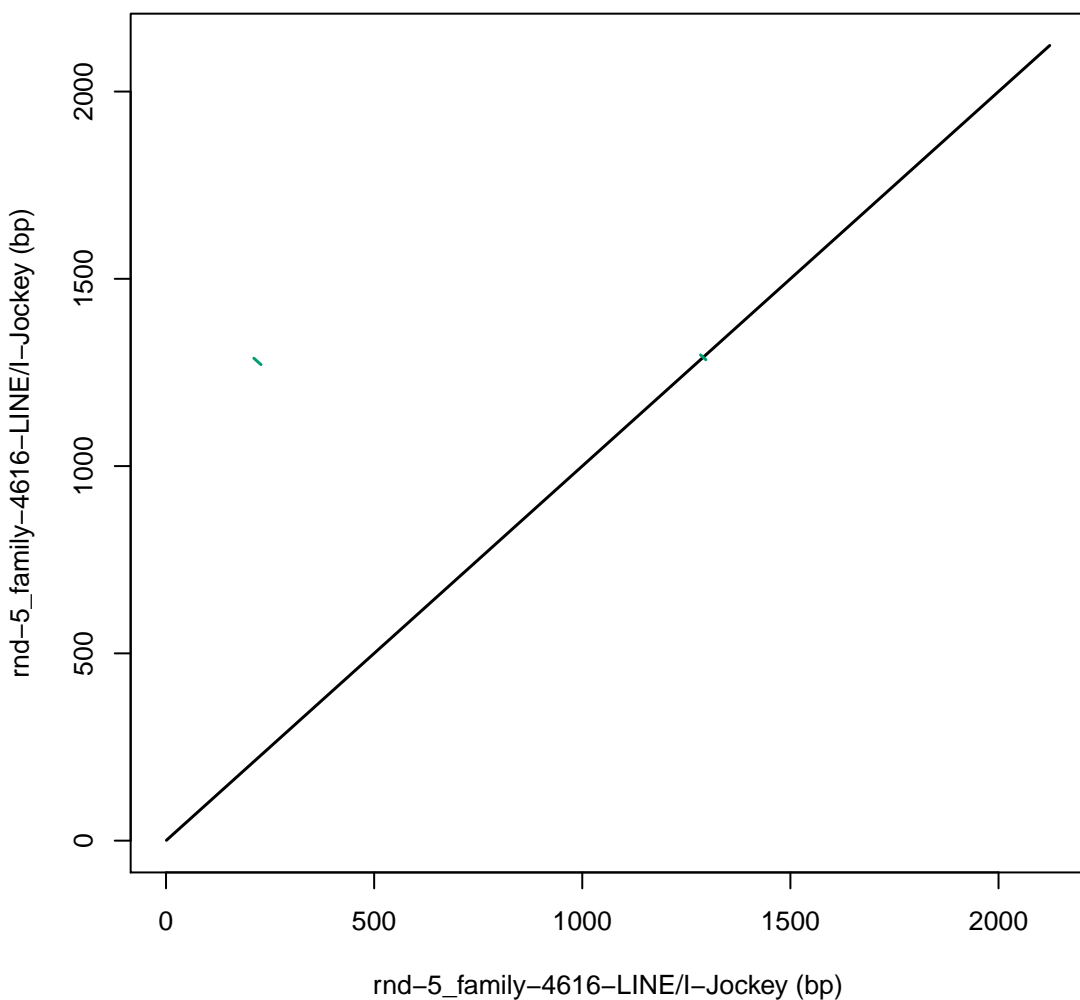
TE: rnd-5_family-4616-LINE/I-Jockey
consensus size: 2123bp; fragments: 84; full length: 0 (≥ 1910.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

