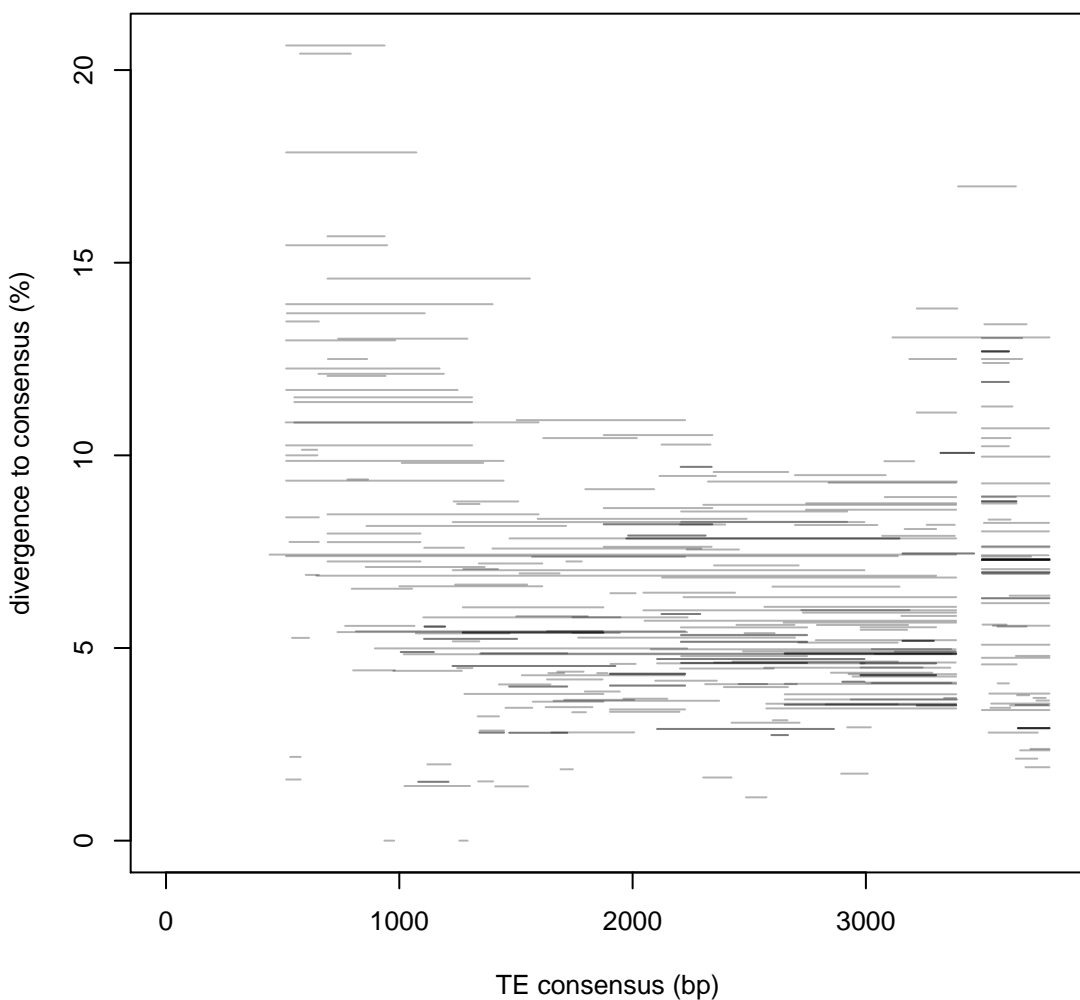
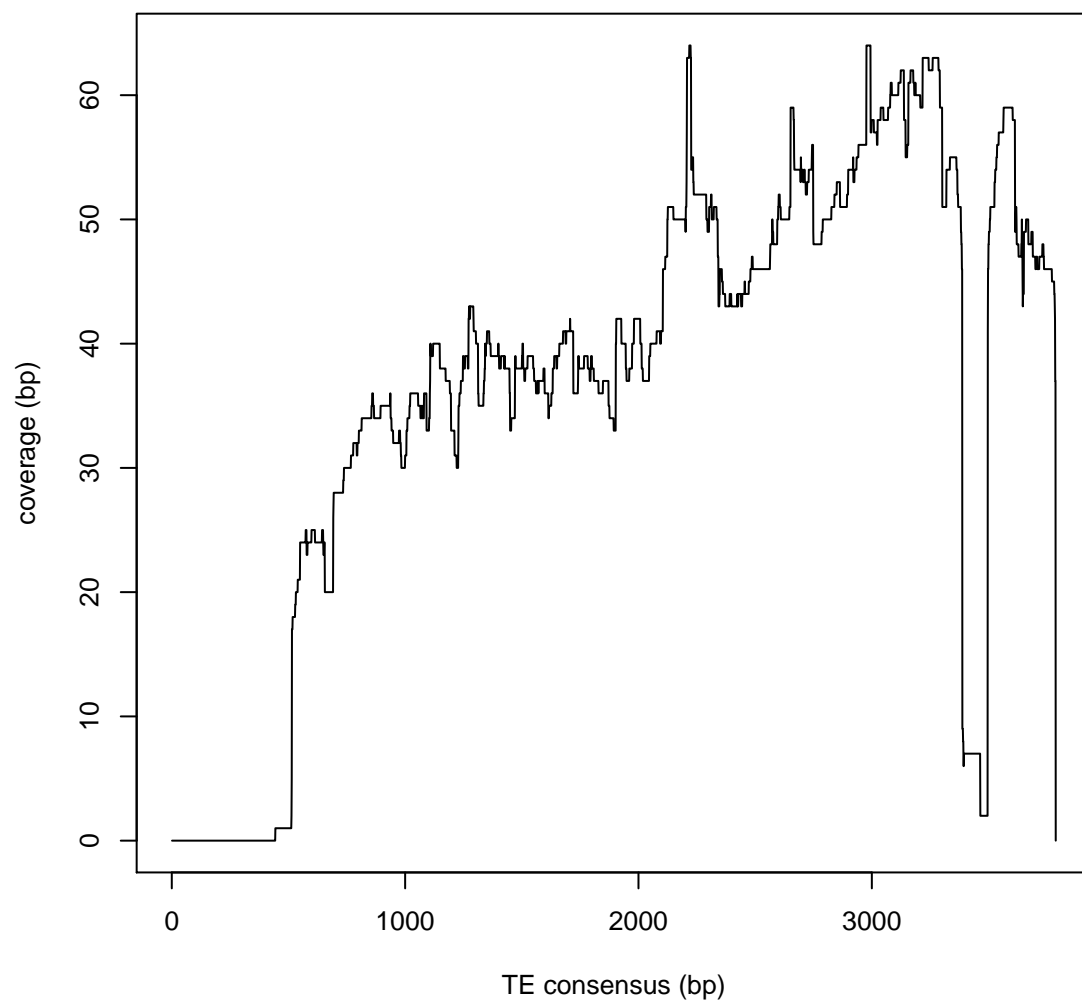


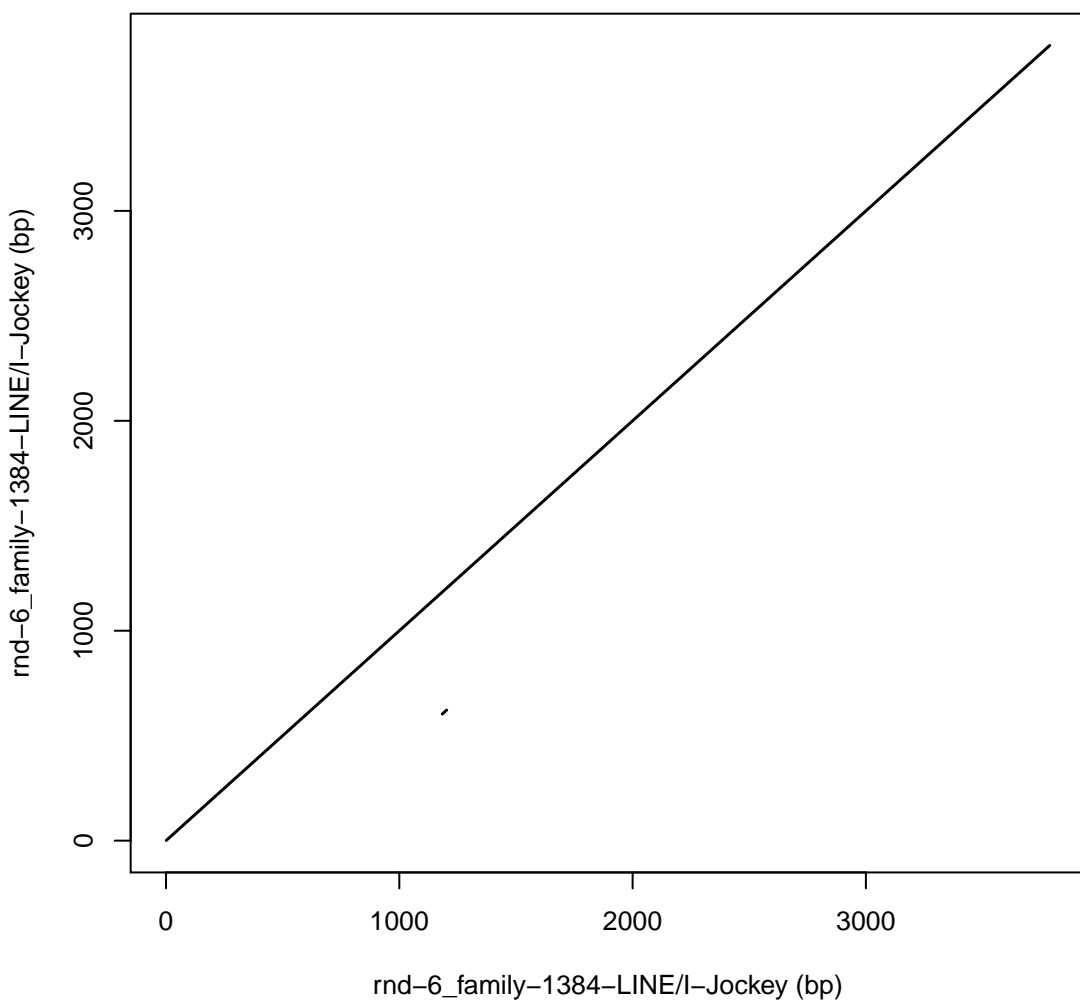
TE: rnd-6_family-1384-LINE/I-Jockey
consensus size: 3788bp; fragments: 333; full length: 0 (≥ 3409.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

