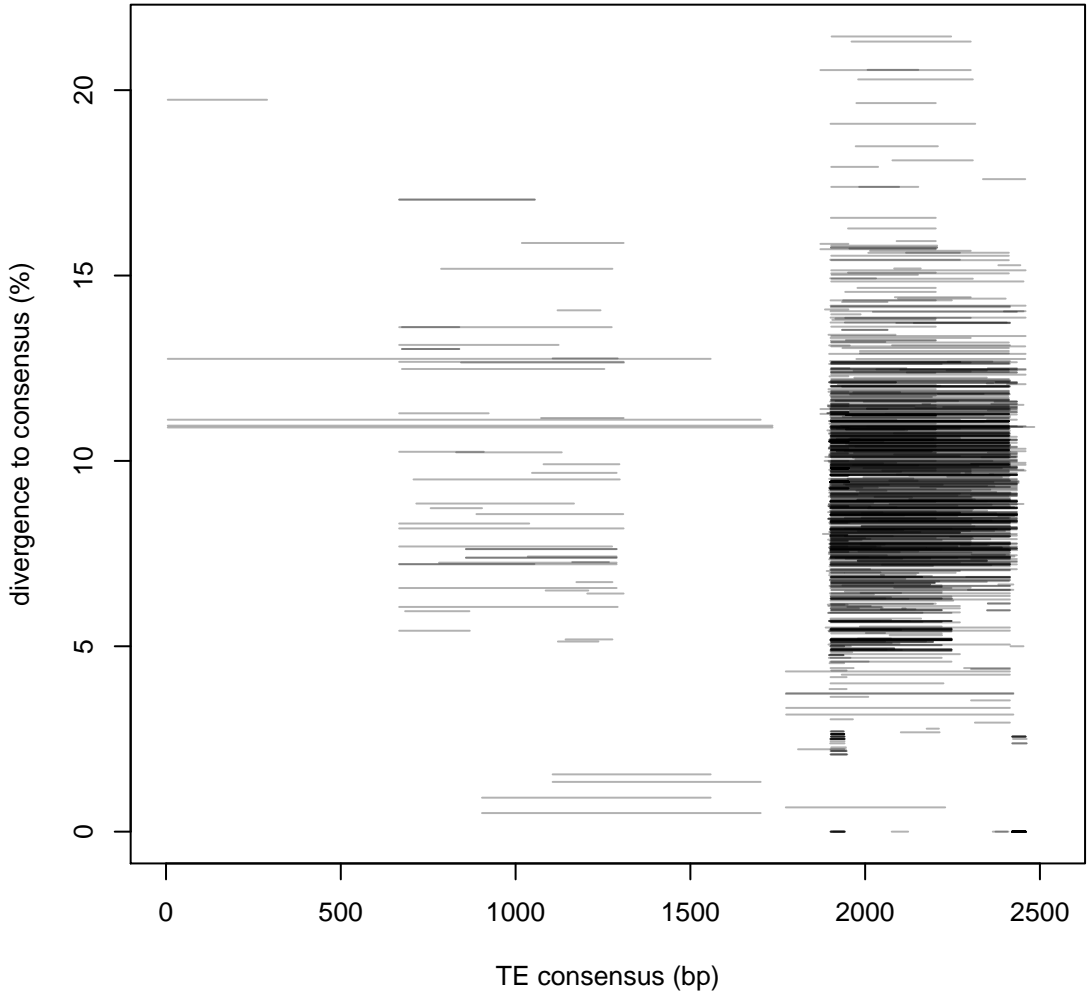
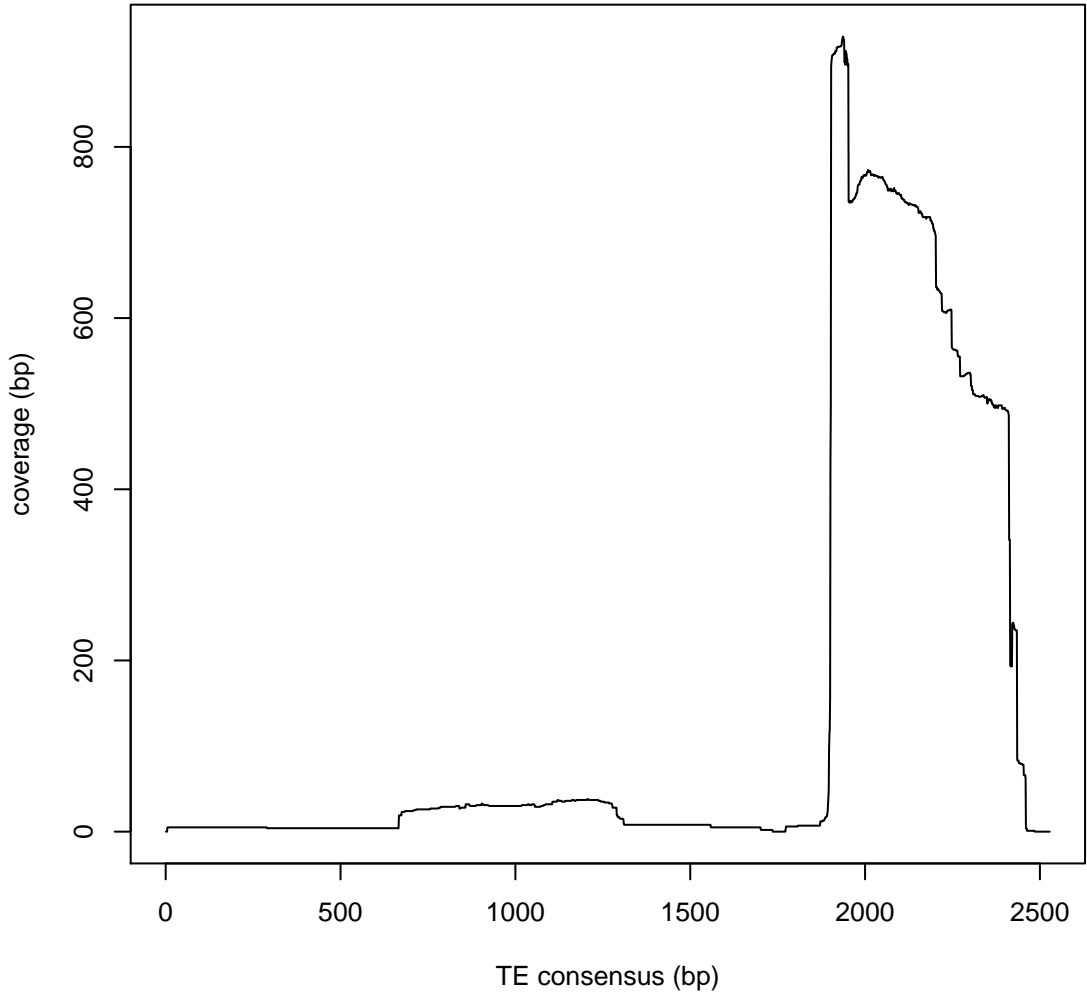


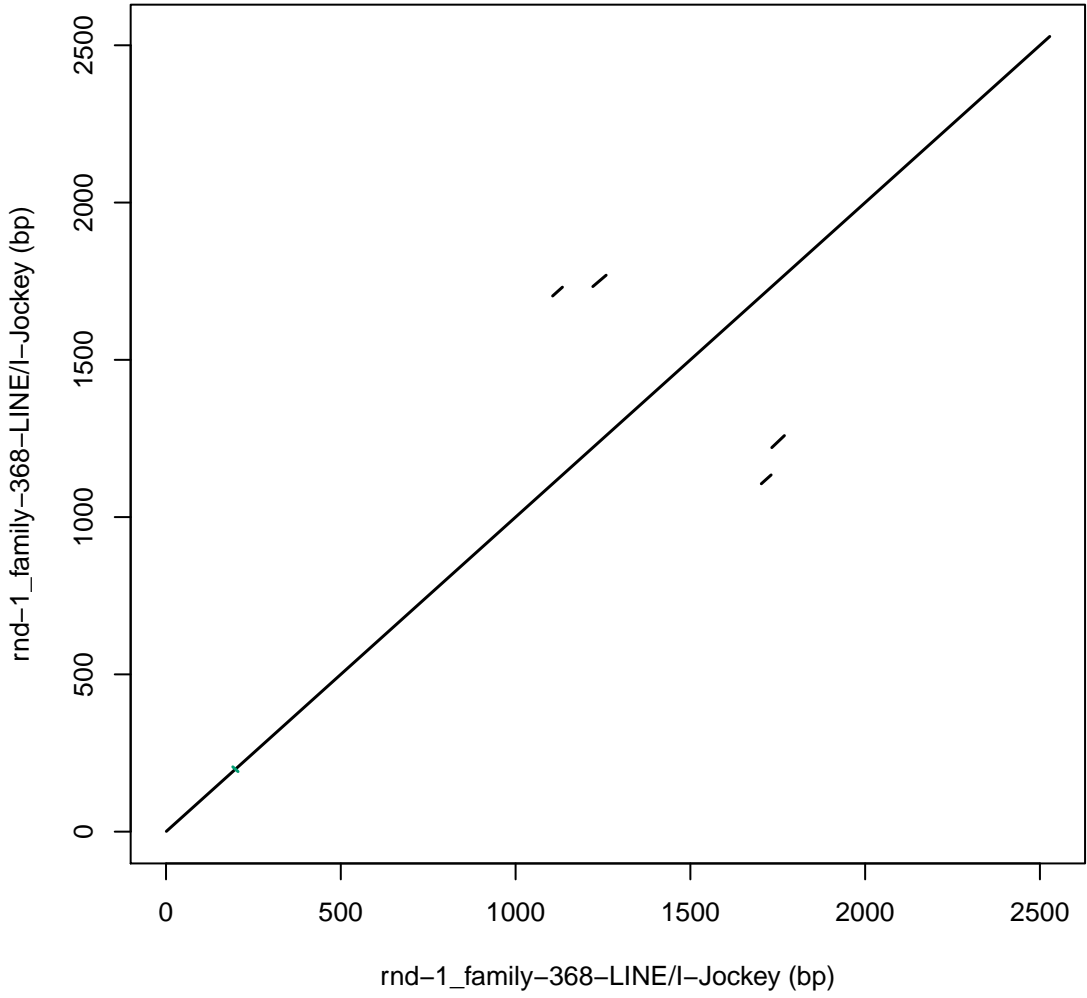
TE: rnd-1_family-368-LINE/I-Jockey
consensus size: 2528bp; fragments: 1206; full length: 0 (>=2275.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

