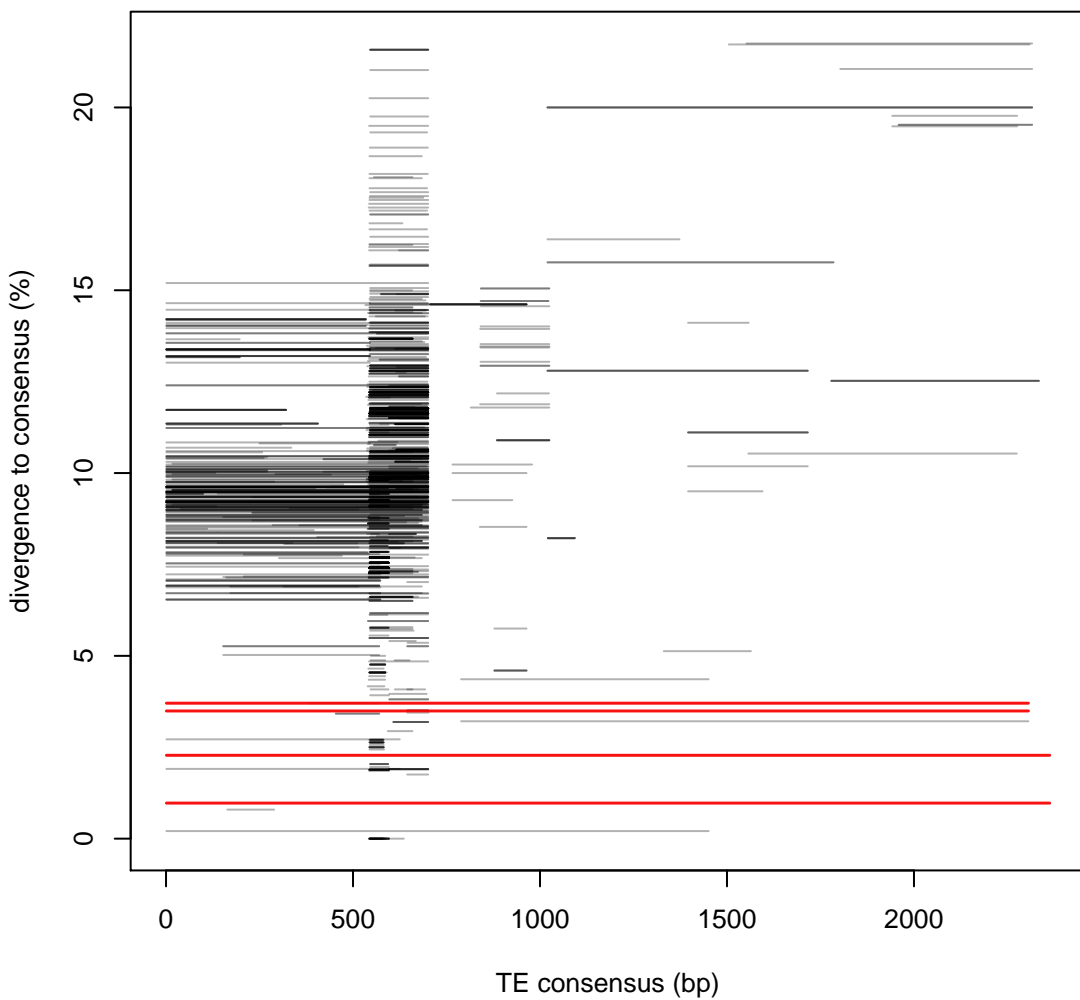
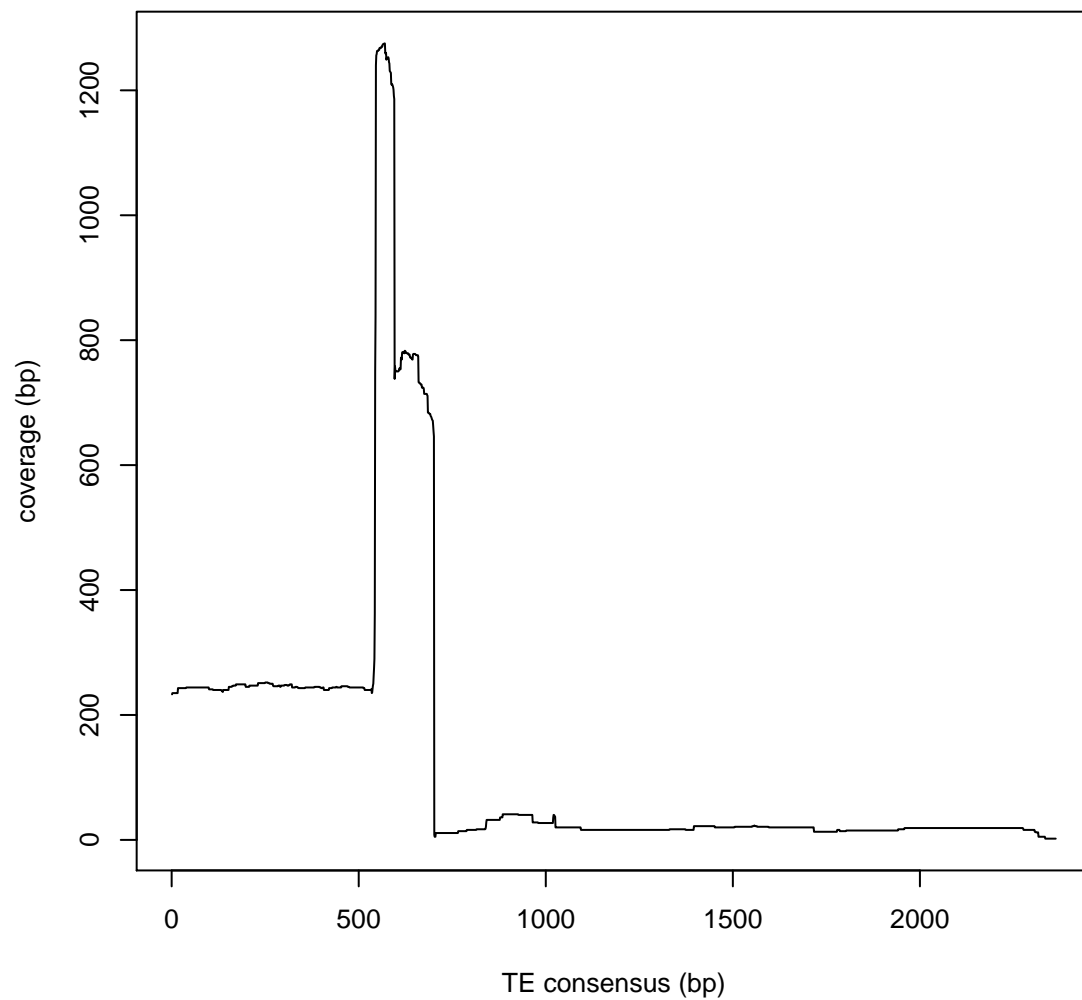


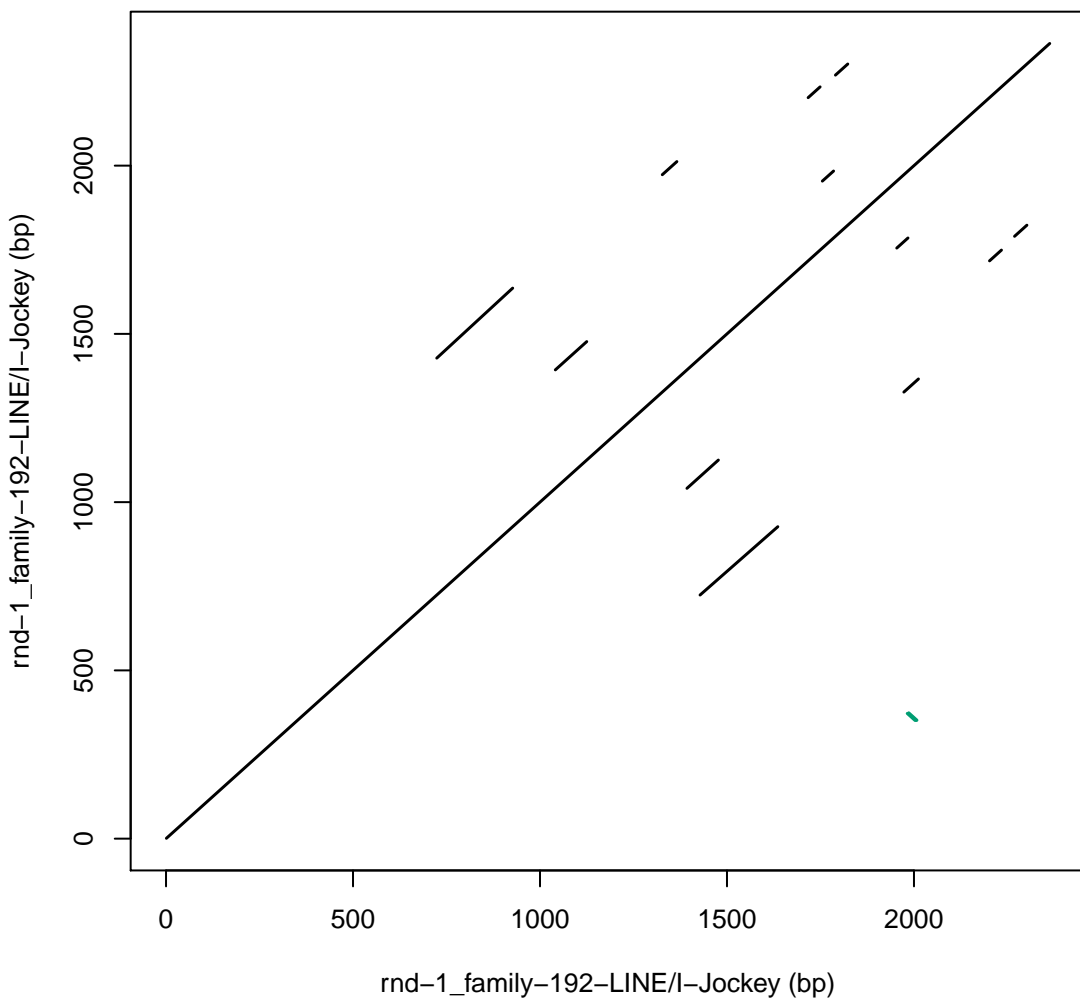
TE: rnd-1_family-192-LINE/I-Jockey
consensus size: 2363bp; fragments: 1516; full length: 4 (≥ 2126.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

