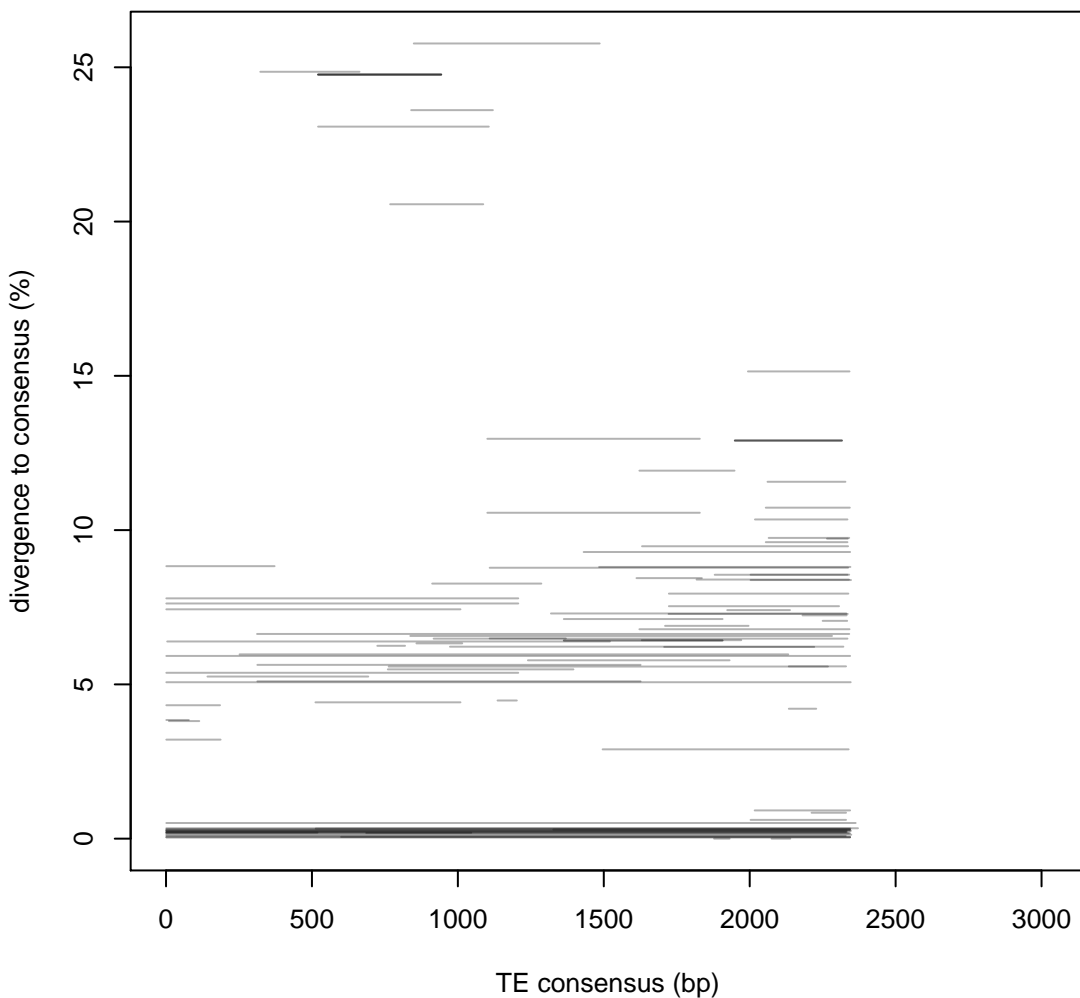
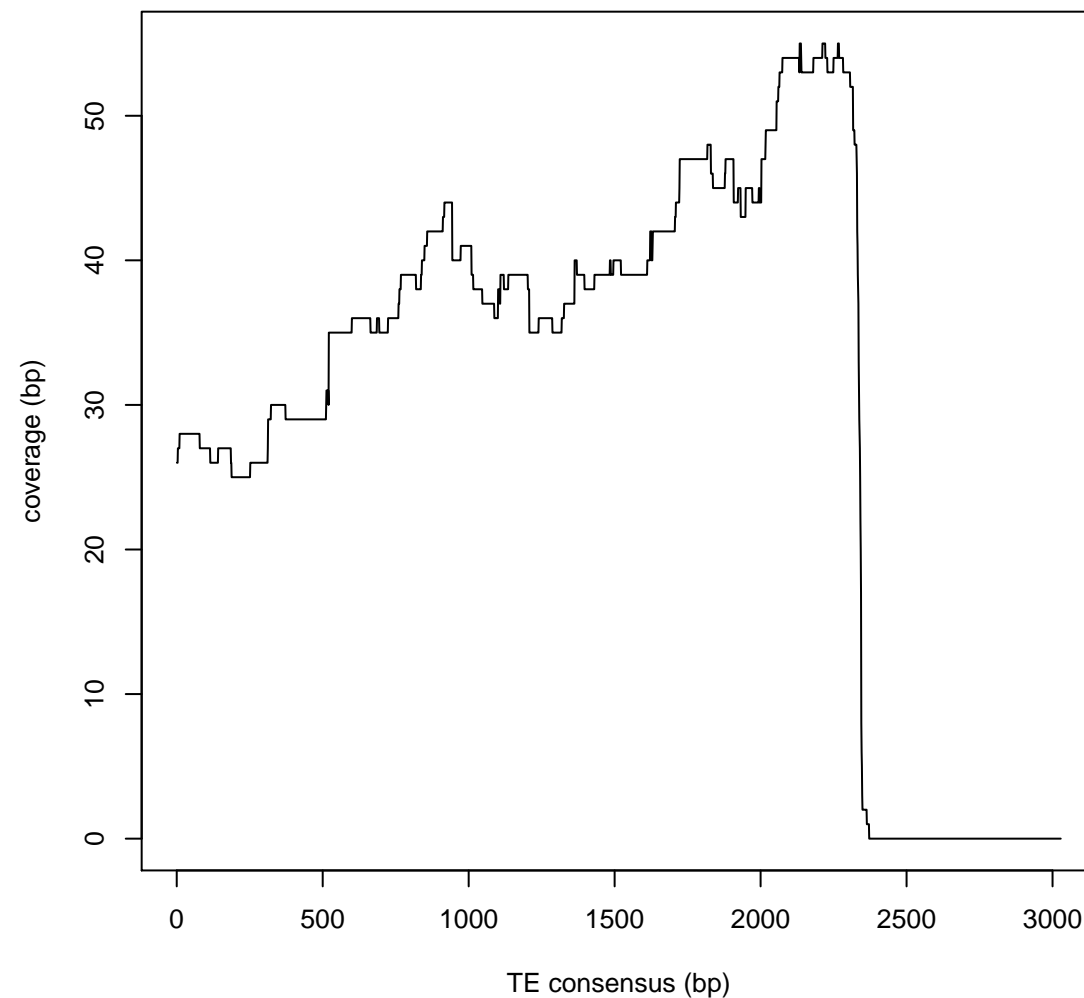


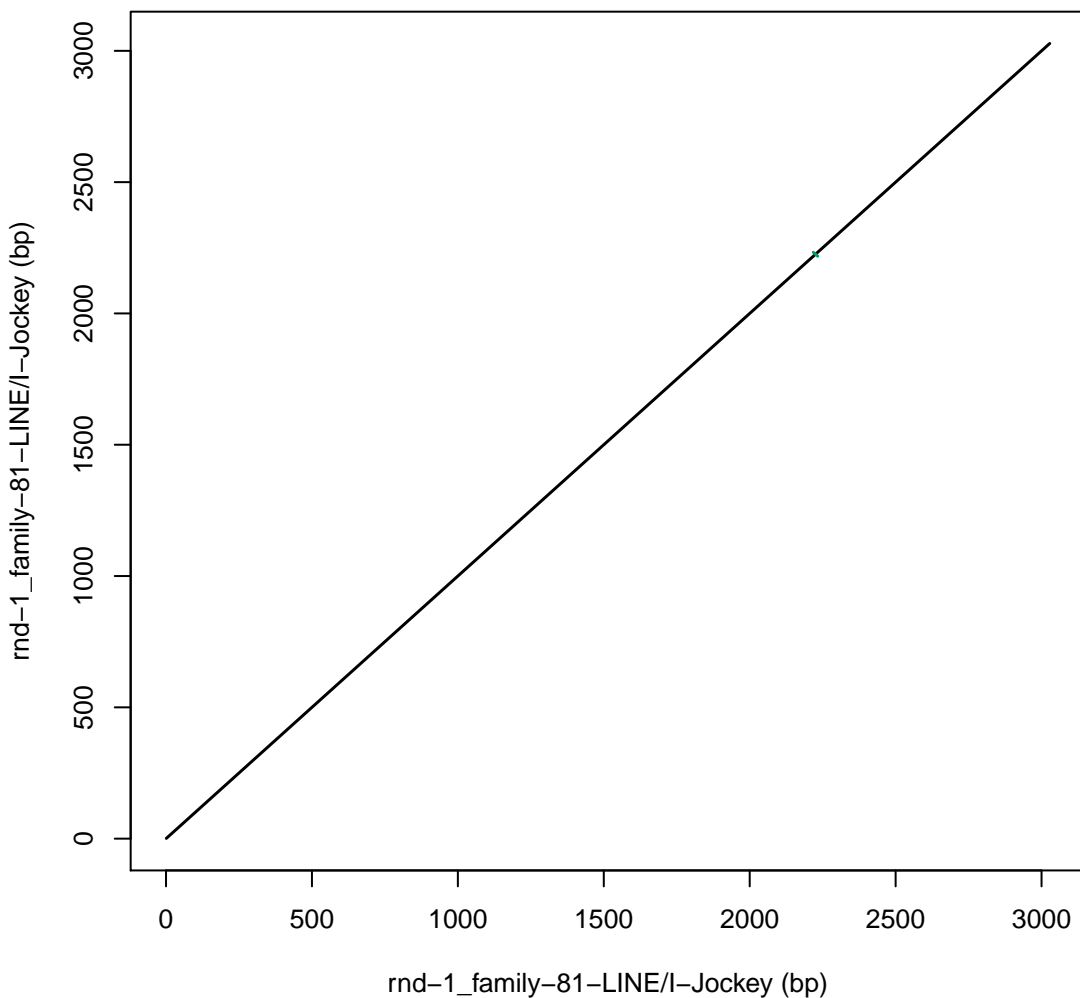
TE: rnd-1_family-81-LINE/I-Jockey
consensus size: 3028bp; fragments: 102; full length: 0 (>=2725.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

