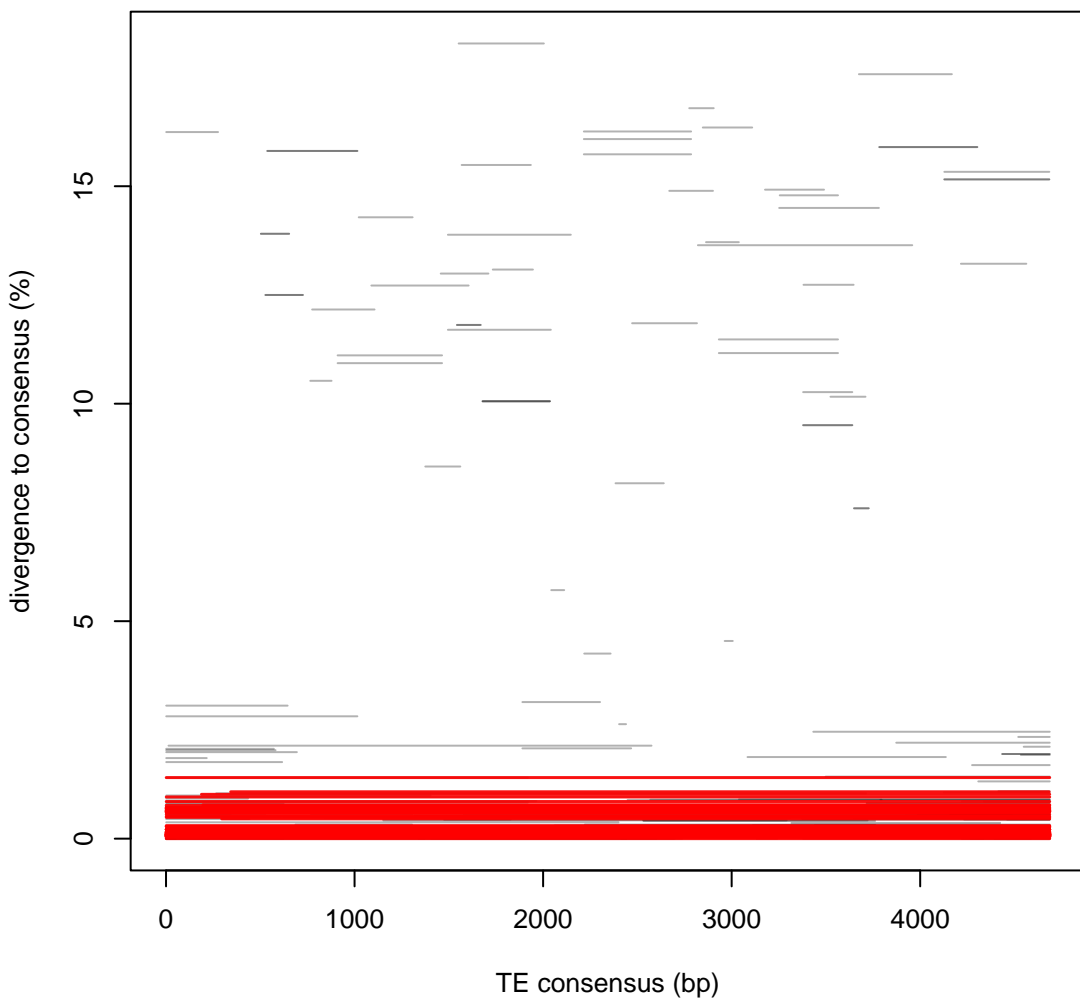
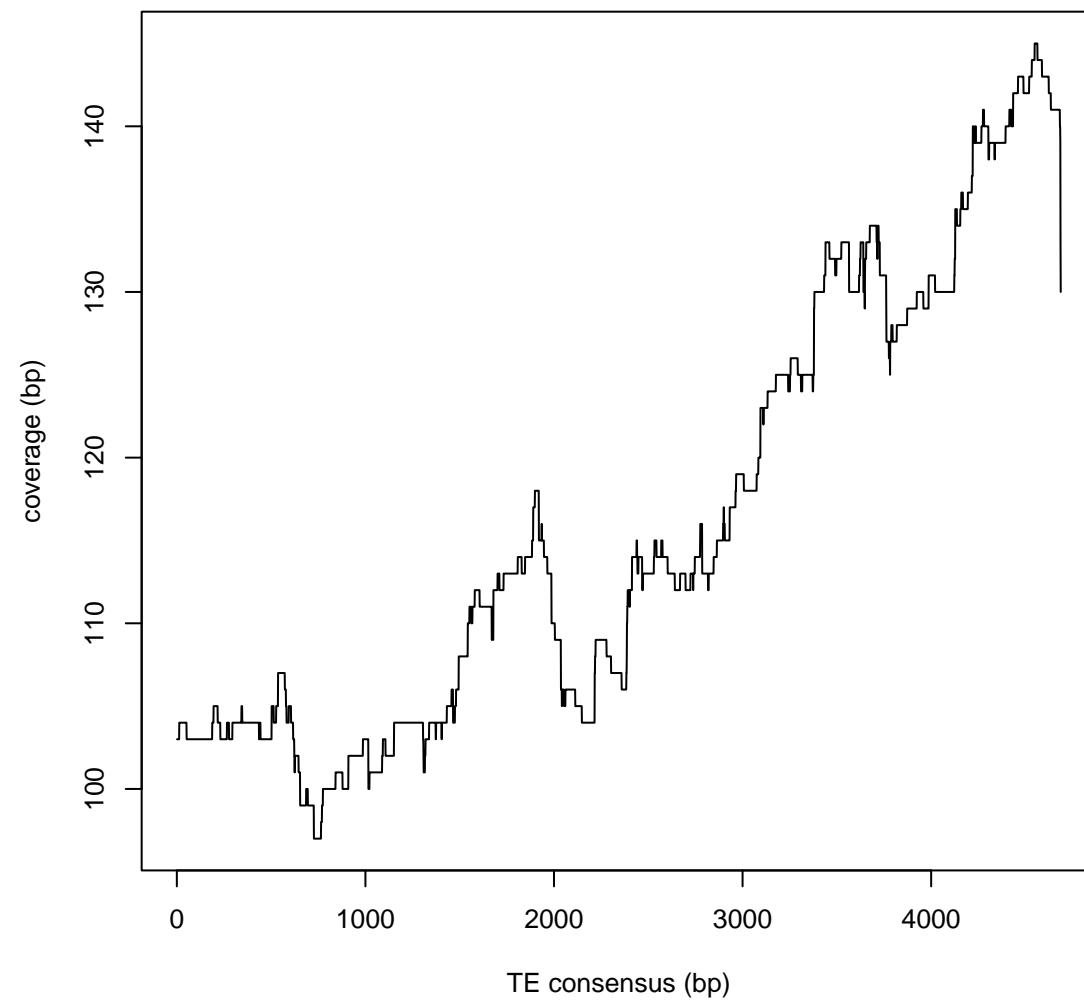


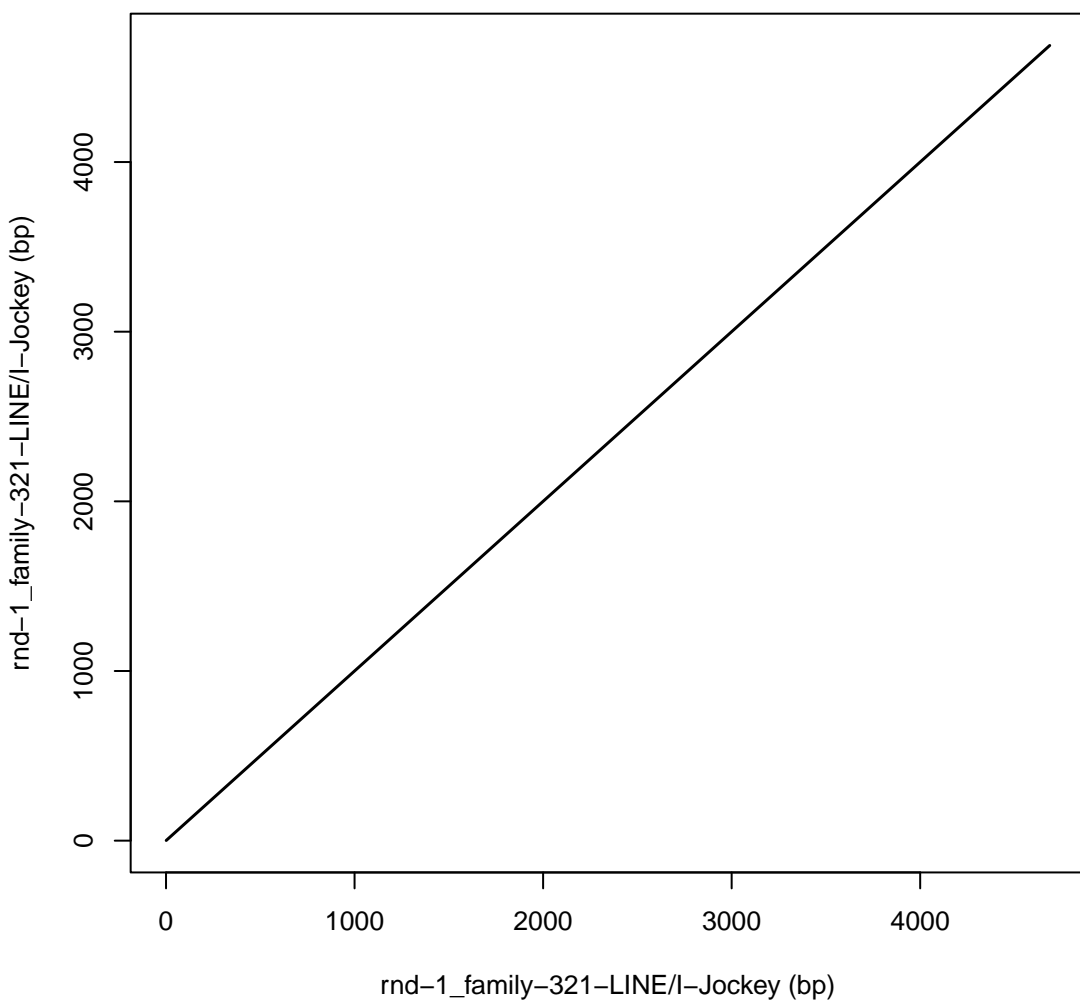
TE: rnd-1_family-321-LINE/I-Jockey
consensus size: 4687bp; fragments: 262; full length: 75 (>=4218.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

