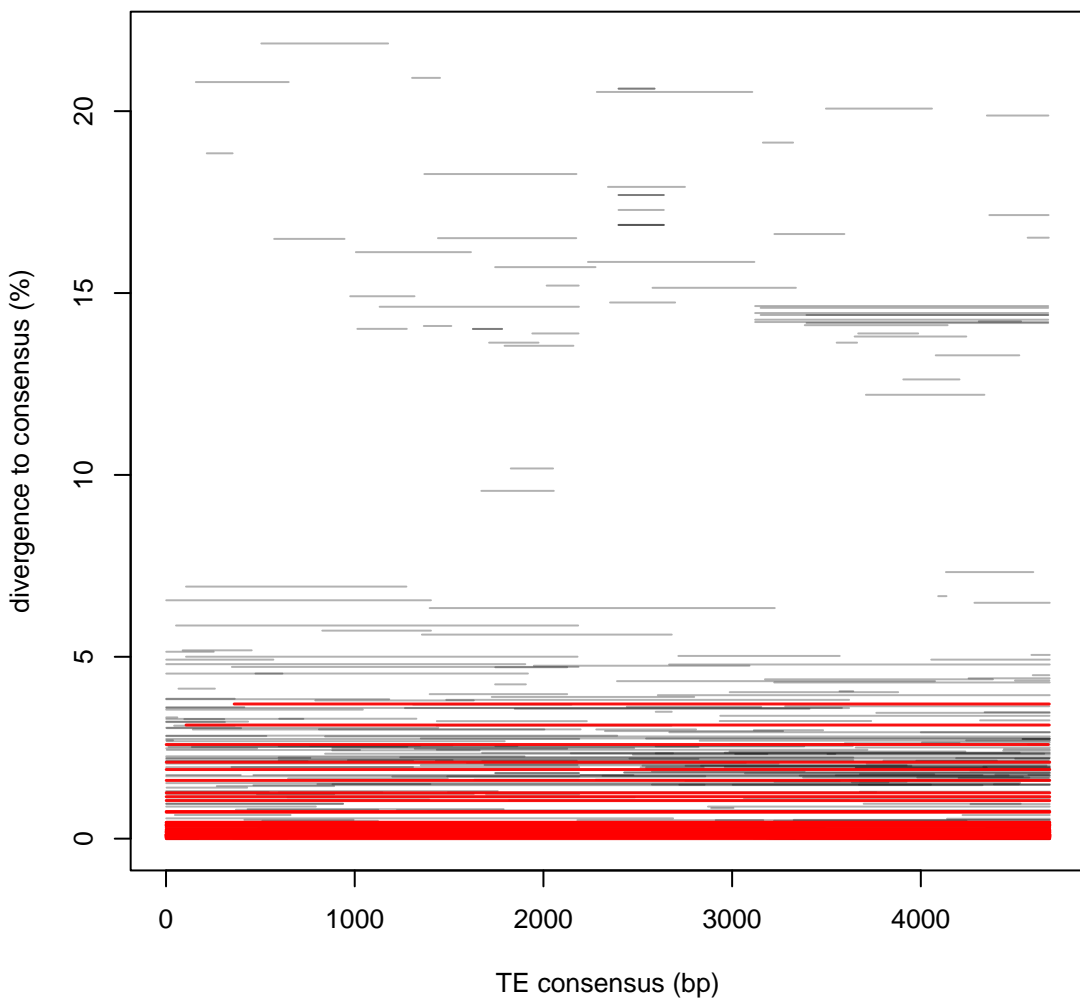
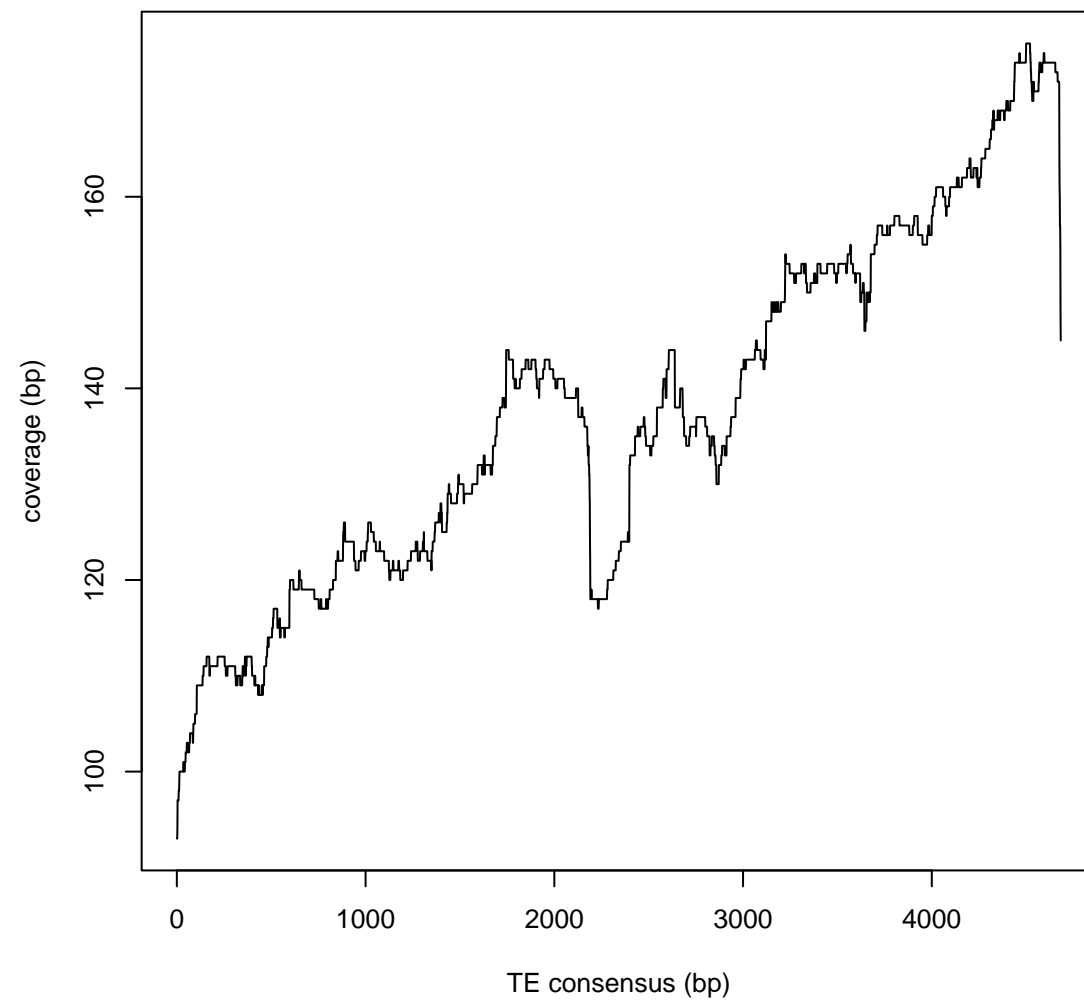


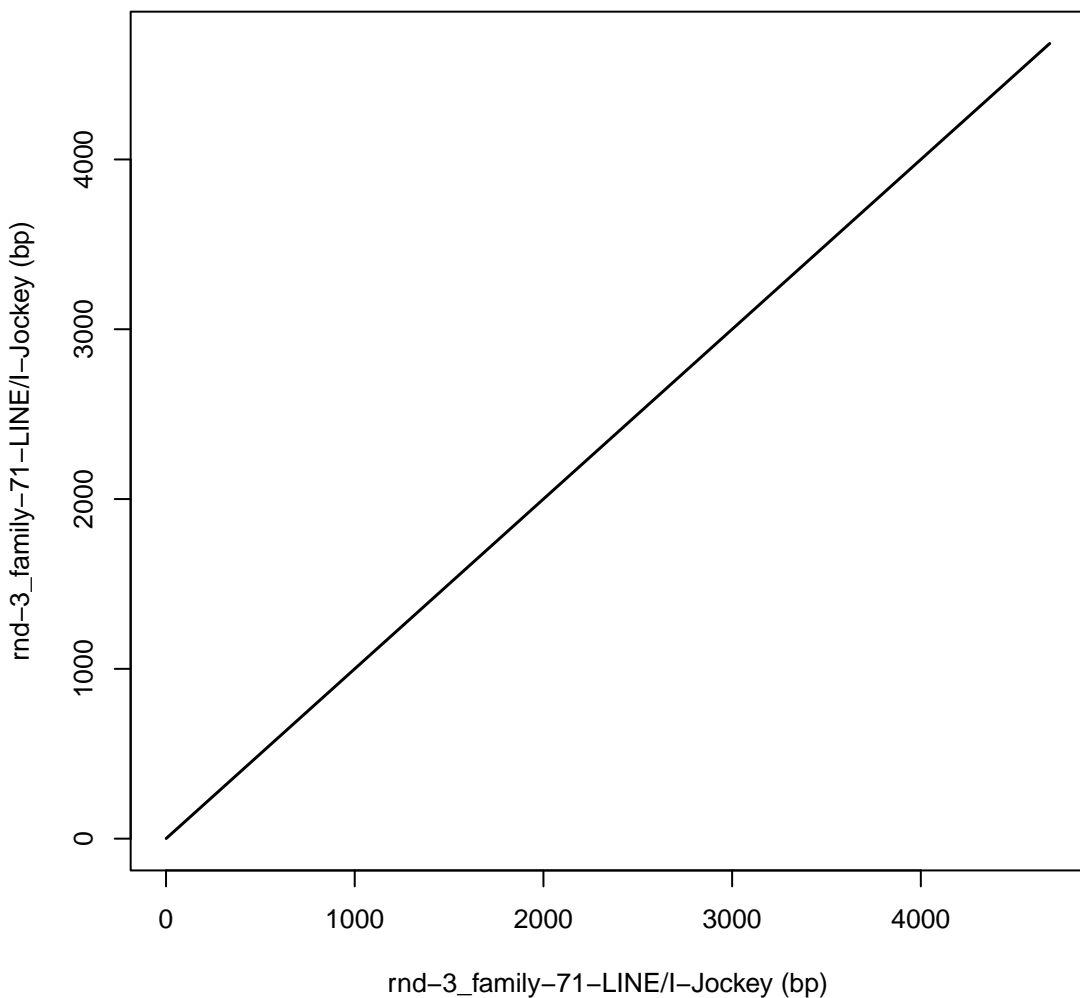
TE: rnd-3_family-71-LINE/I-Jockey
consensus size: 4683bp; fragments: 394; full length: 51 (>=4214.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

