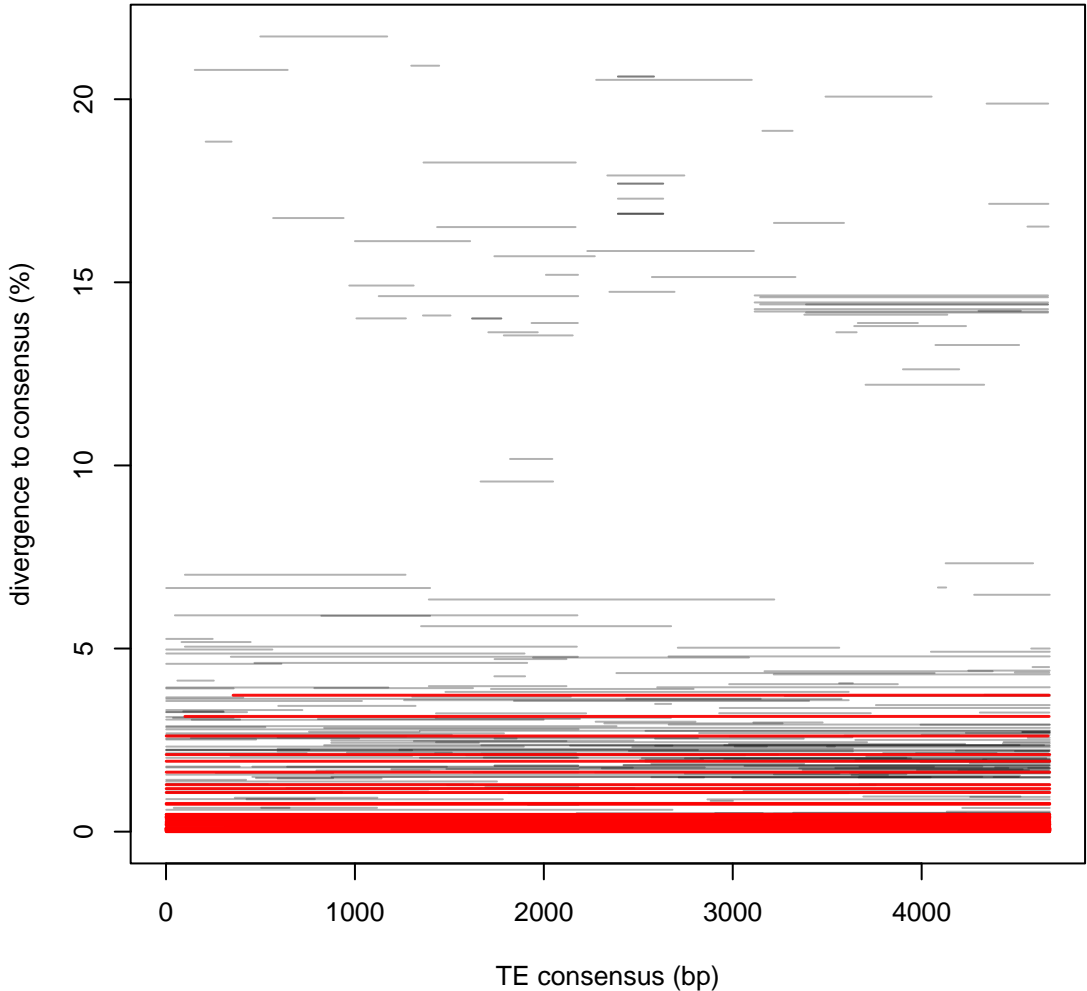
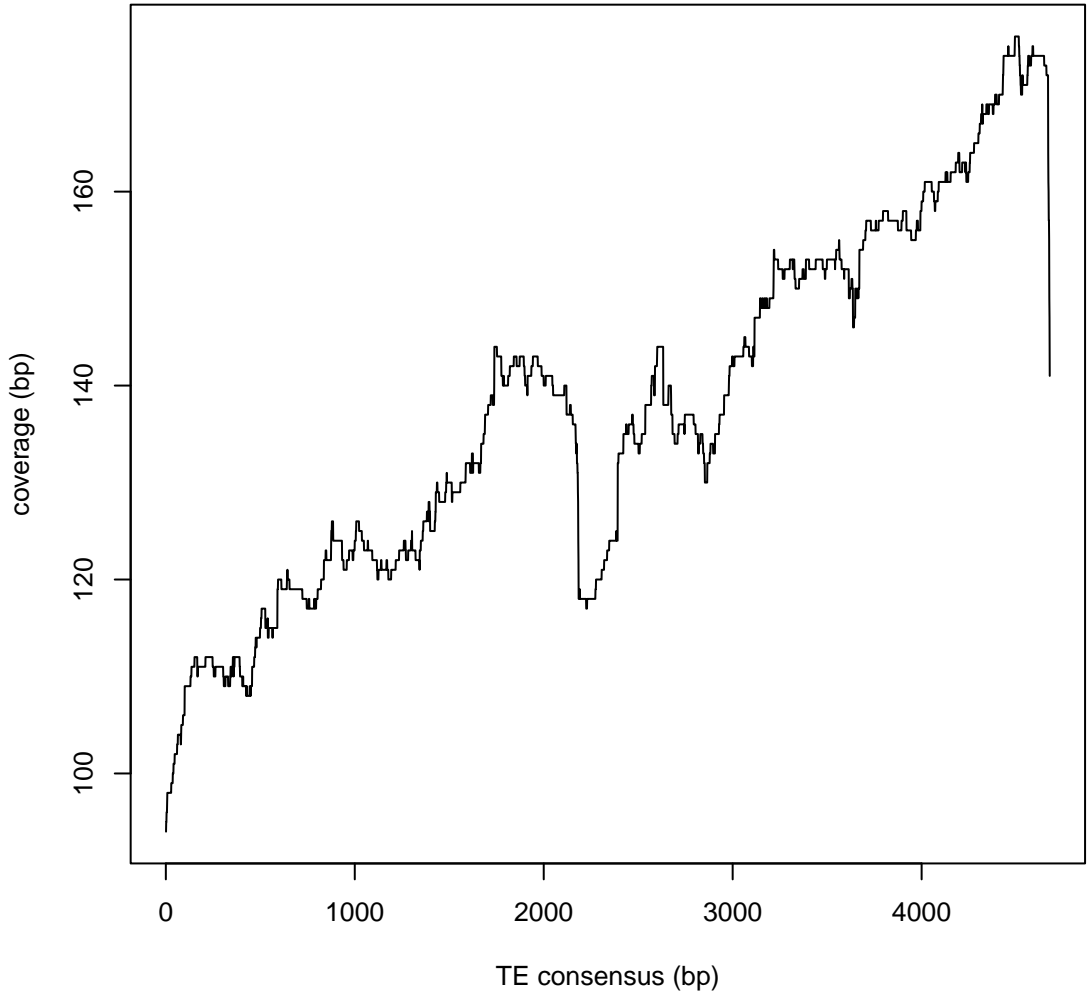


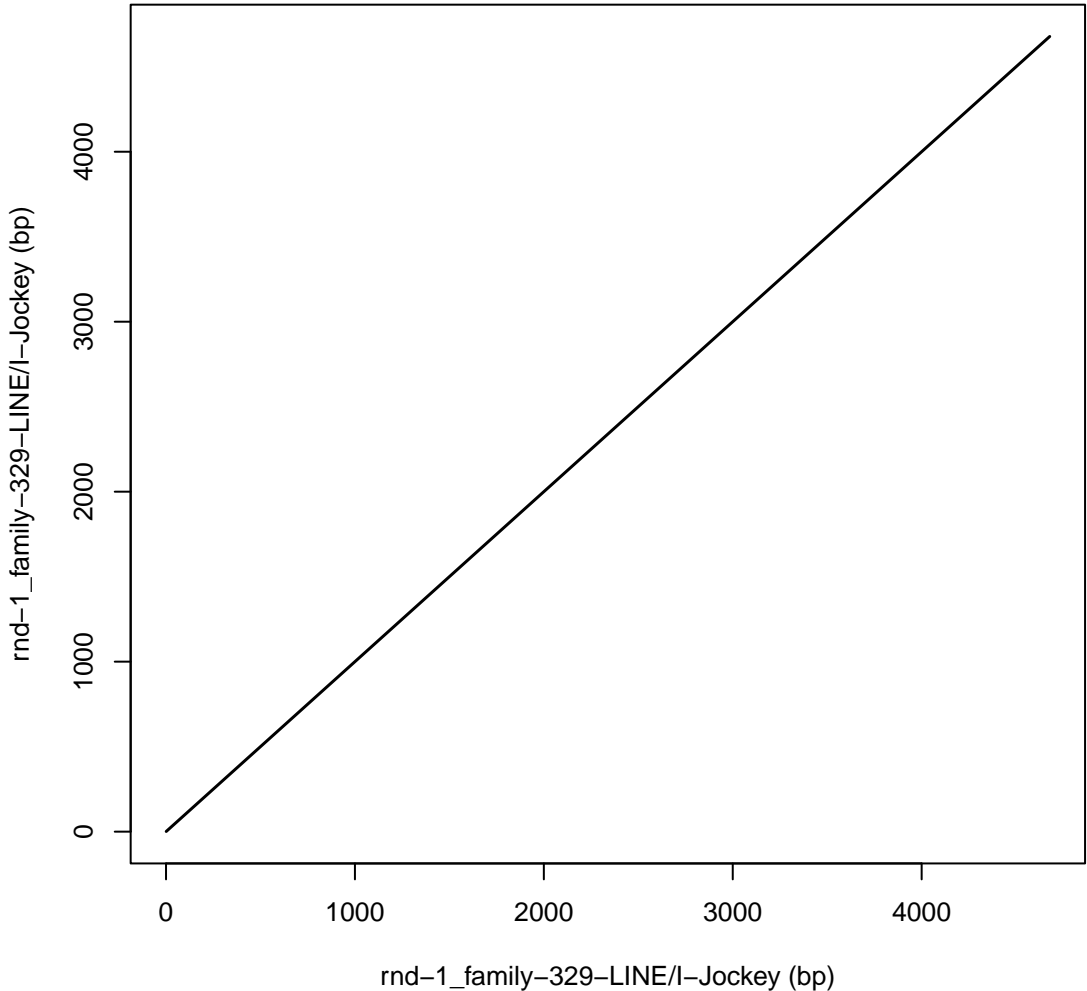
TE: rnd-1_family-329-LINE/I-Jockey
consensus size: 4678bp; fragments: 392; full length: 51 (>=4210.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

