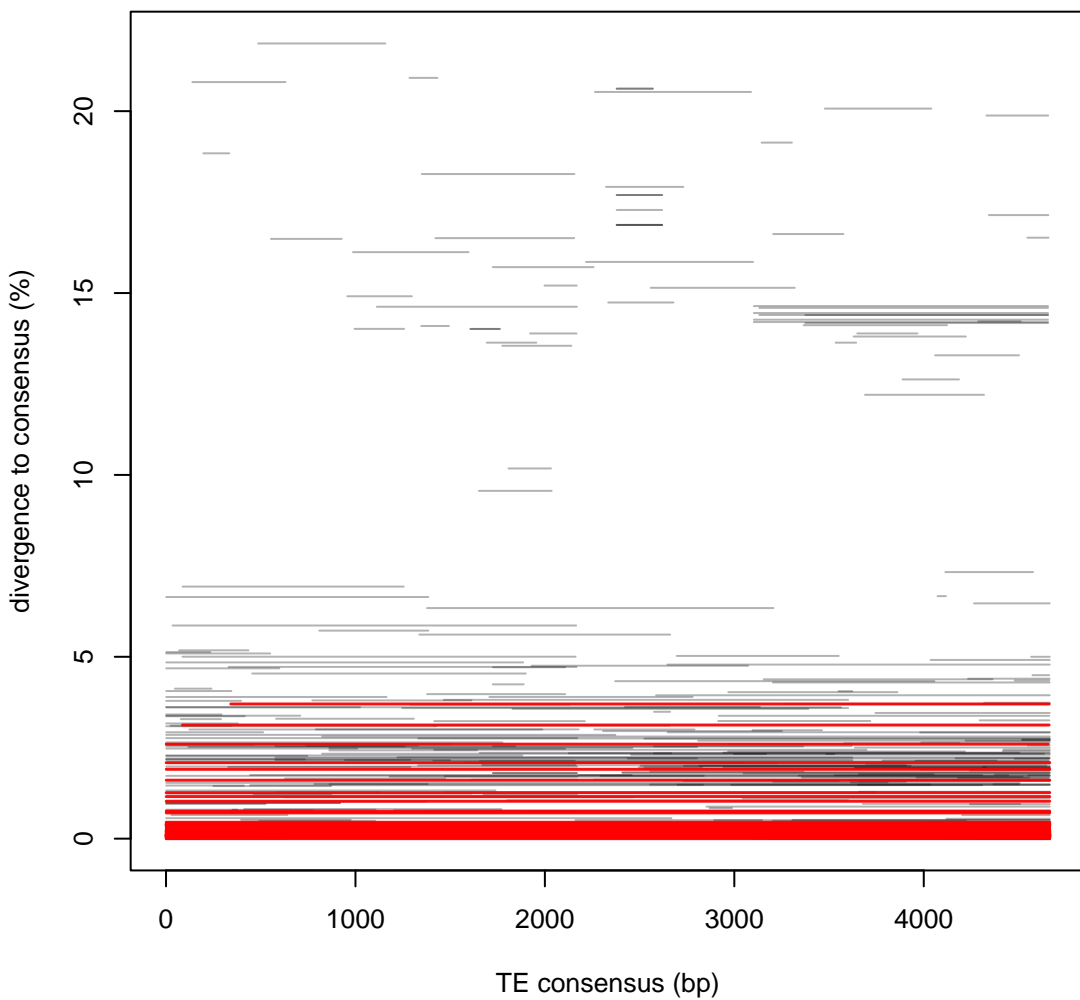
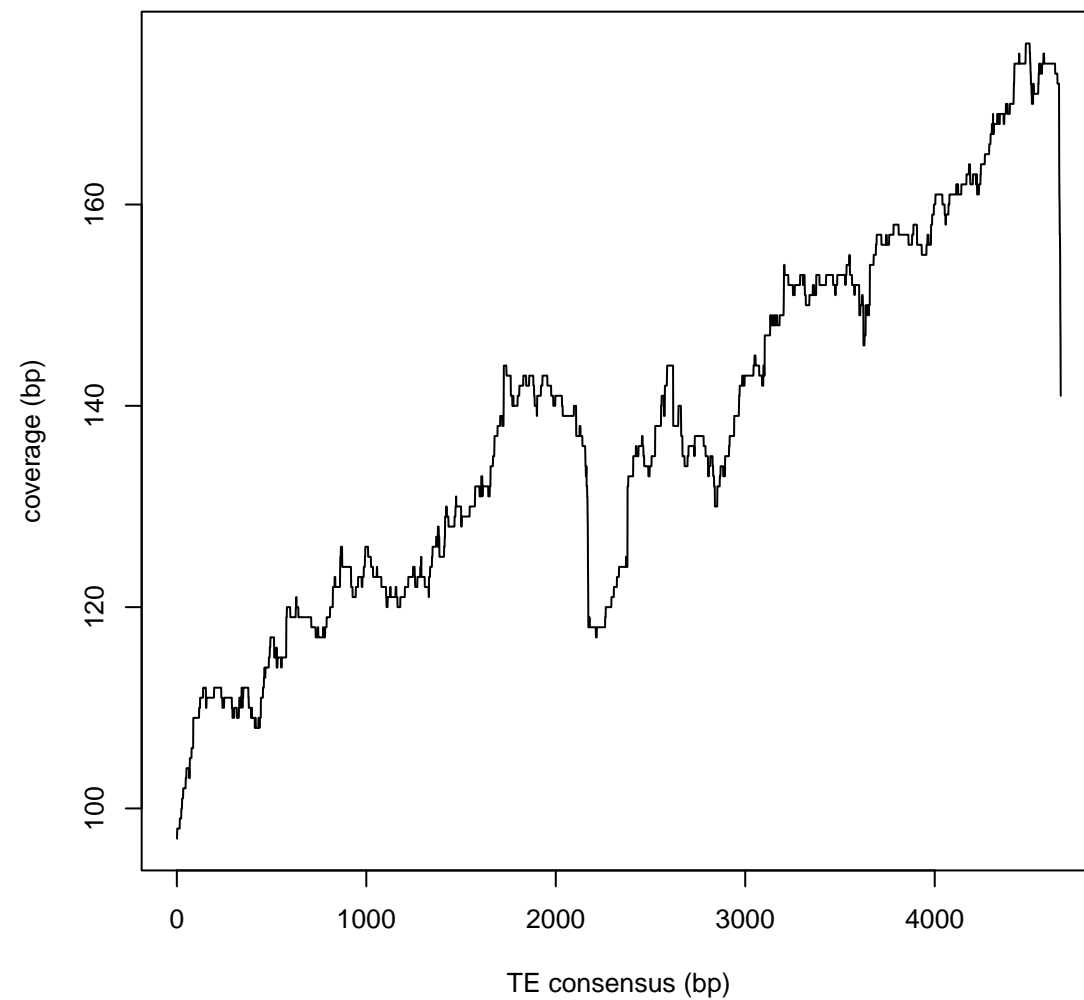


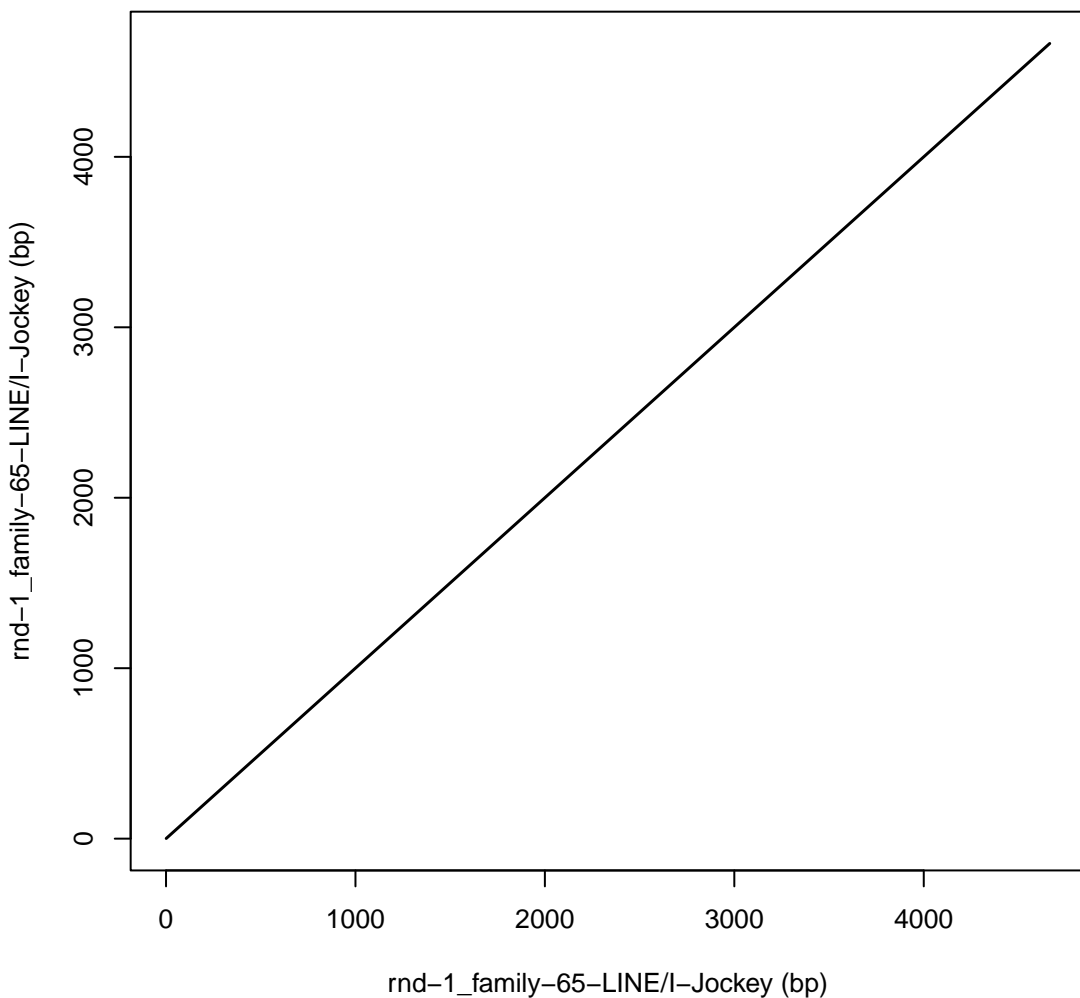
TE: rnd-1_family-65-LINE/I-Jockey
consensus size: 4665bp; fragments: 392; full length: 52 (>=4198.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

