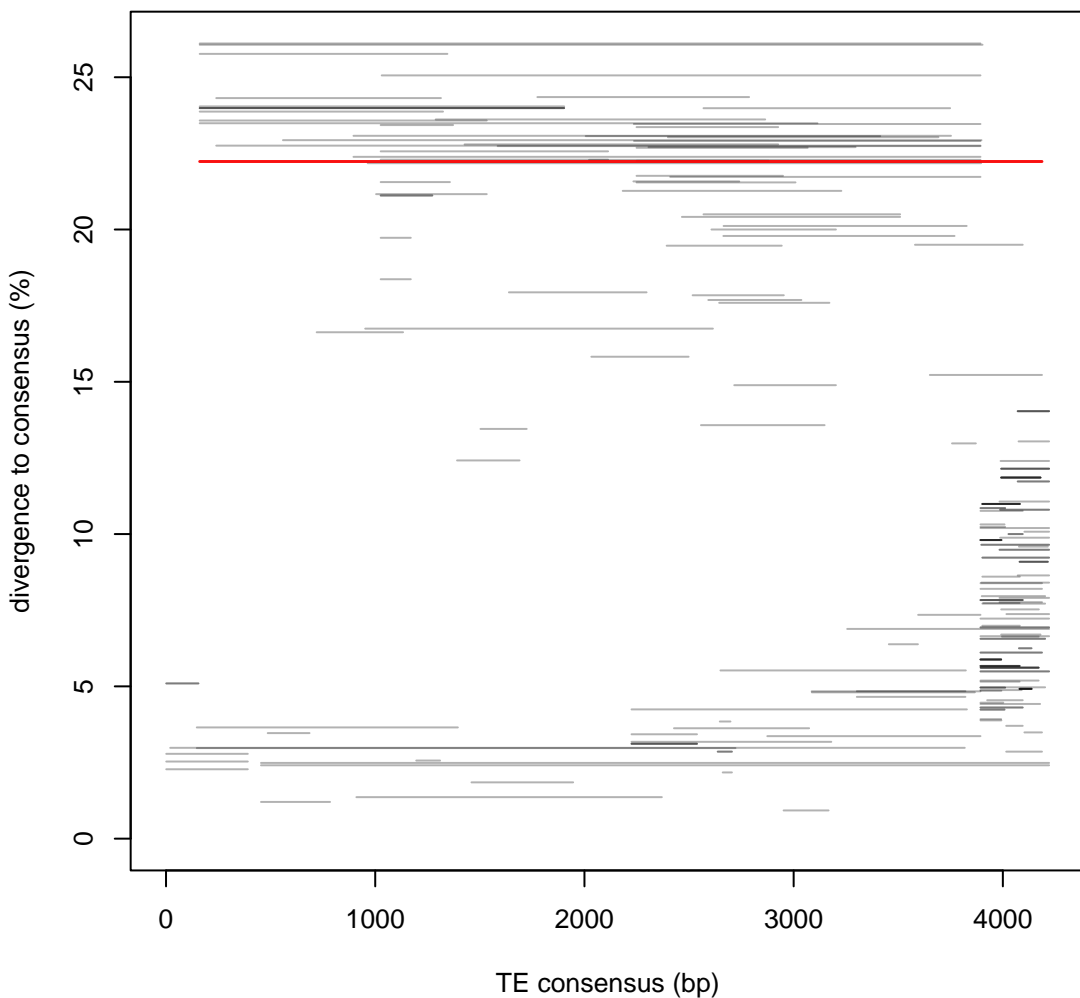
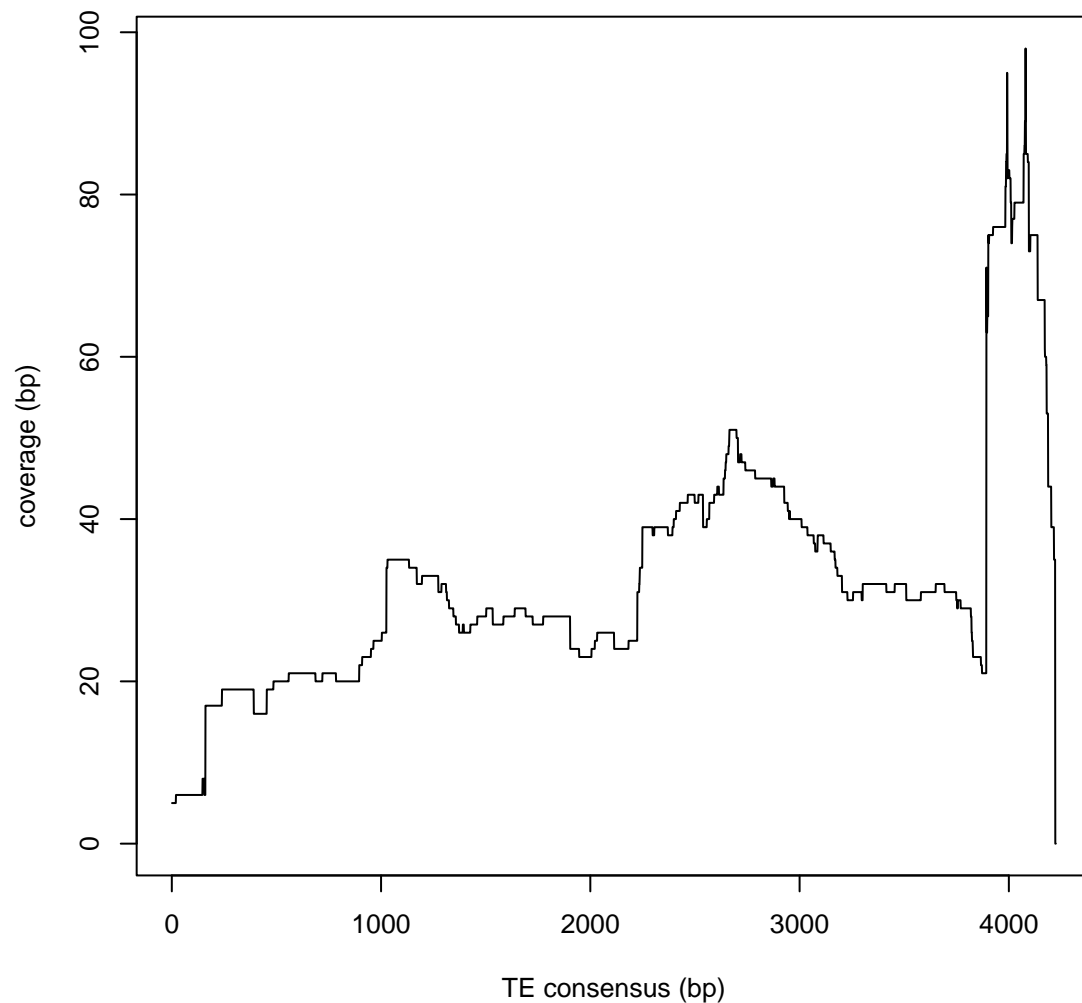


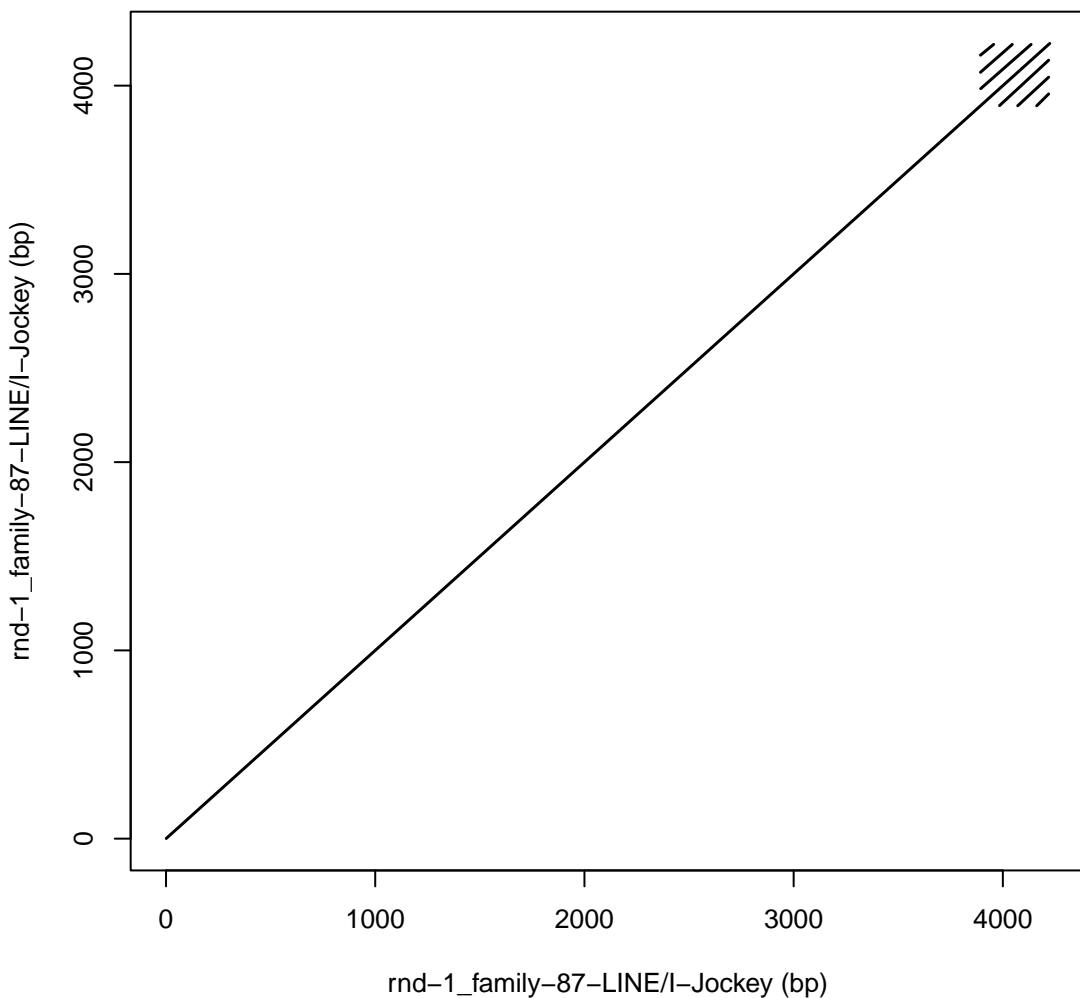
TE: rnd-1_family-87-LINE/I-Jockey
consensus size: 4224bp; fragments: 218; full length: 1 (>=3801.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

