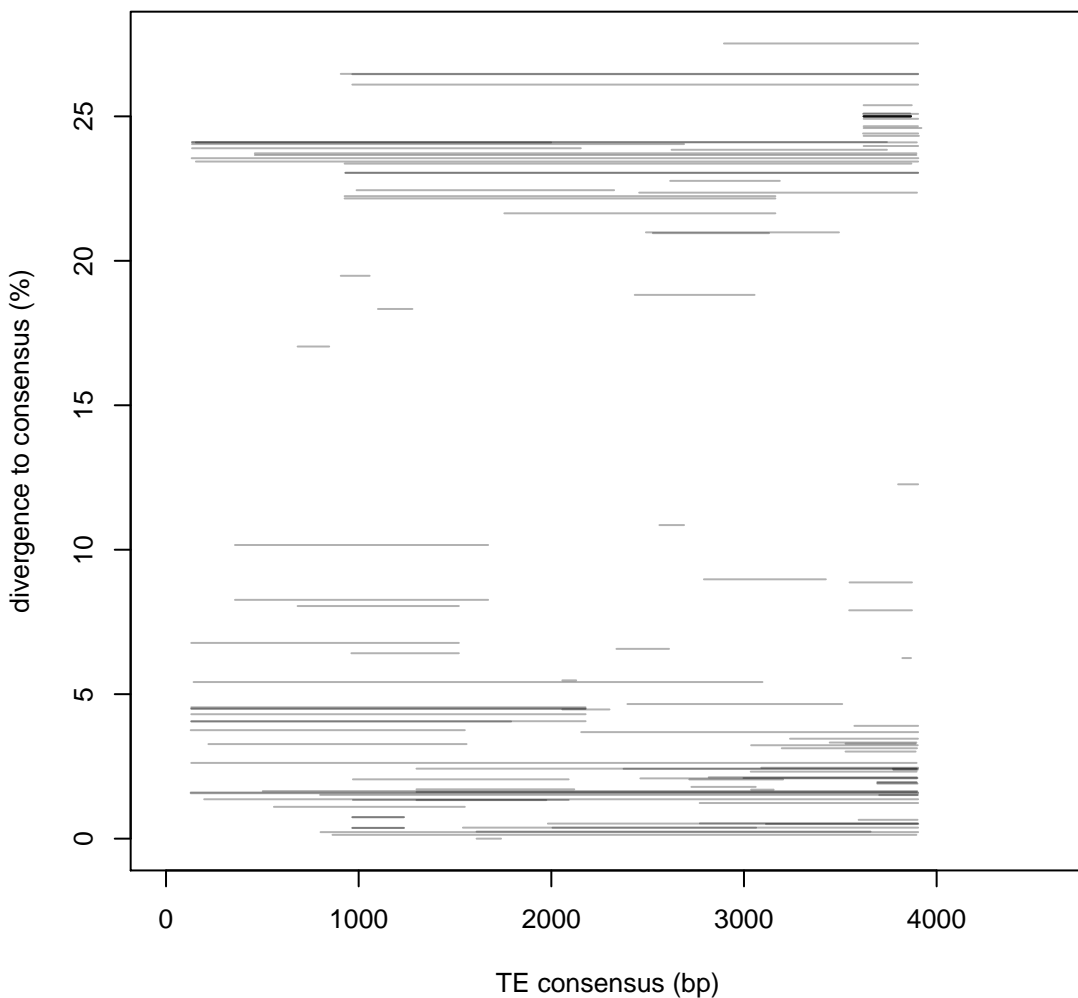
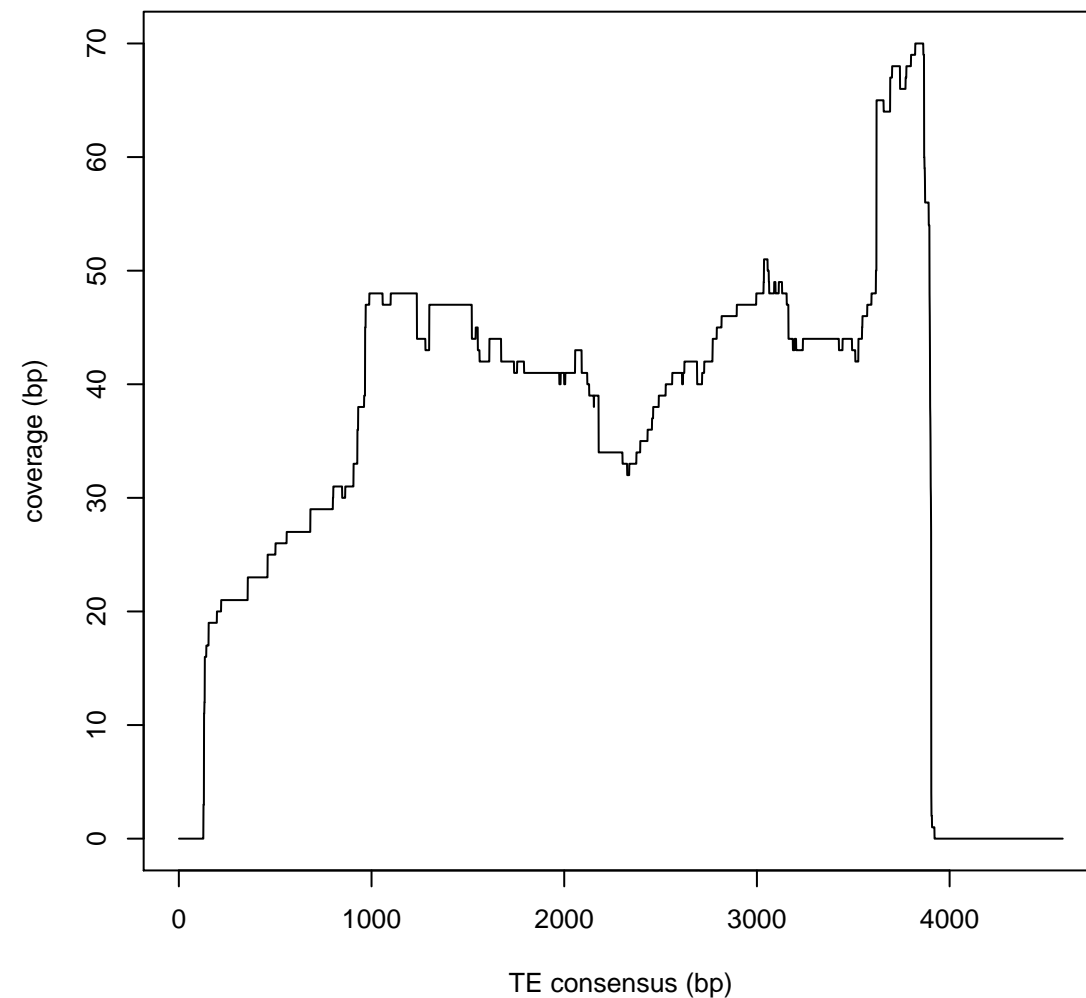


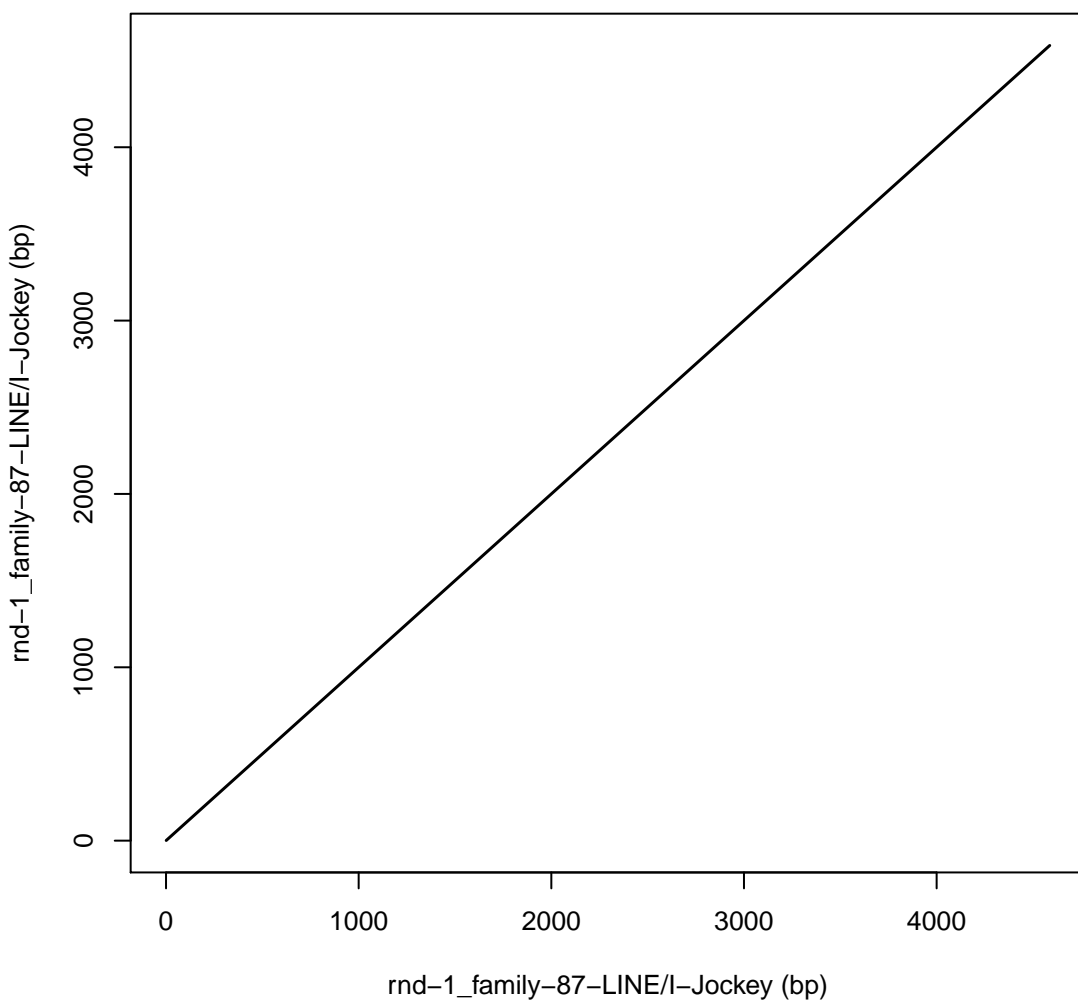
TE: rnd-1_family-87-LINE/I-Jockey
consensus size: 4587bp; fragments: 121; full length: 0 (>=4128.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

