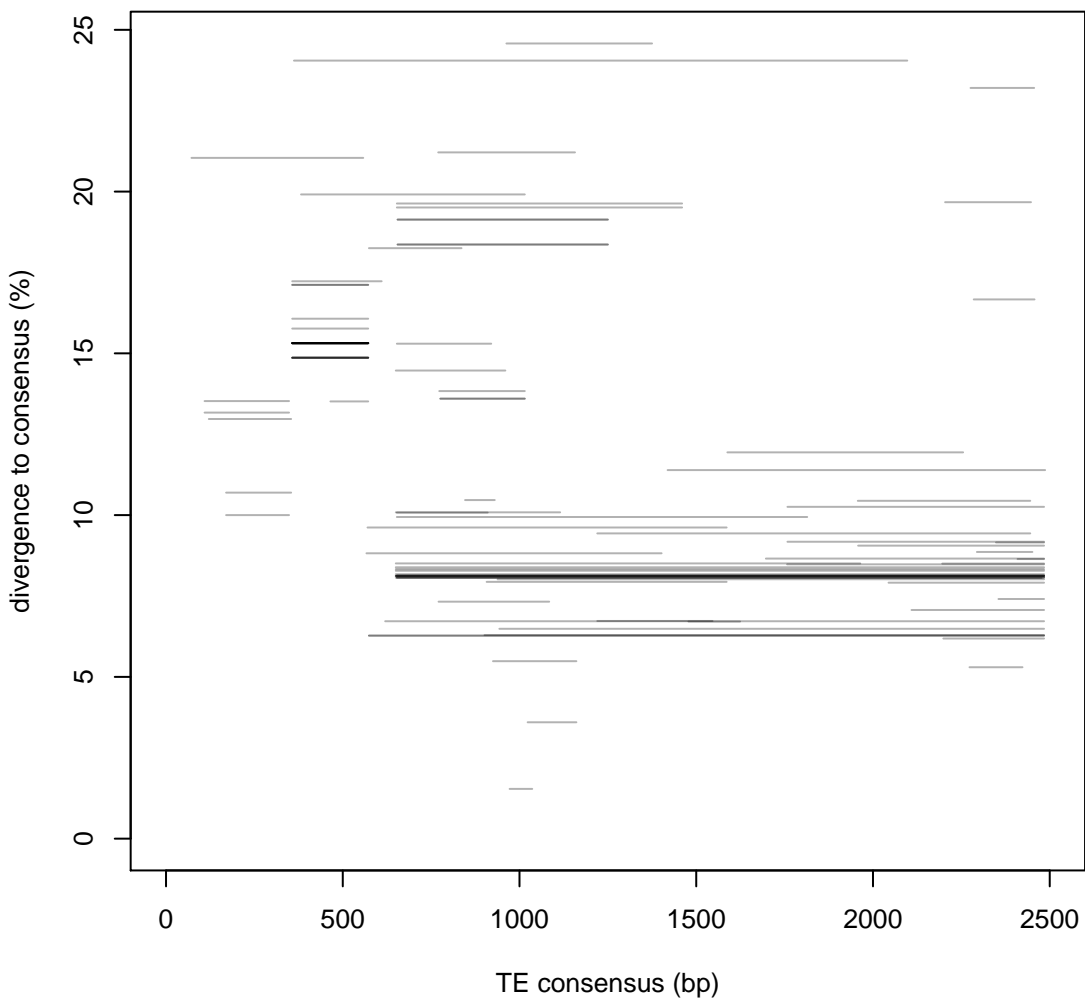
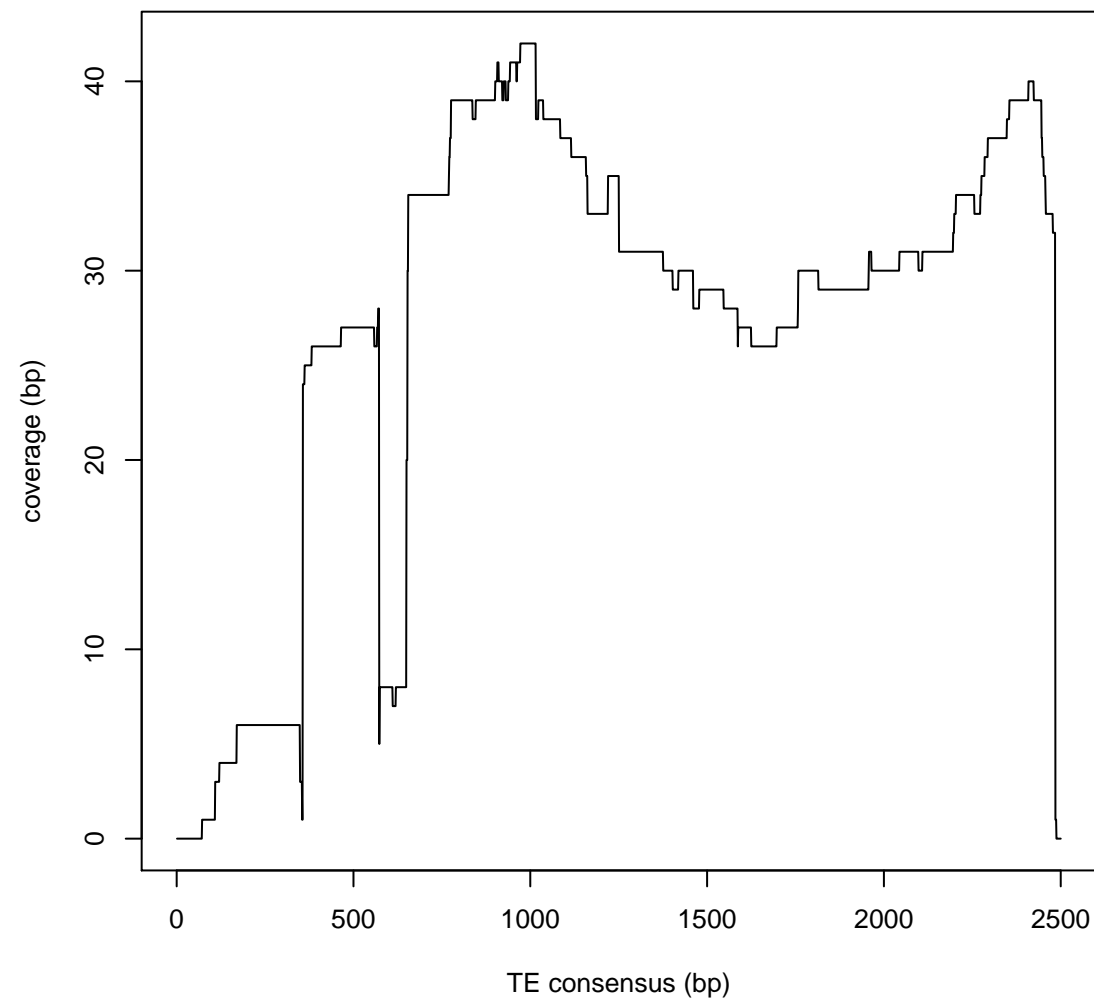


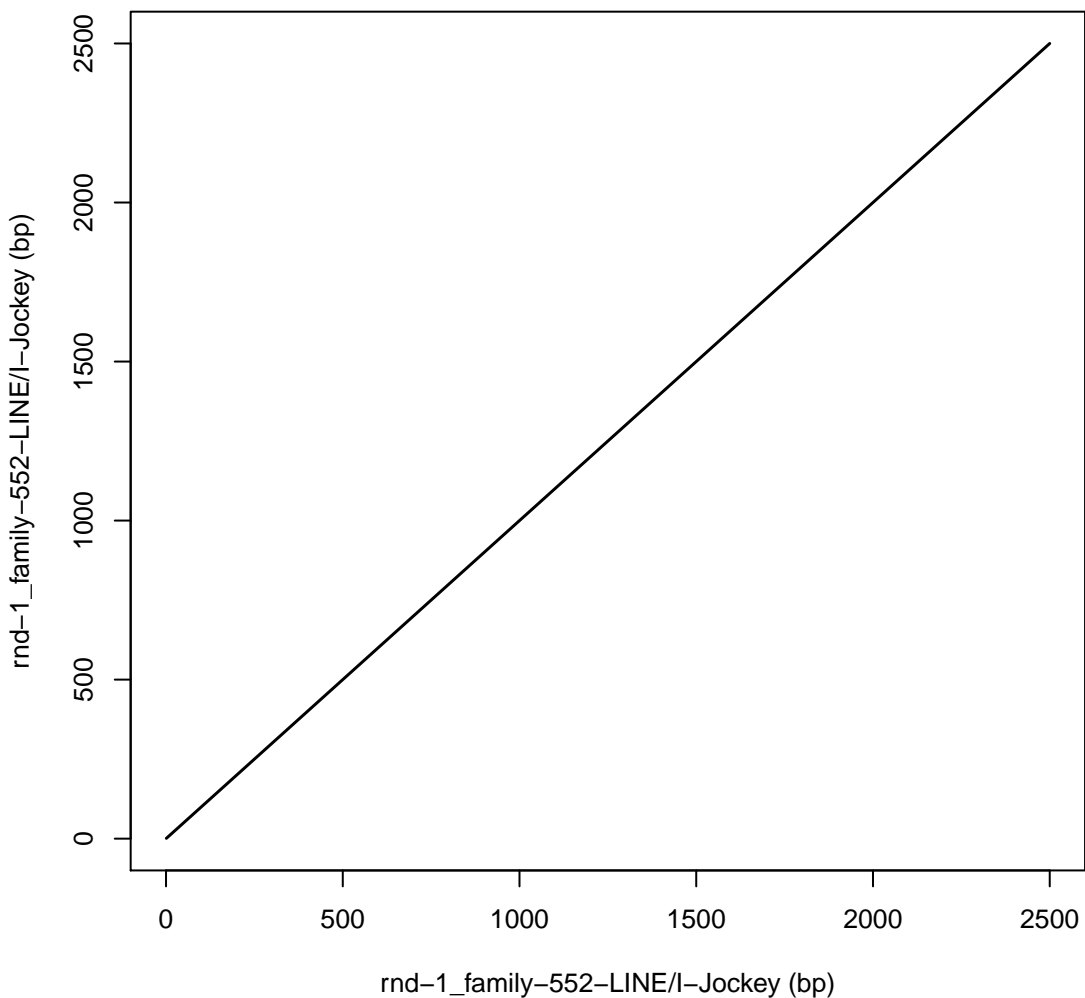
TE: rnd-1_family-552-LINE/I-Jockey
consensus size: 2500bp; fragments: 101; full length: 0 (>=2250bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

