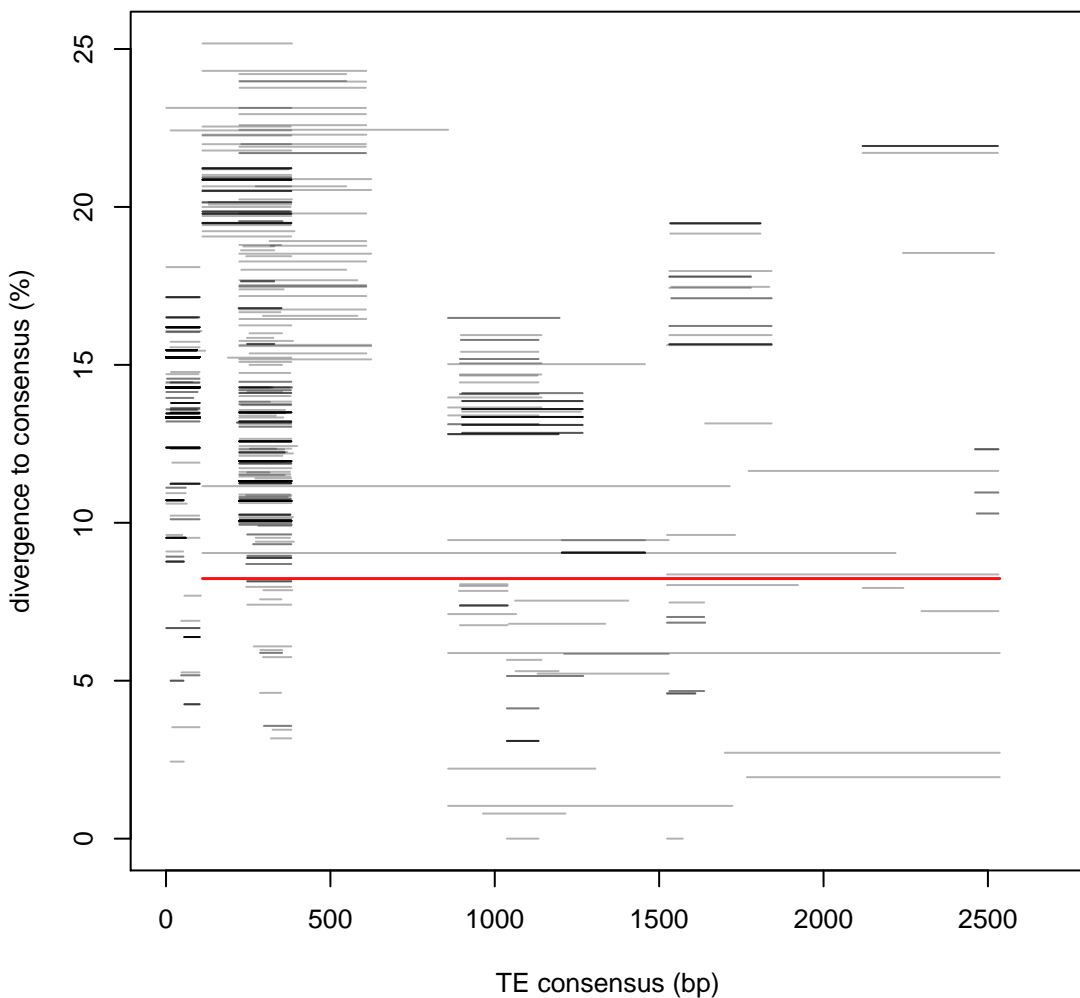
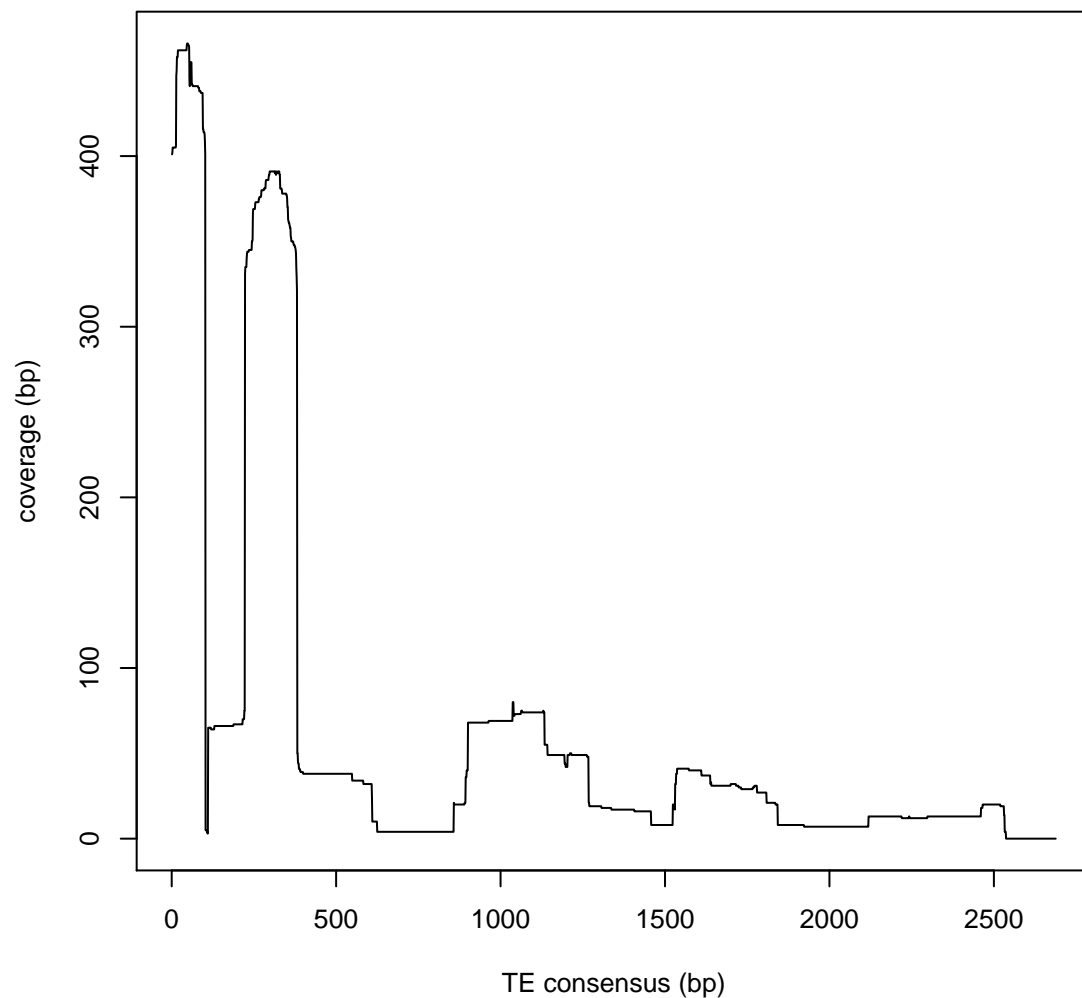


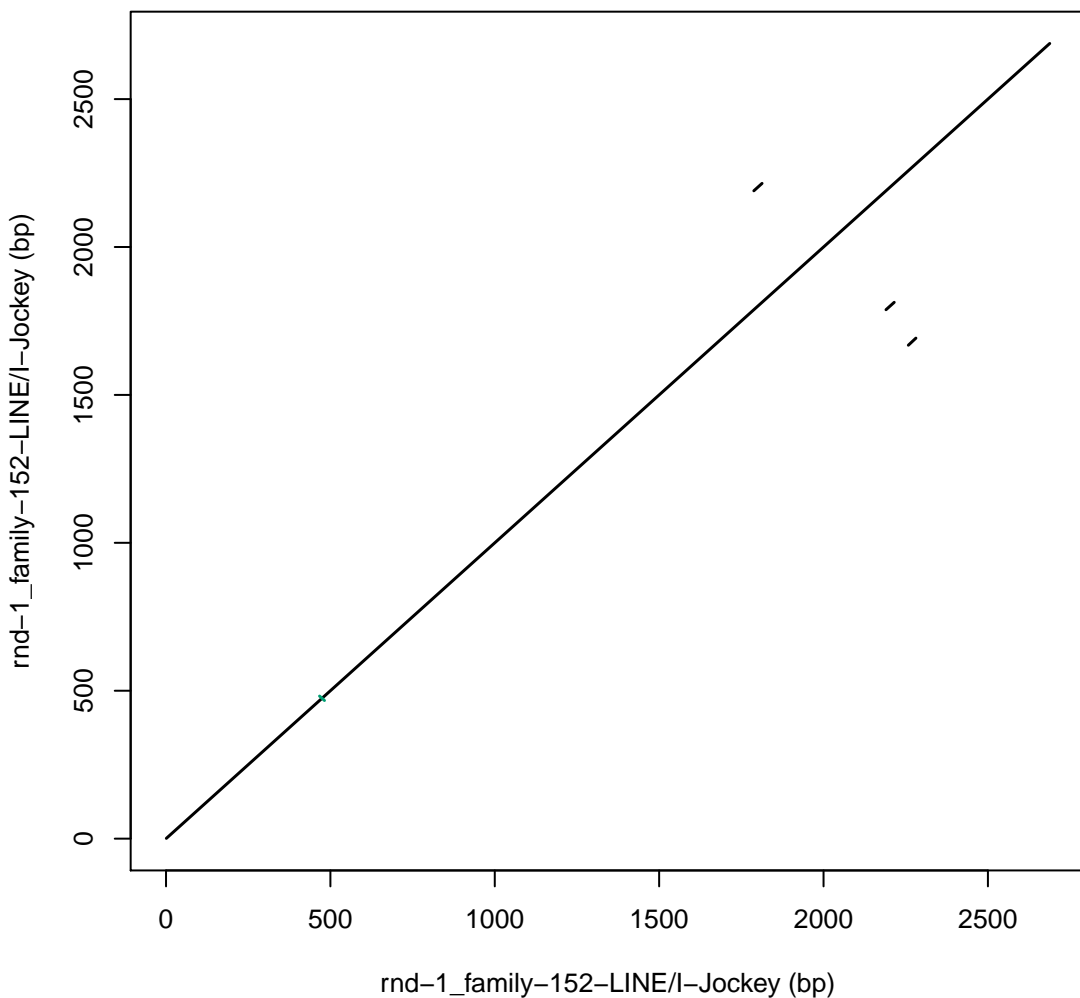
TE: rnd-1_family-152-LINE/l-Jockey
consensus size: 2688bp; fragments: 1017; full length: 1 (>=2419.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

