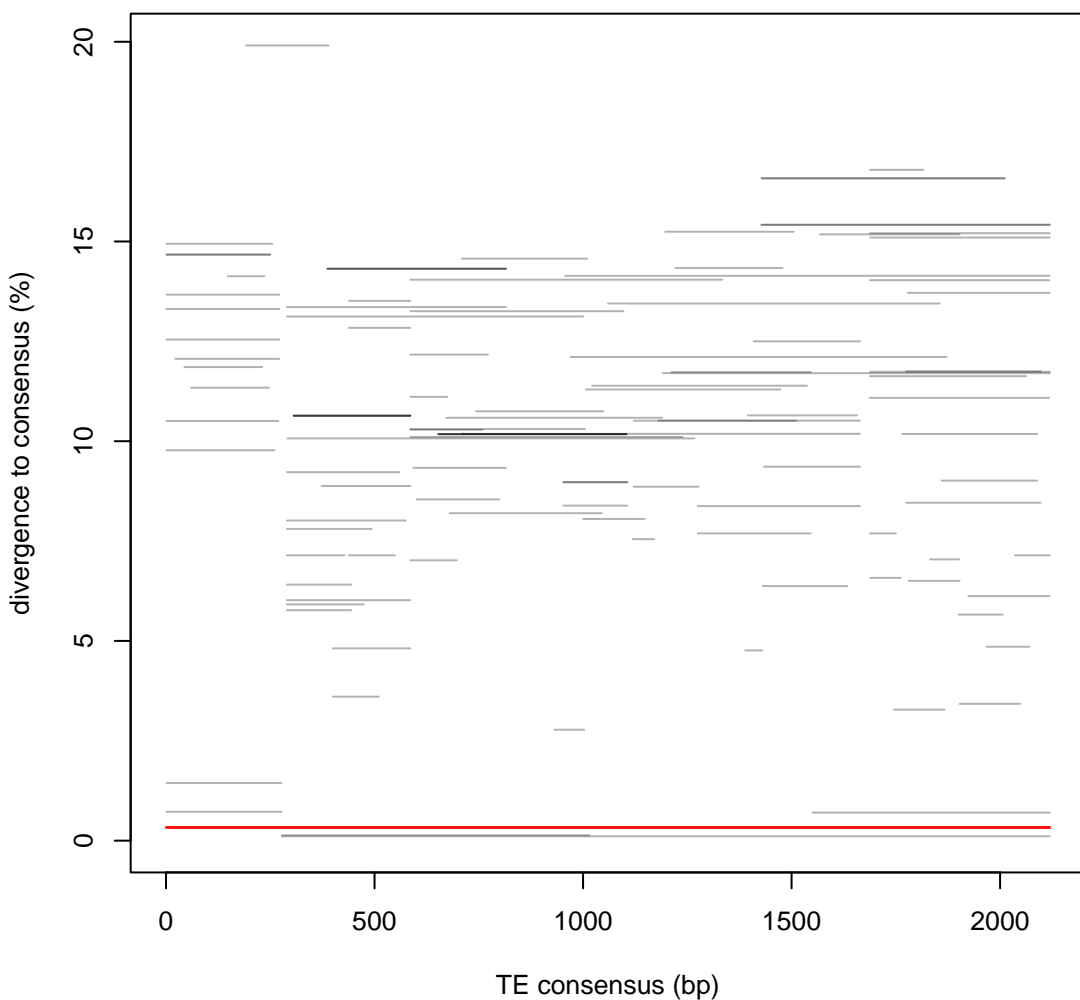
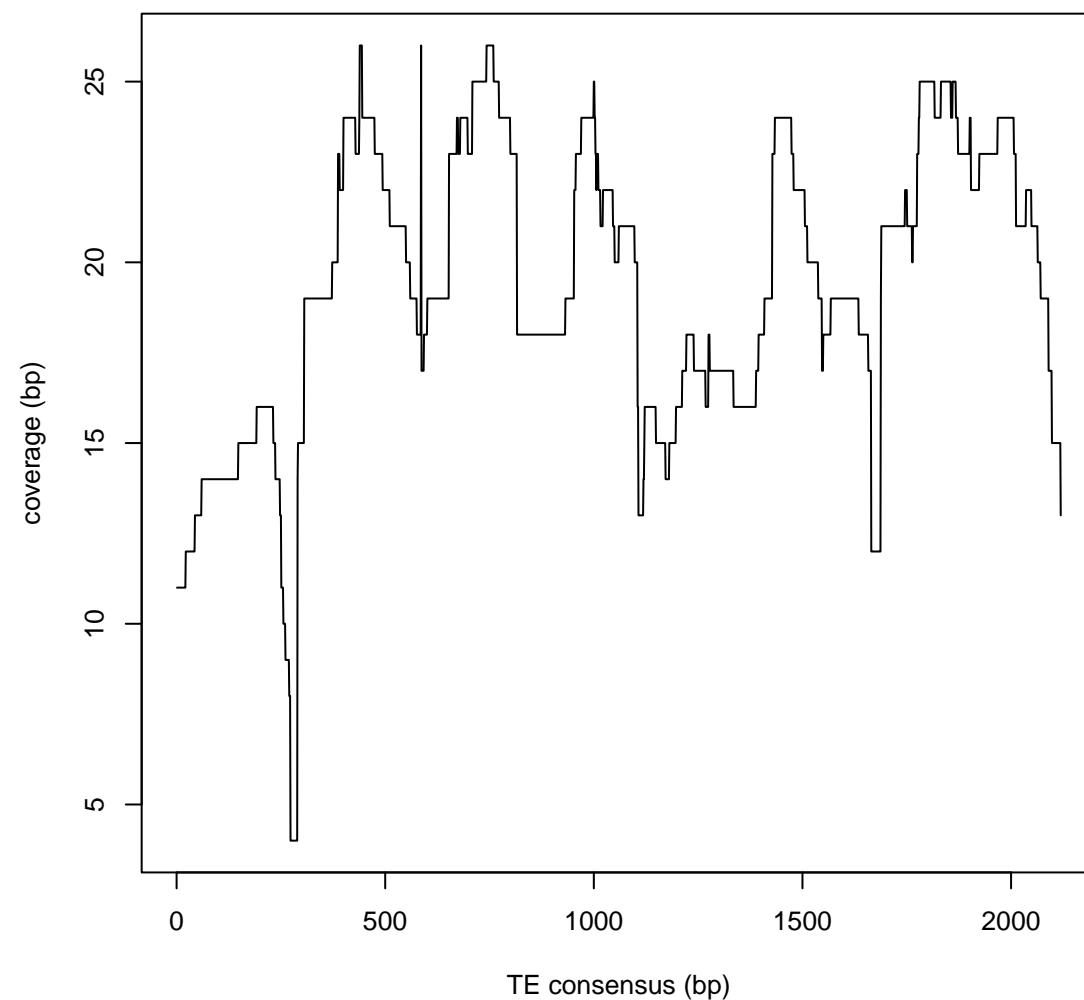


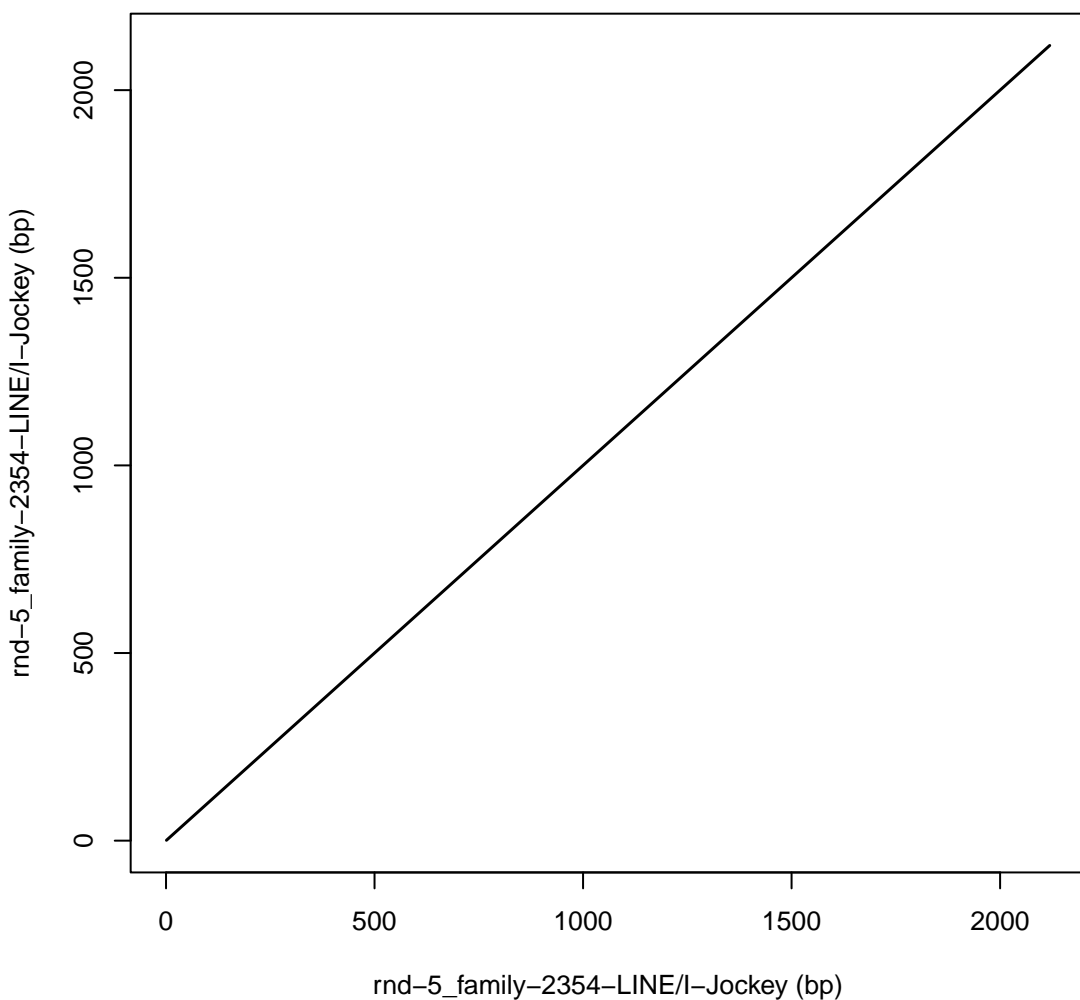
TE: rnd-5\_family-2354-LINE/I-Jockey  
consensus size: 2119bp; fragments: 114; full length: 1 (>=1907.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

