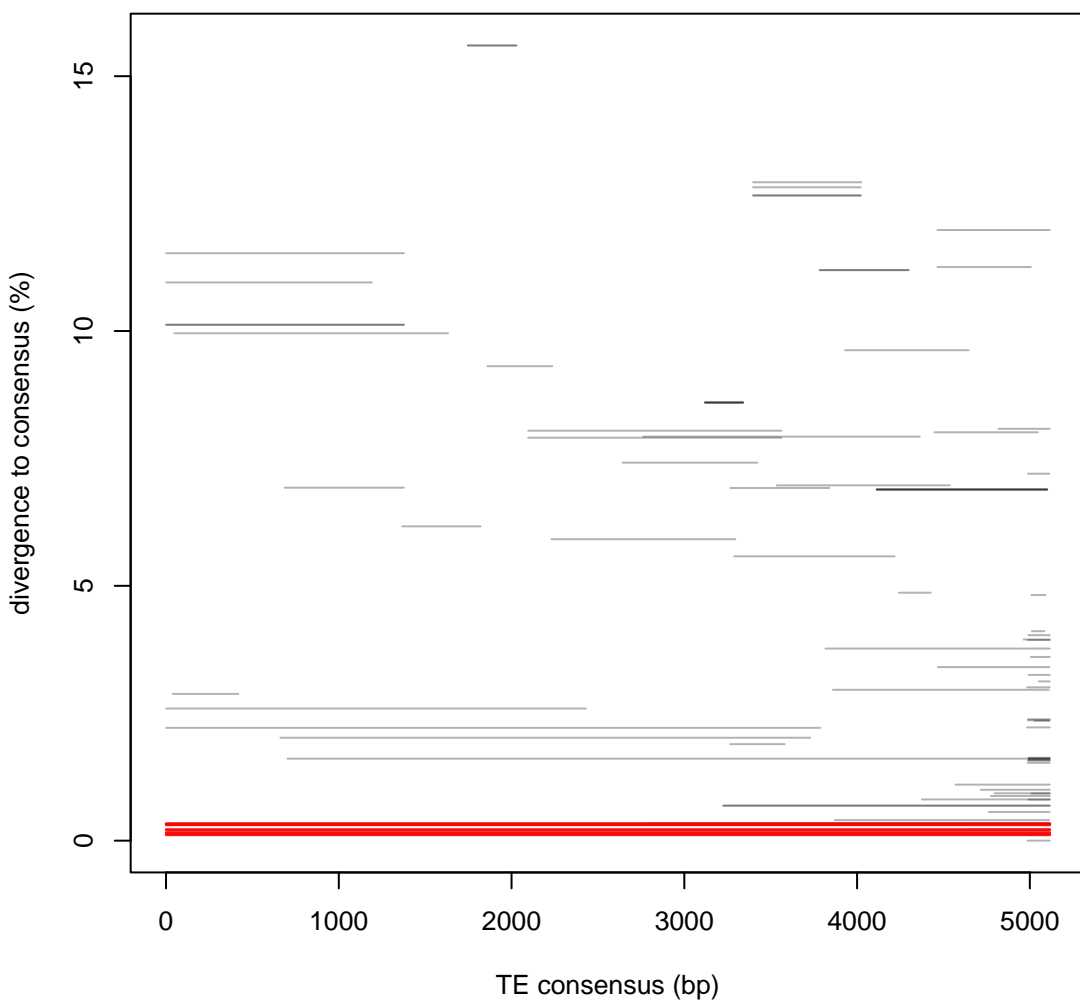
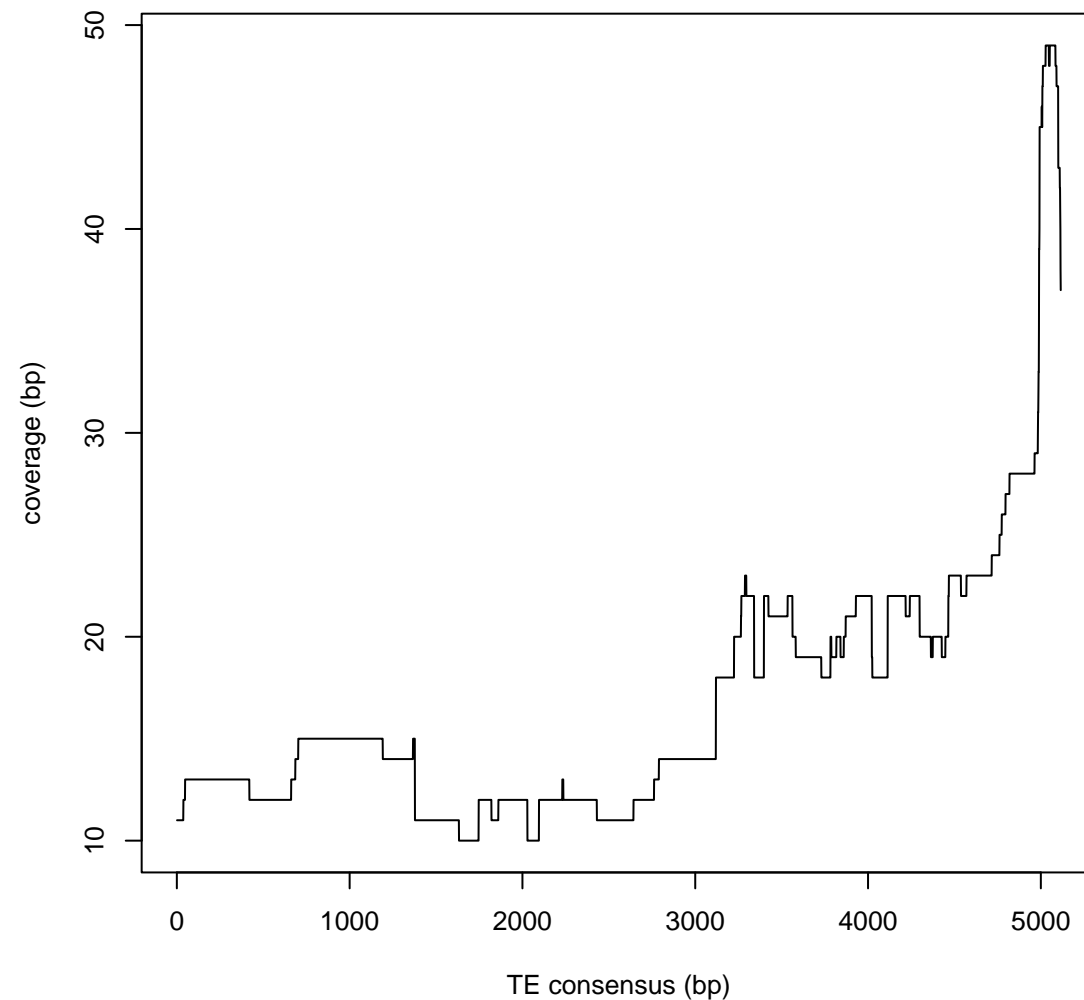


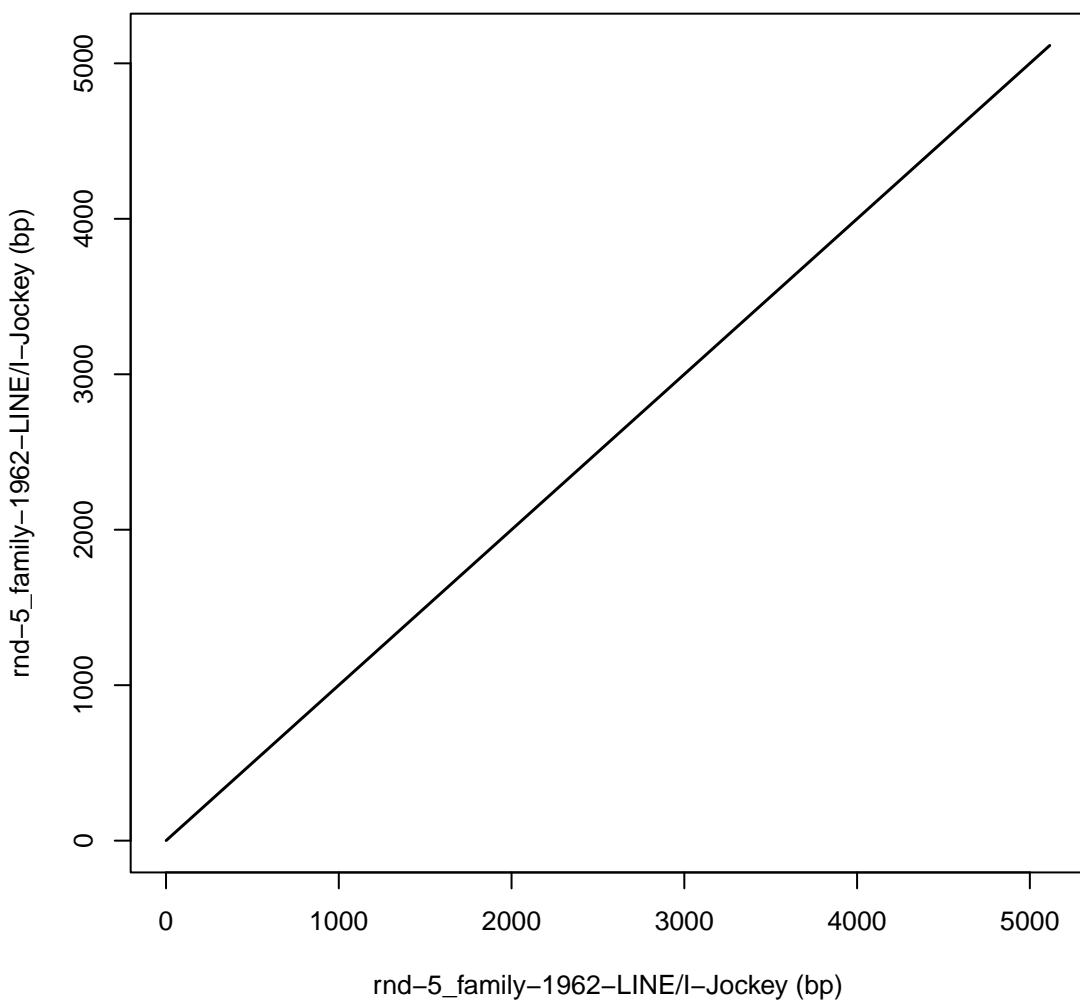
TE: rnd-5\_family-1962-LINE/I-Jockey  
consensus size: 5115bp; fragments: 86; full length: 5 (>=4603.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

