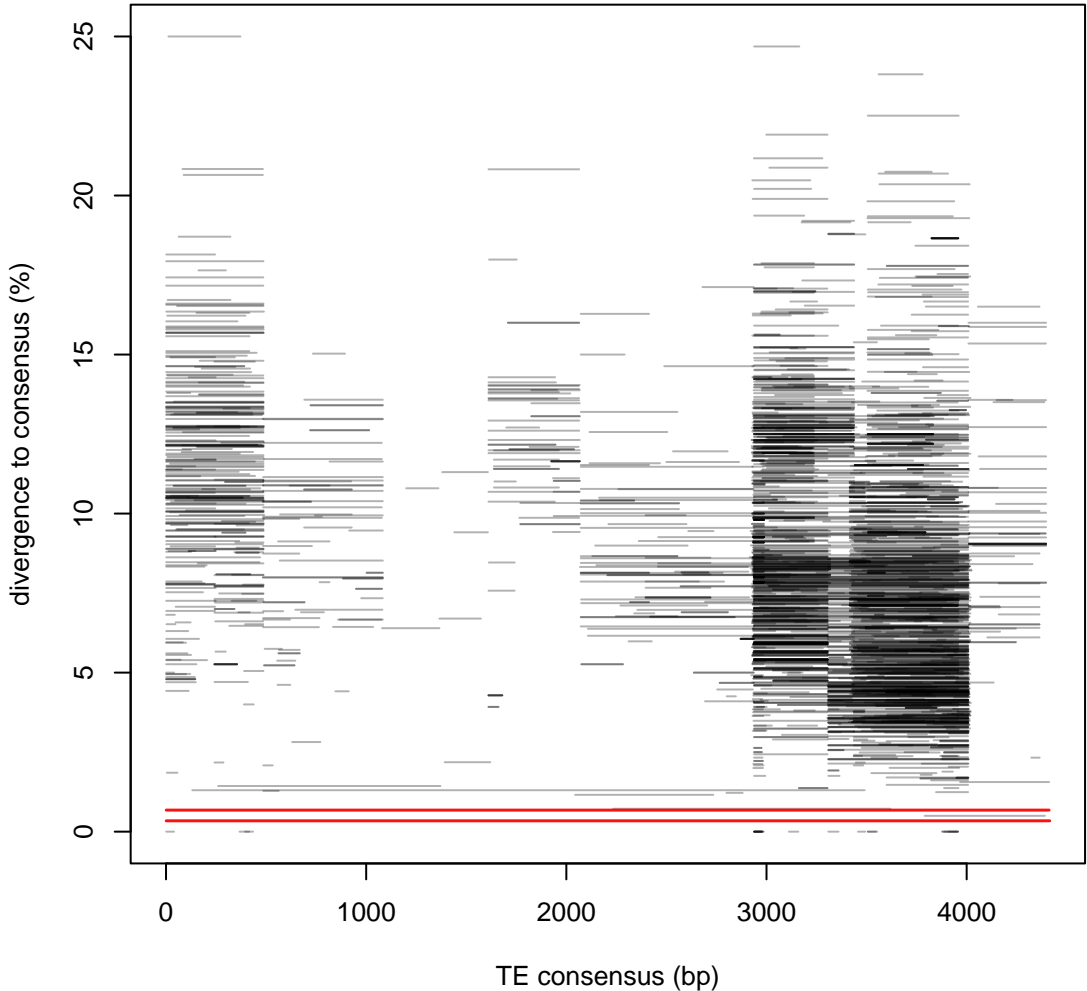
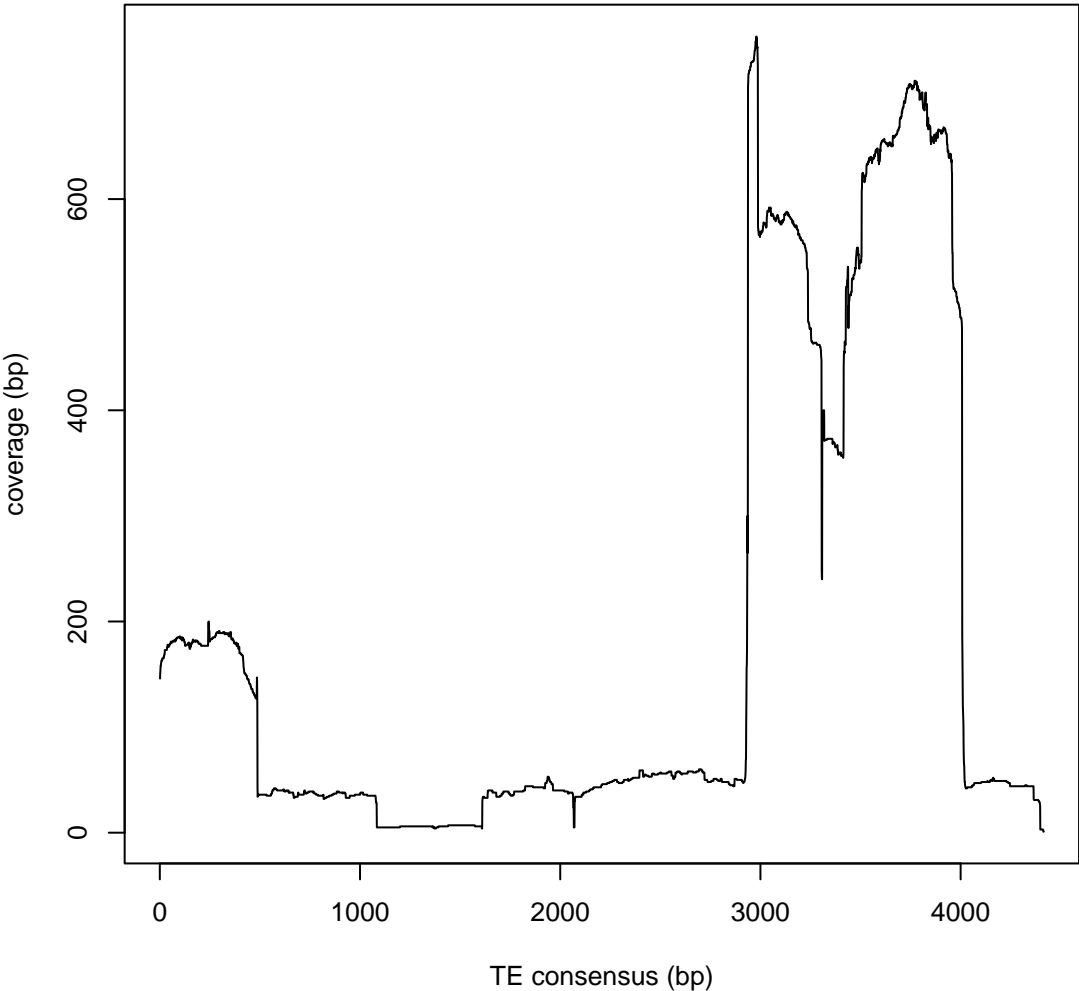


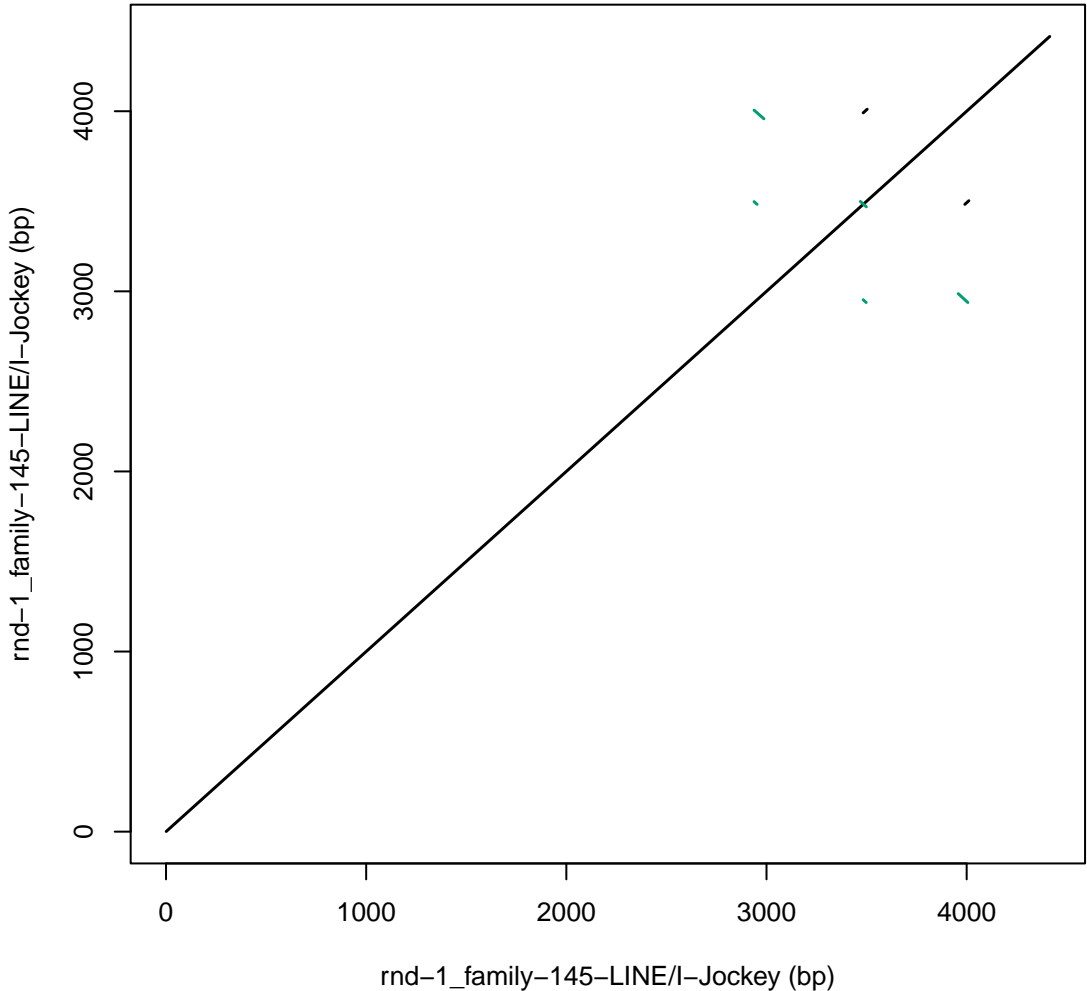
TE: rnd-1\_family-145-LINE/I-Jockey  
consensus size: 4415bp; fragments: 2343; full length: 2 (>=3973.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

