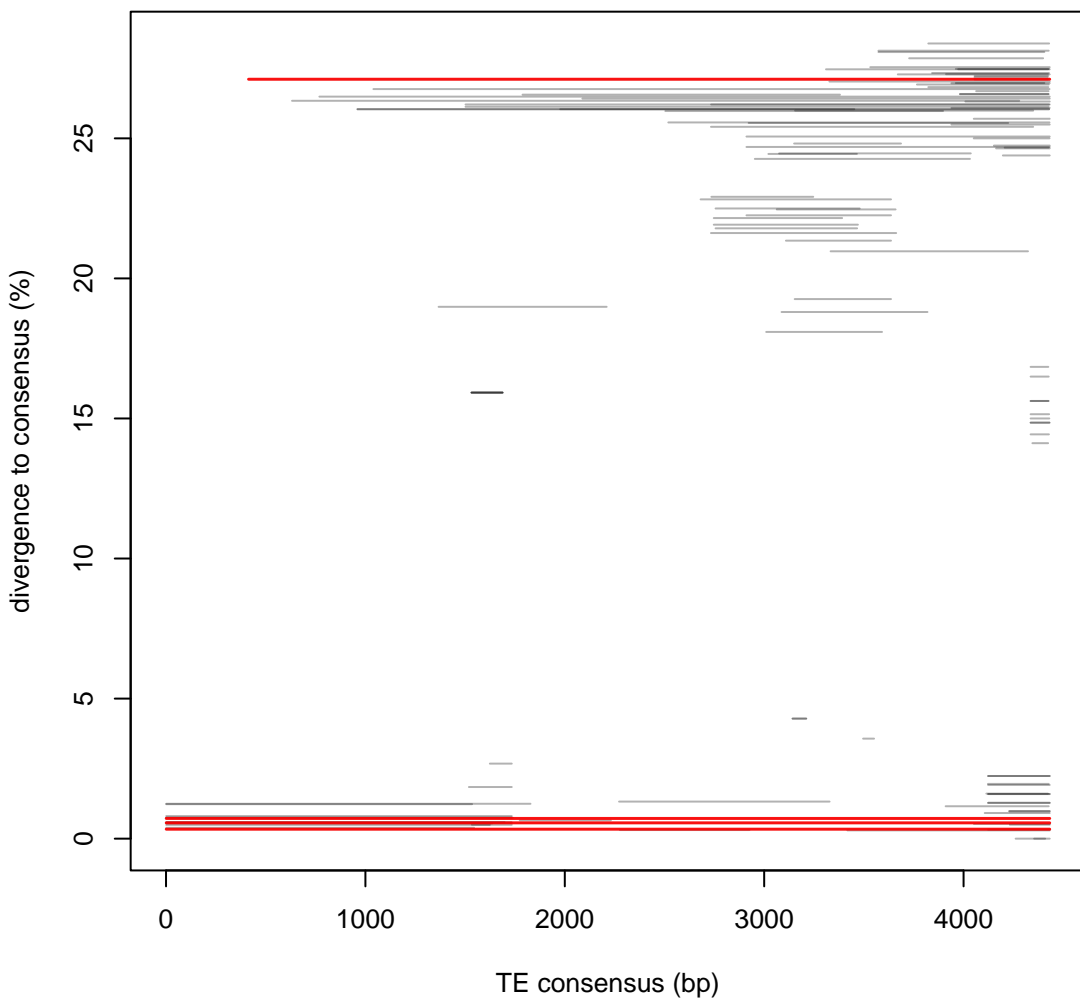
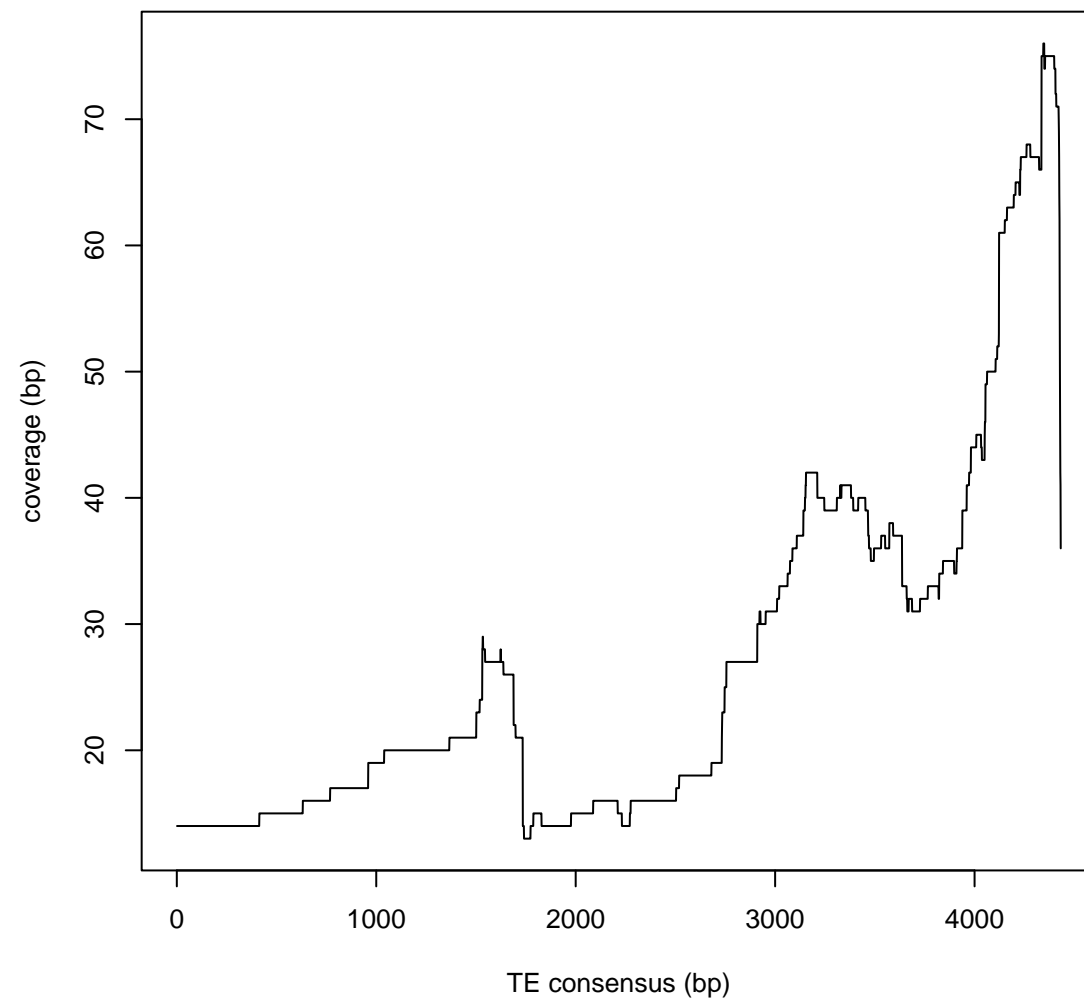


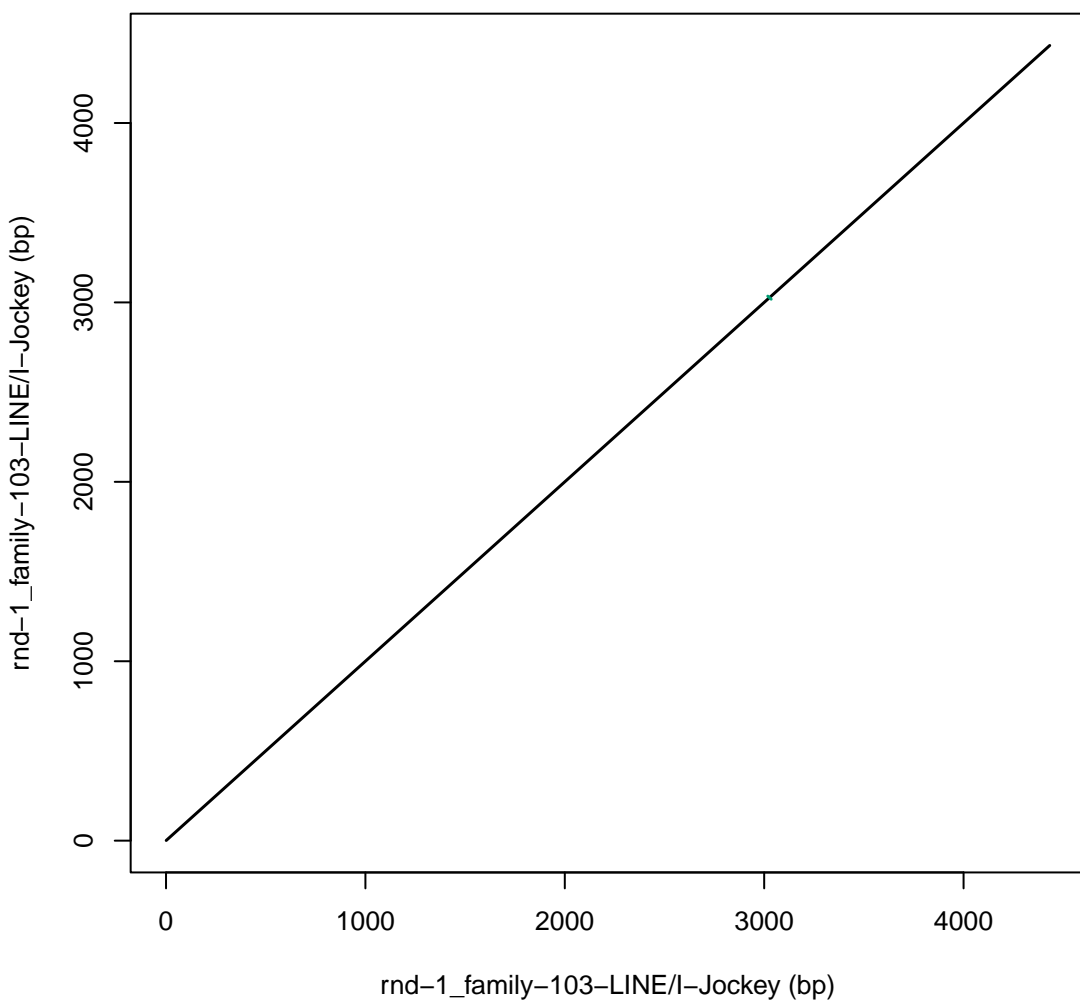
TE: rnd-1\_family-103-LINE/I-Jockey  
consensus size: 4432bp; fragments: 125; full length: 4 (>=3988.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

