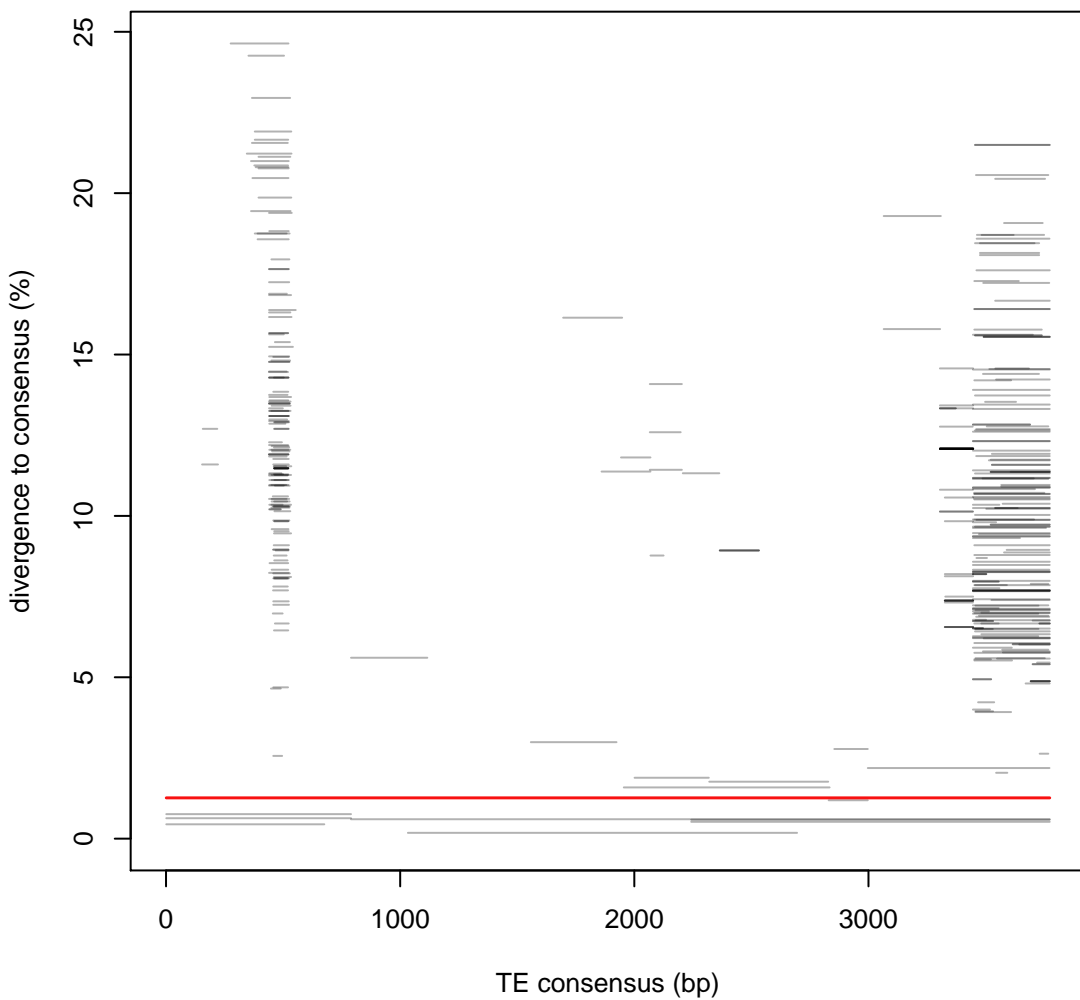
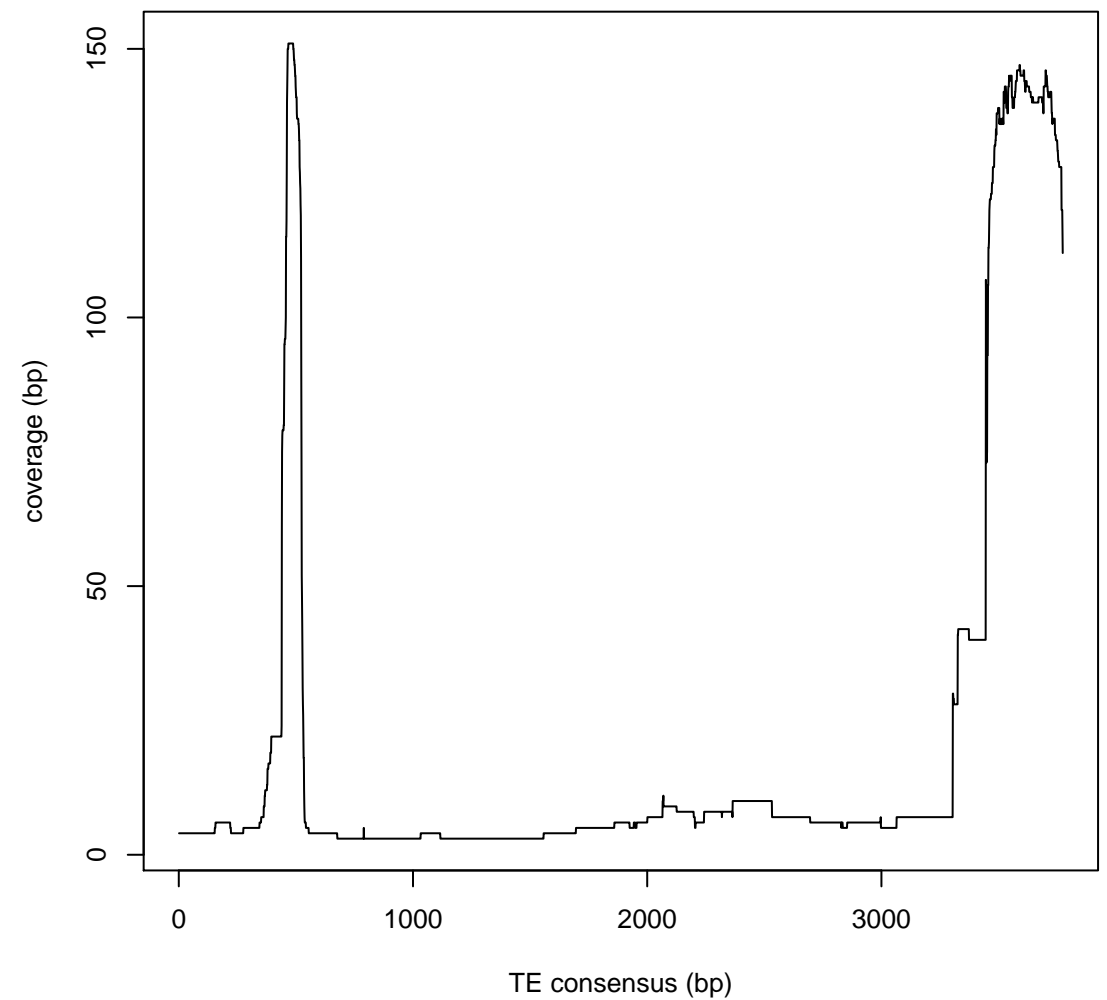


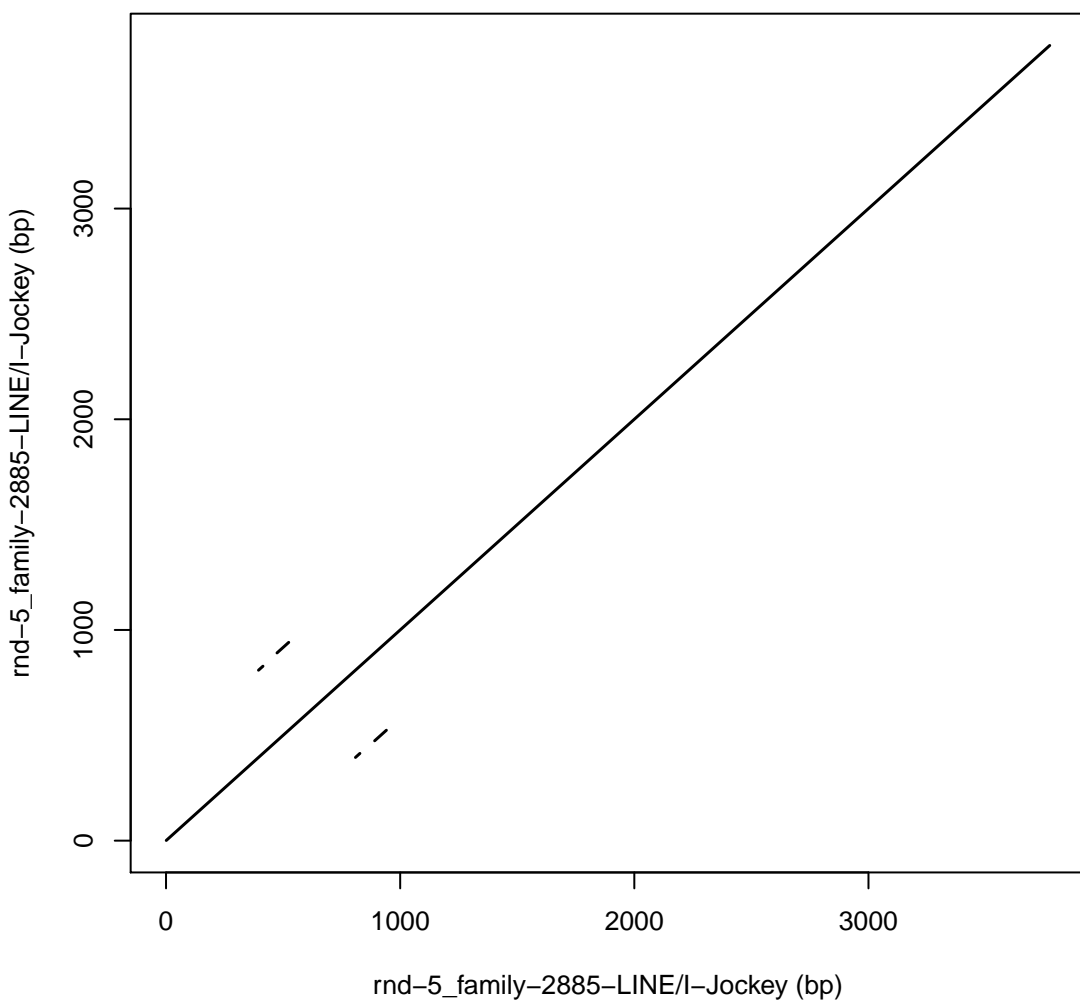
TE: rnd-5\_family-2885-LINE/I-Jockey  
consensus size: 3774bp; fragments: 398; full length: 1 (>=3396.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

