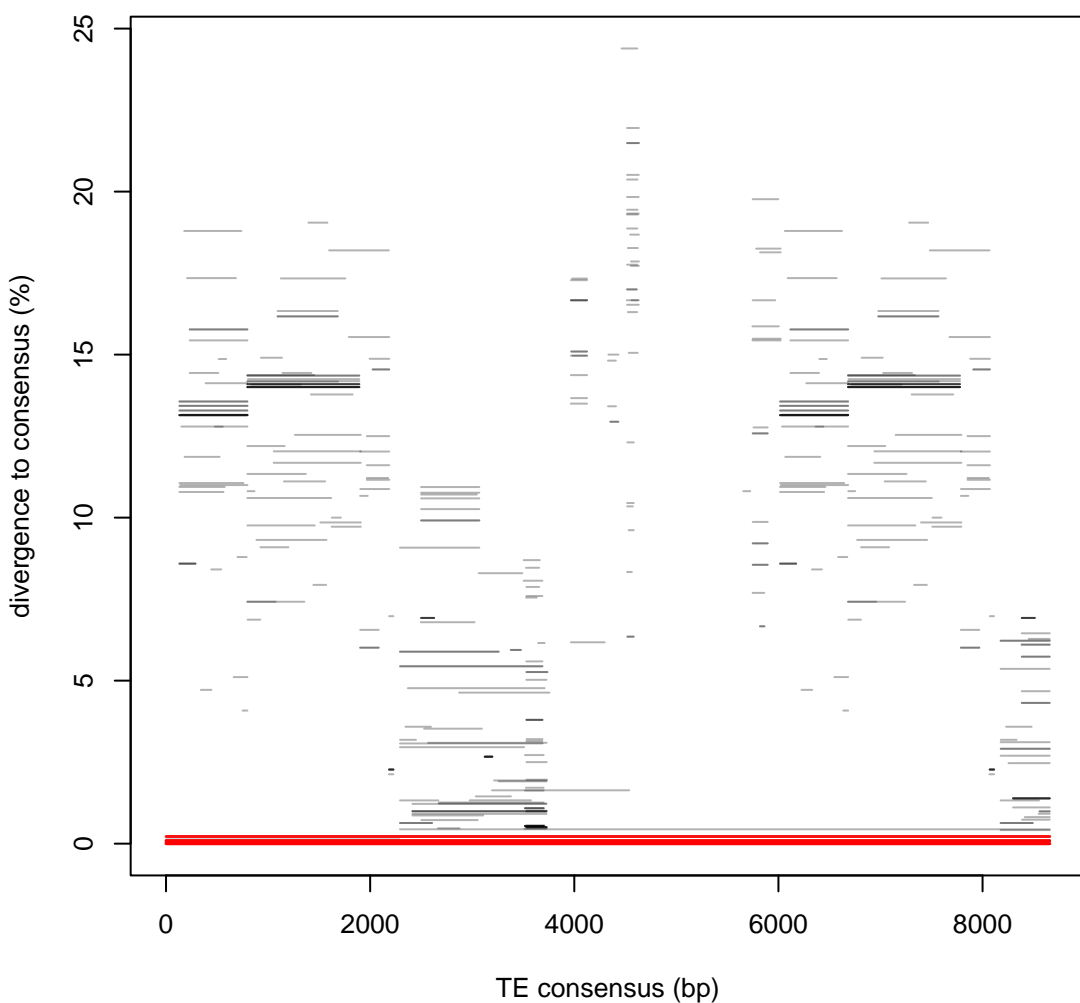
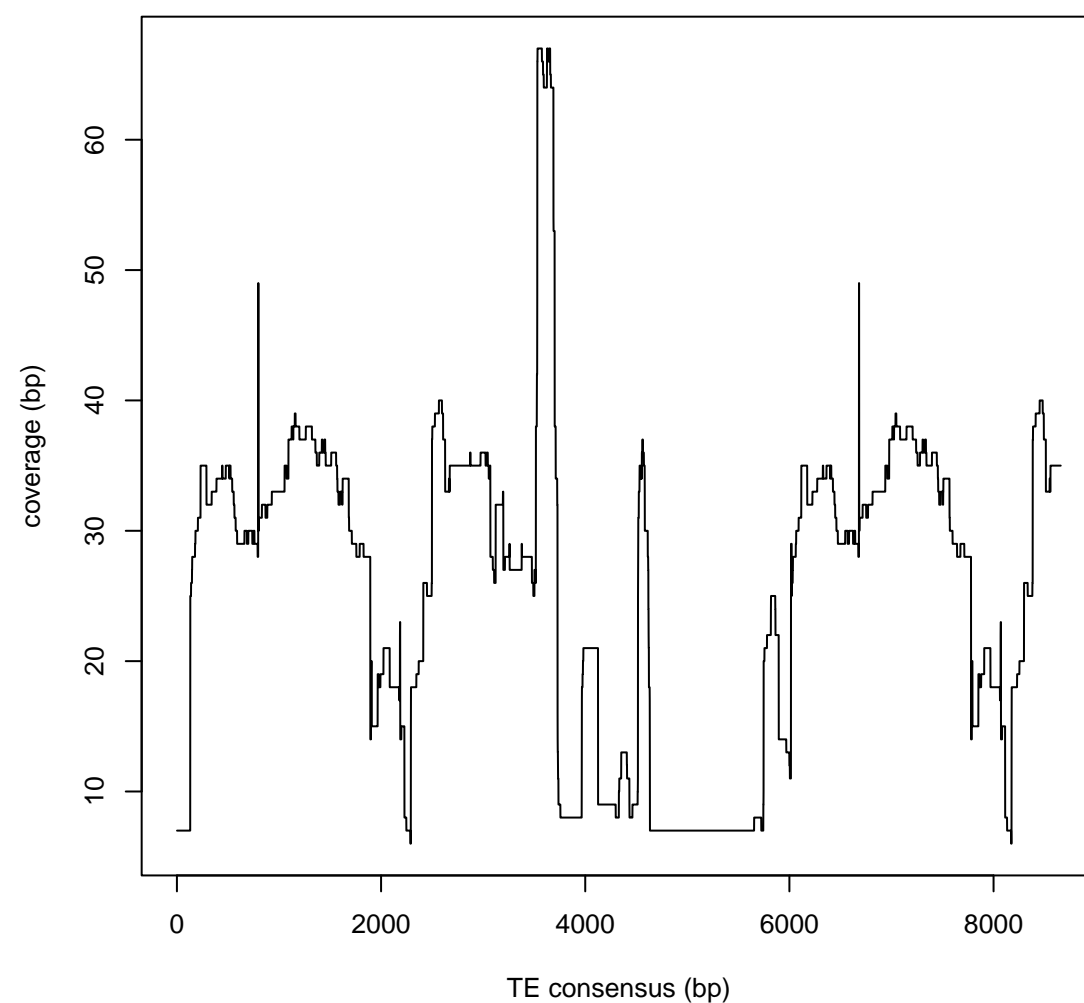


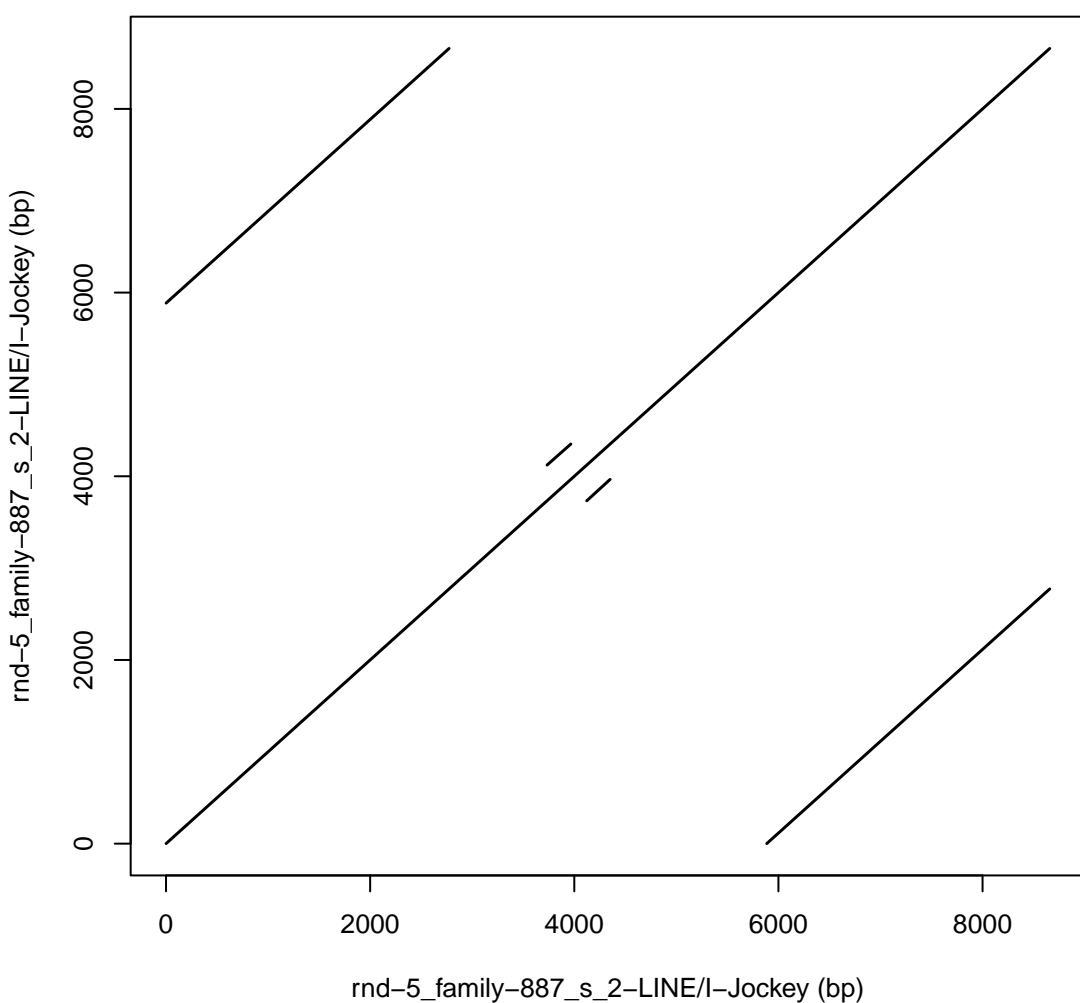
TE: rnd-5\_family-887\_s\_2-LINE/I-Jockey  
consensus size: 8658bp; fragments: 418; full length: 6 (>=7792.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

