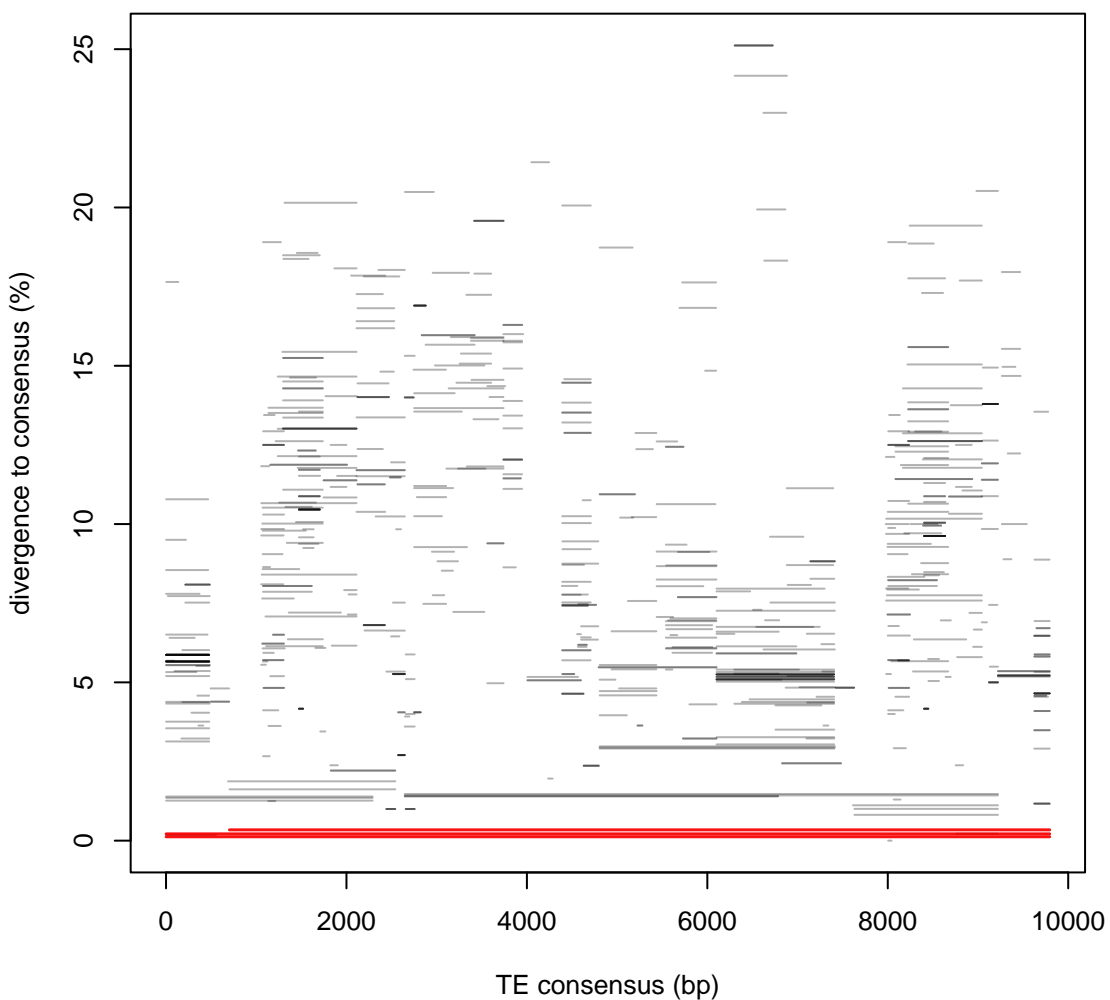
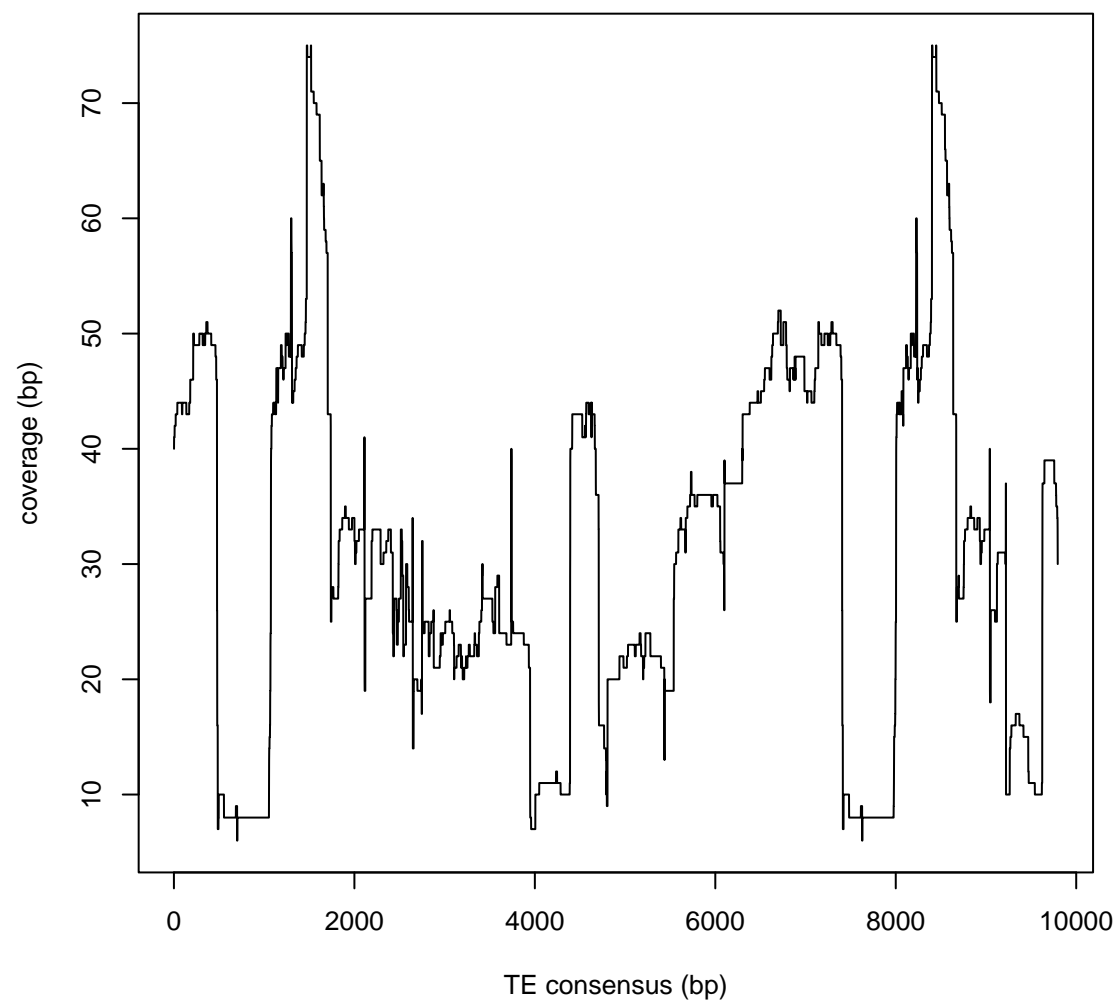


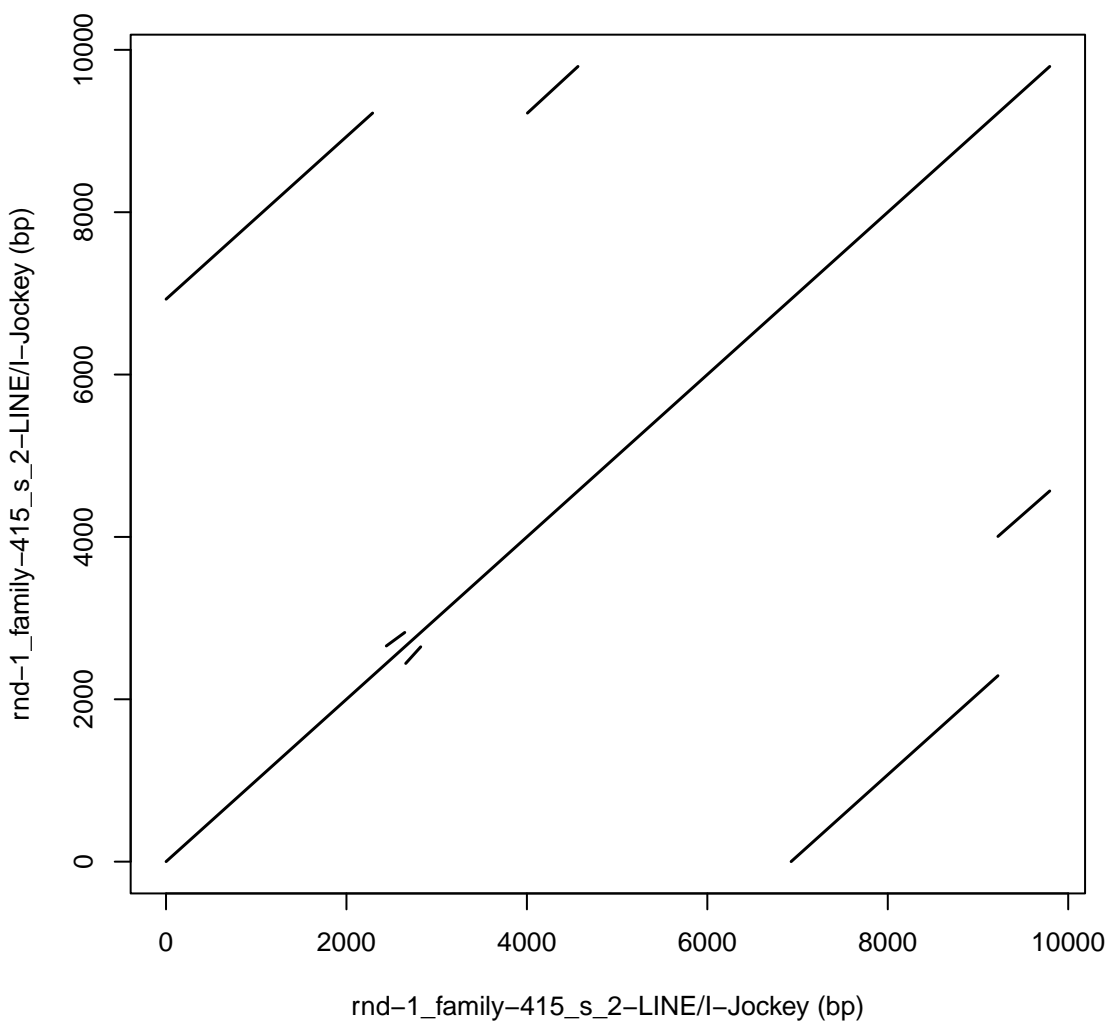
TE: rnd-1\_family-415\_s\_2-LINE/I-Jockey  
consensus size: 9795bp; fragments: 671; full length: 3 (>=8815.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

