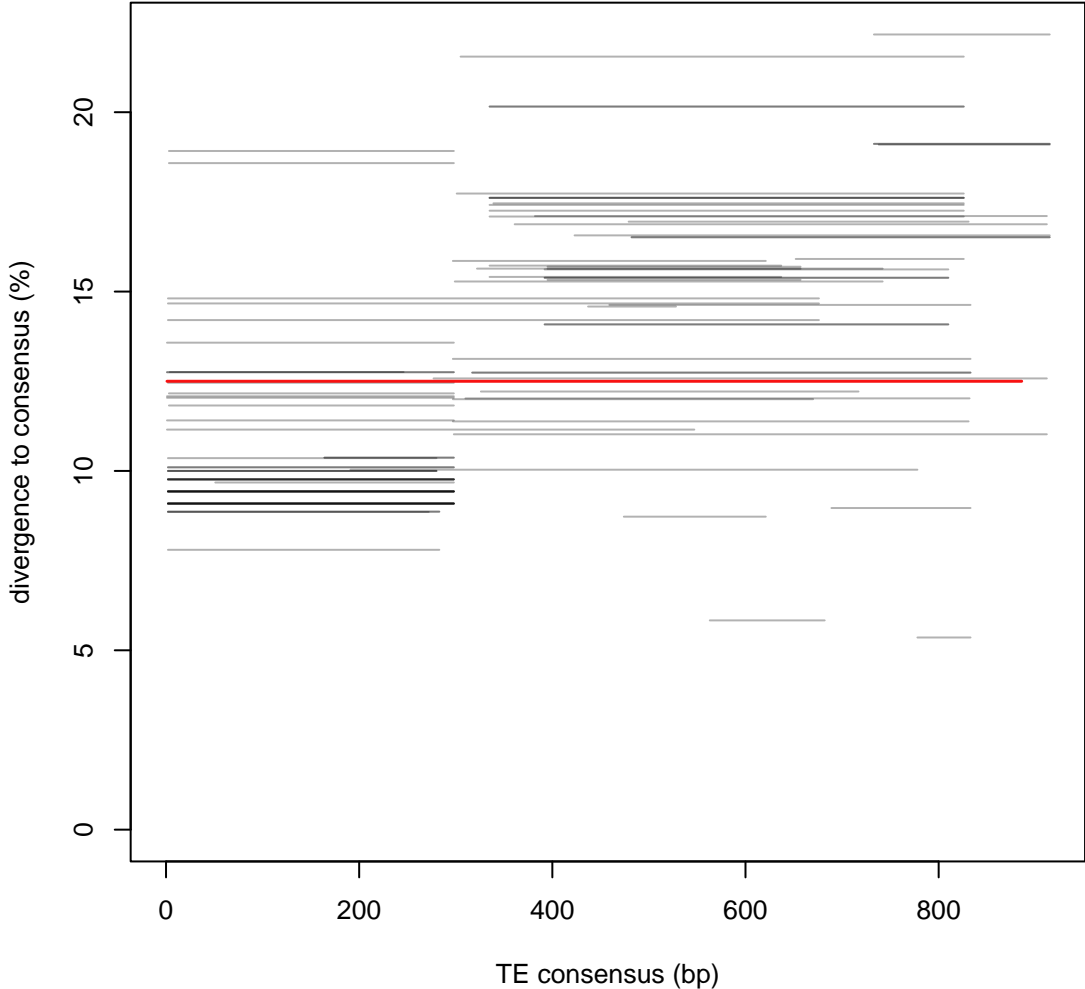
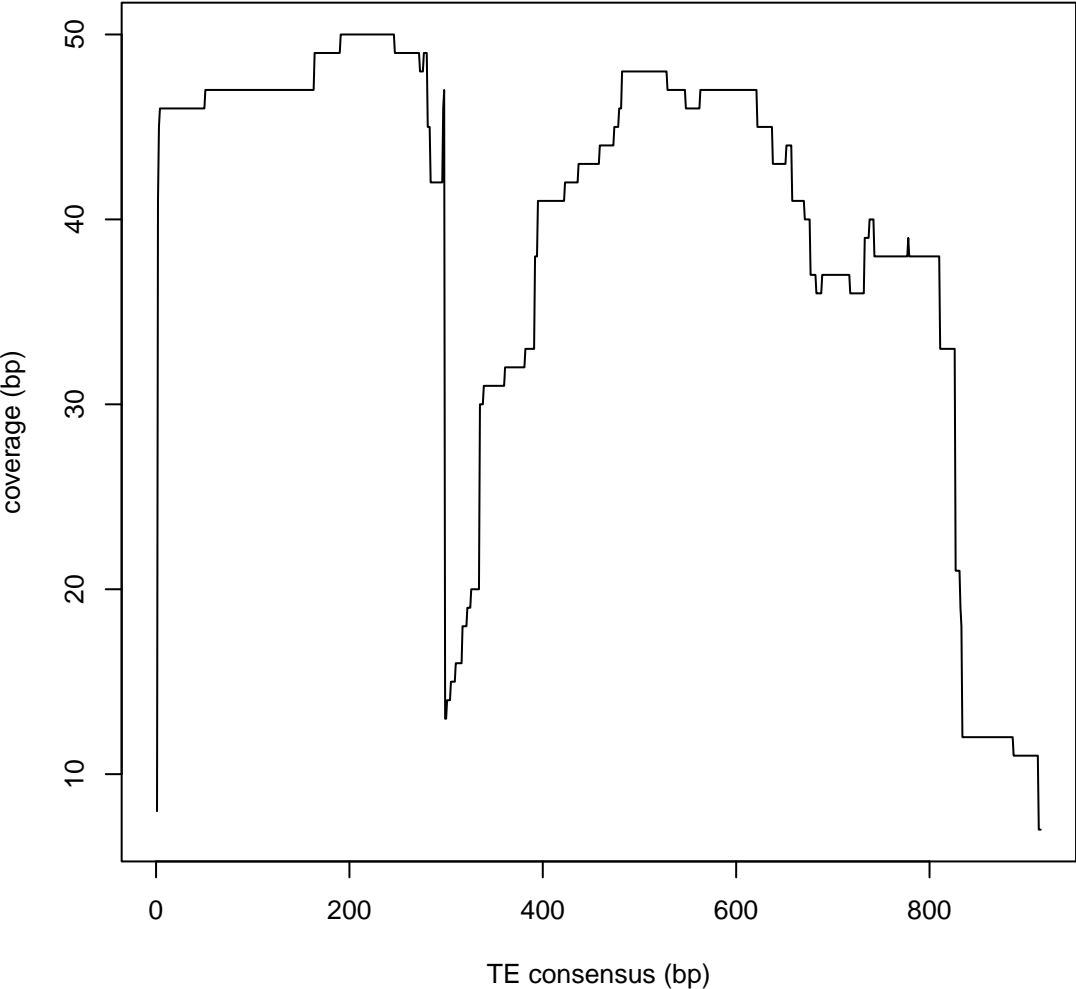


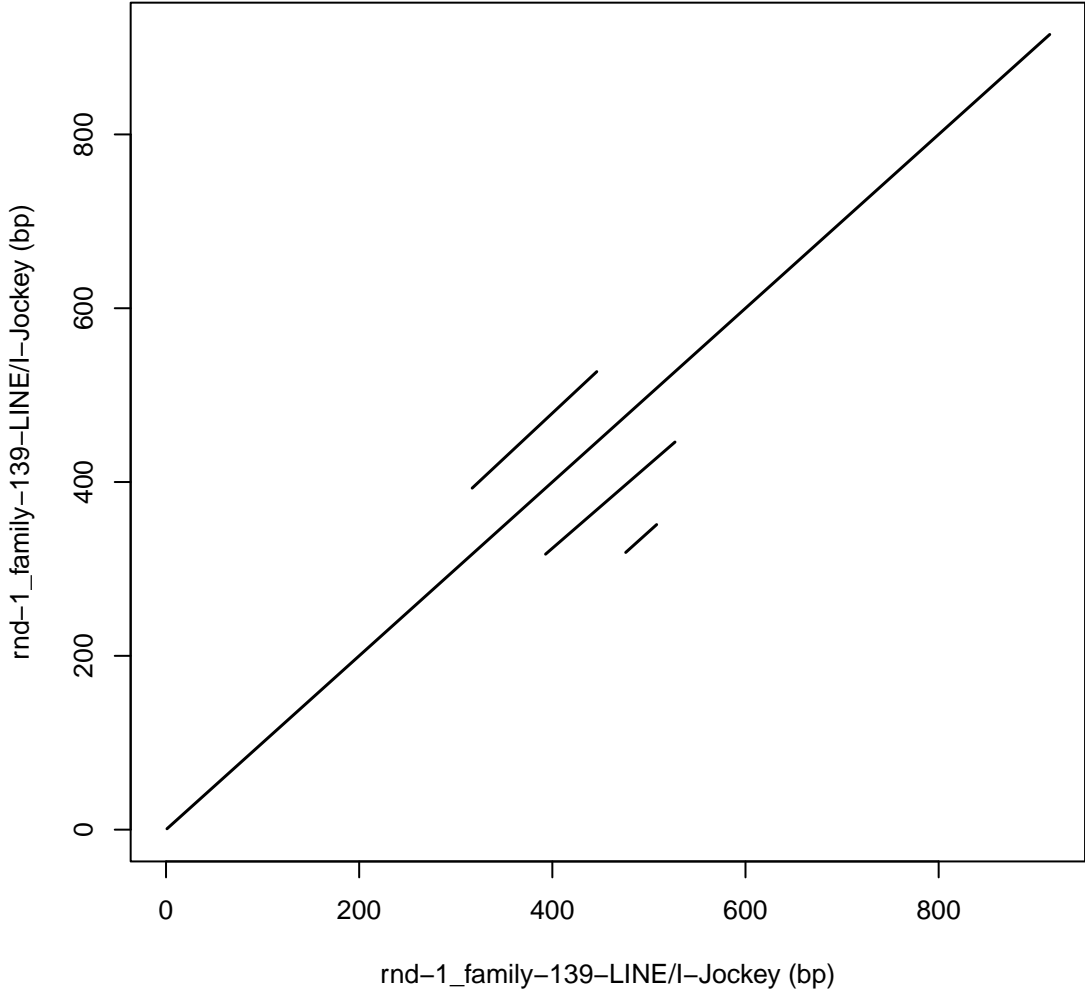
TE: rnd-1\_family-139-LINE/I-Jockey  
consensus size: 915bp; fragments: 100; full length: 1 (>=823.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

