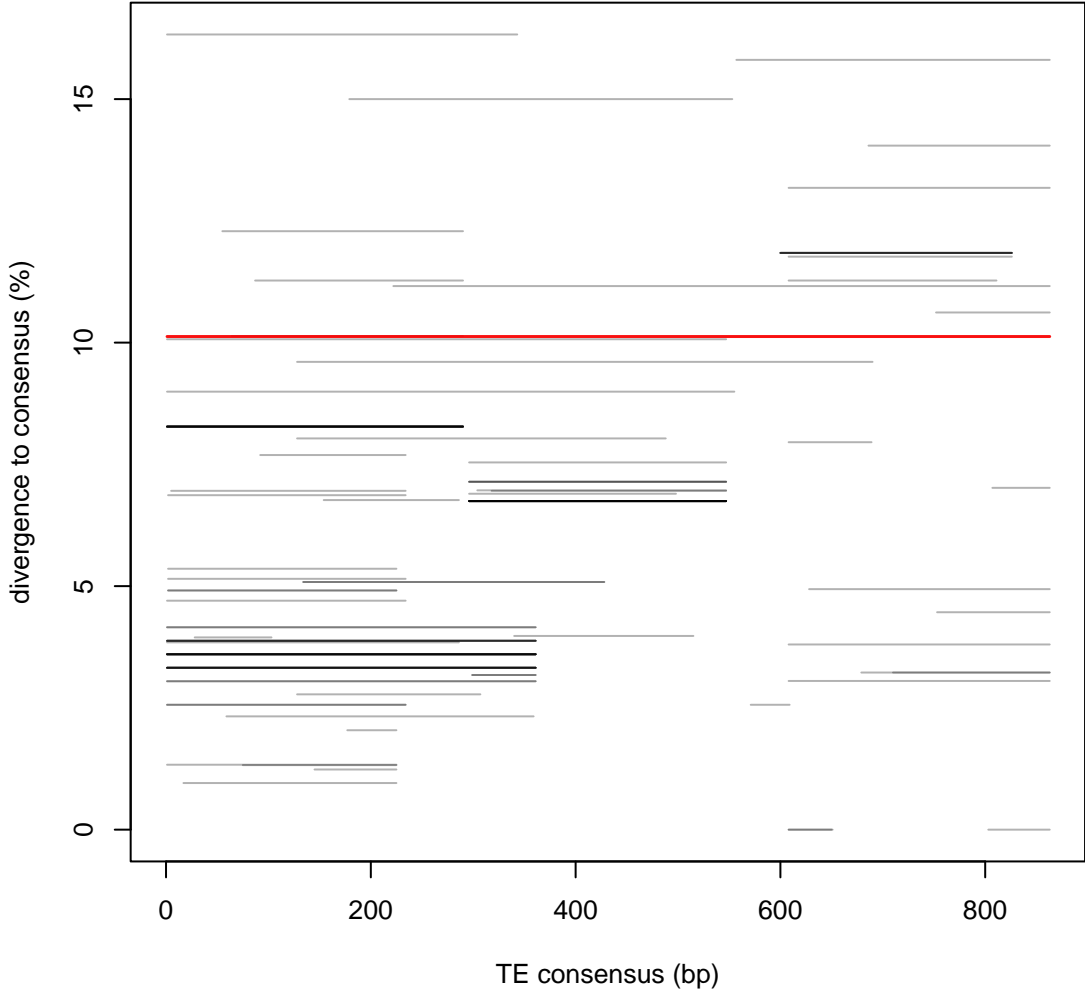
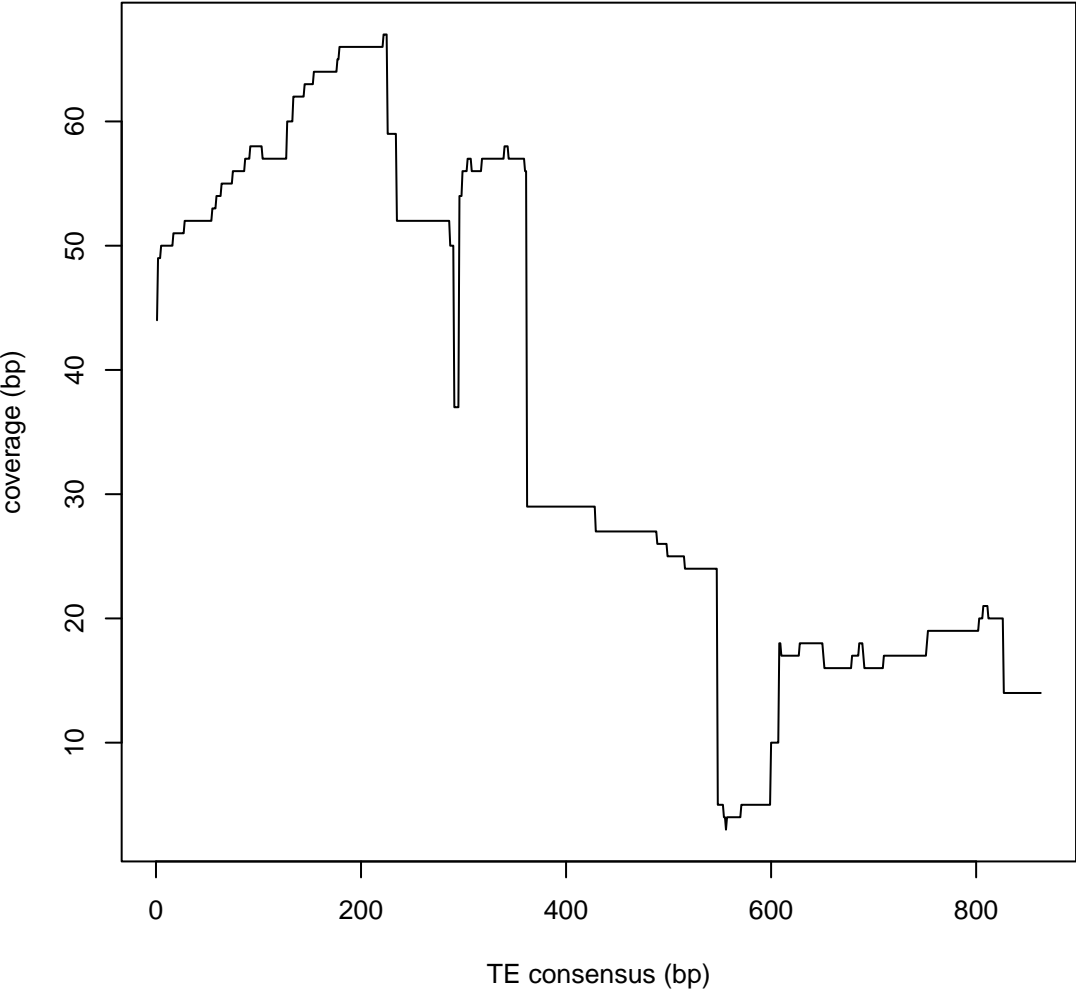


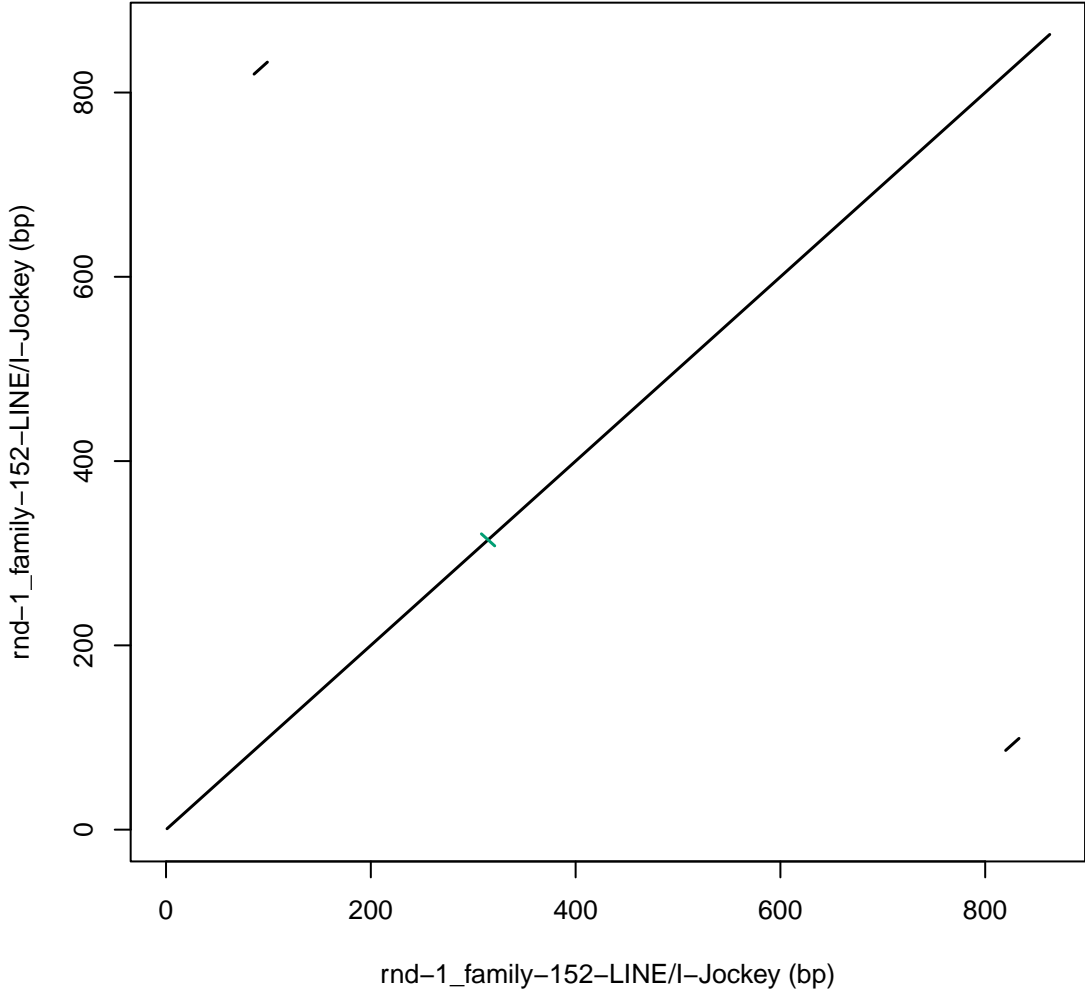
TE: rnd-1\_family-152-LINE/I-Jockey  
consensus size: 863bp; fragments: 113; full length: 1 (>=776.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

