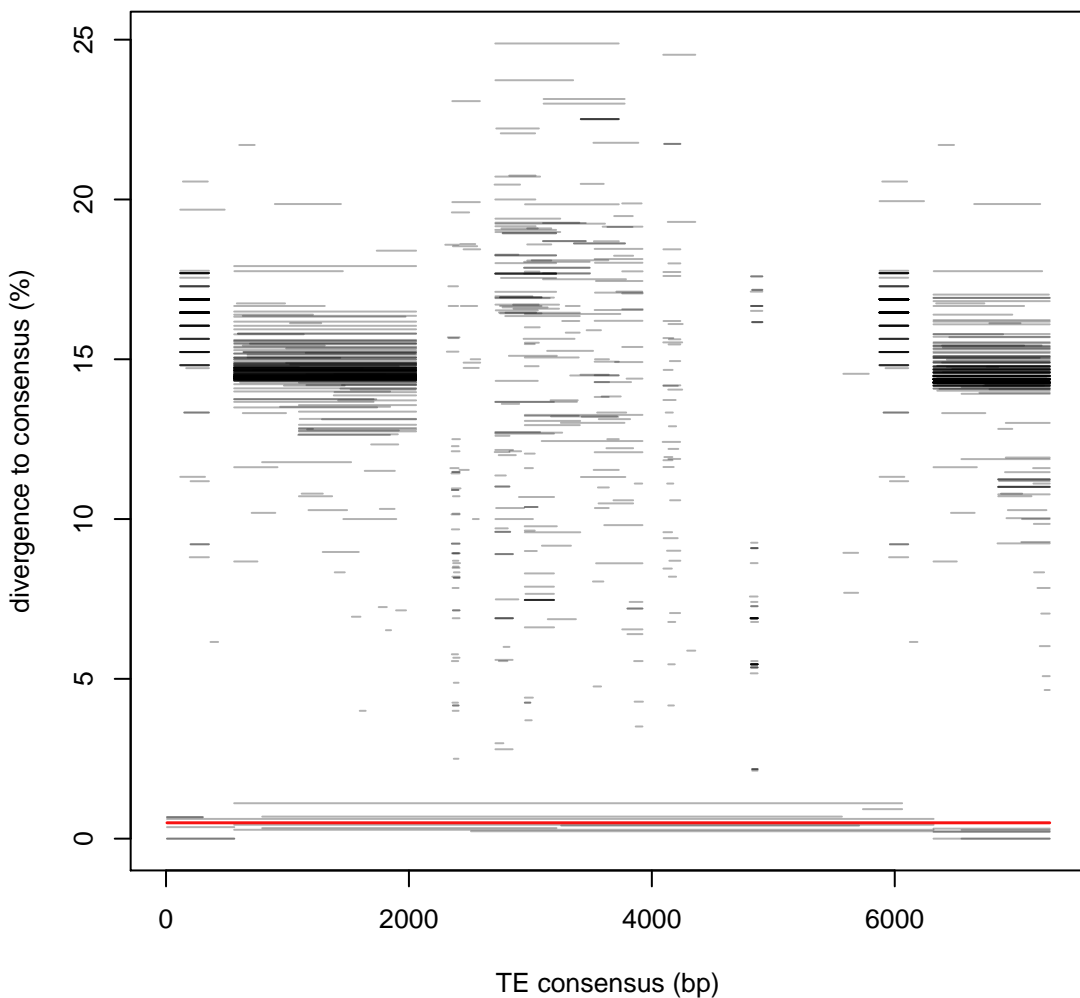
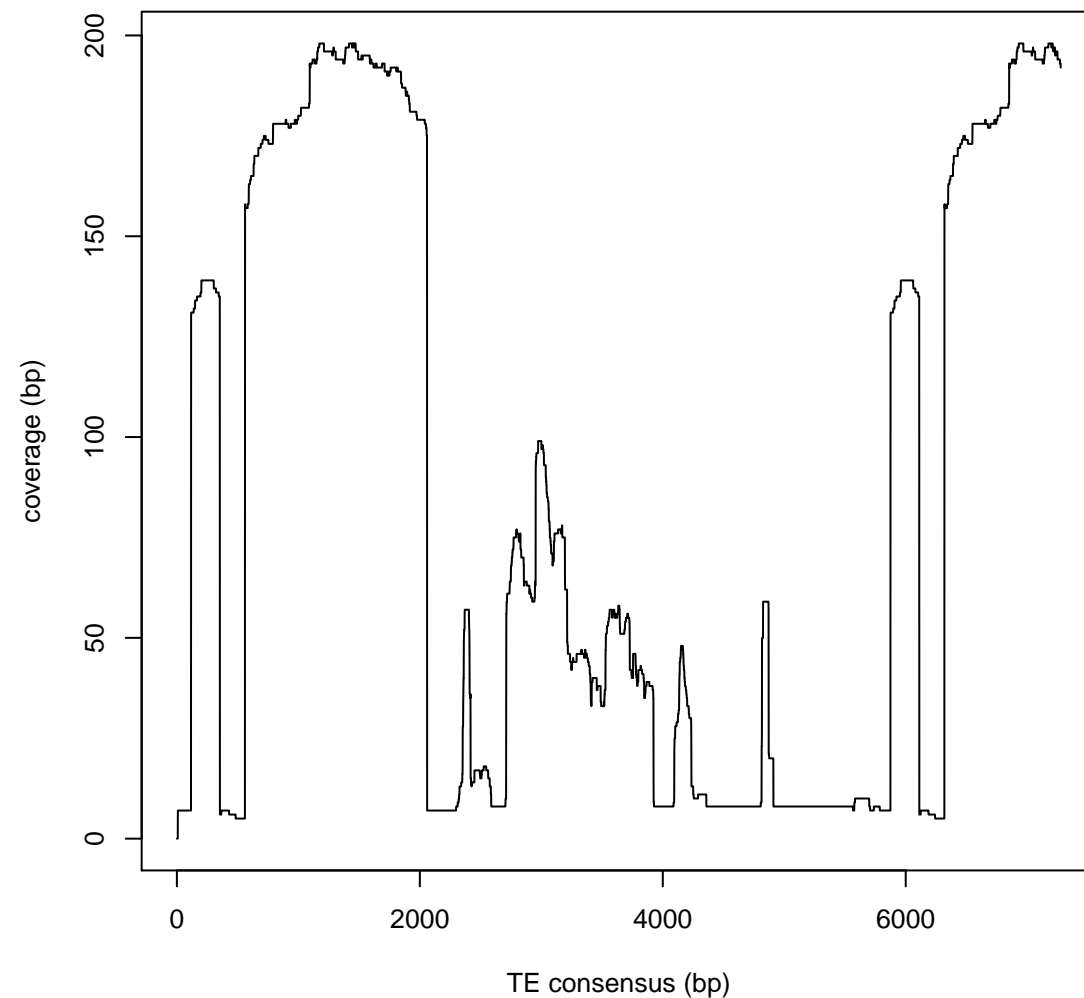


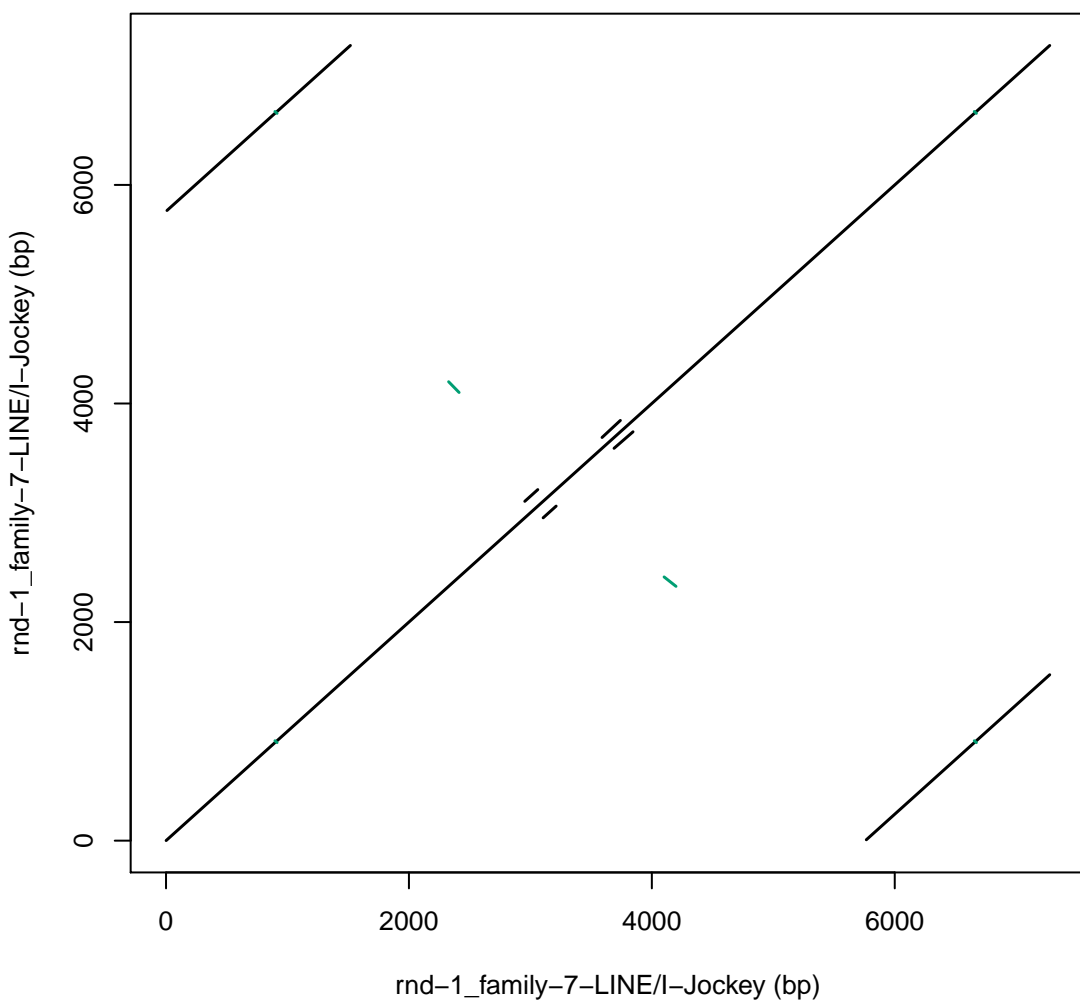
TE: rnd-1\_family-7-LINE/l-Jockey  
consensus size: 7276bp; fragments: 1066; full length: 1 (>=6548.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

