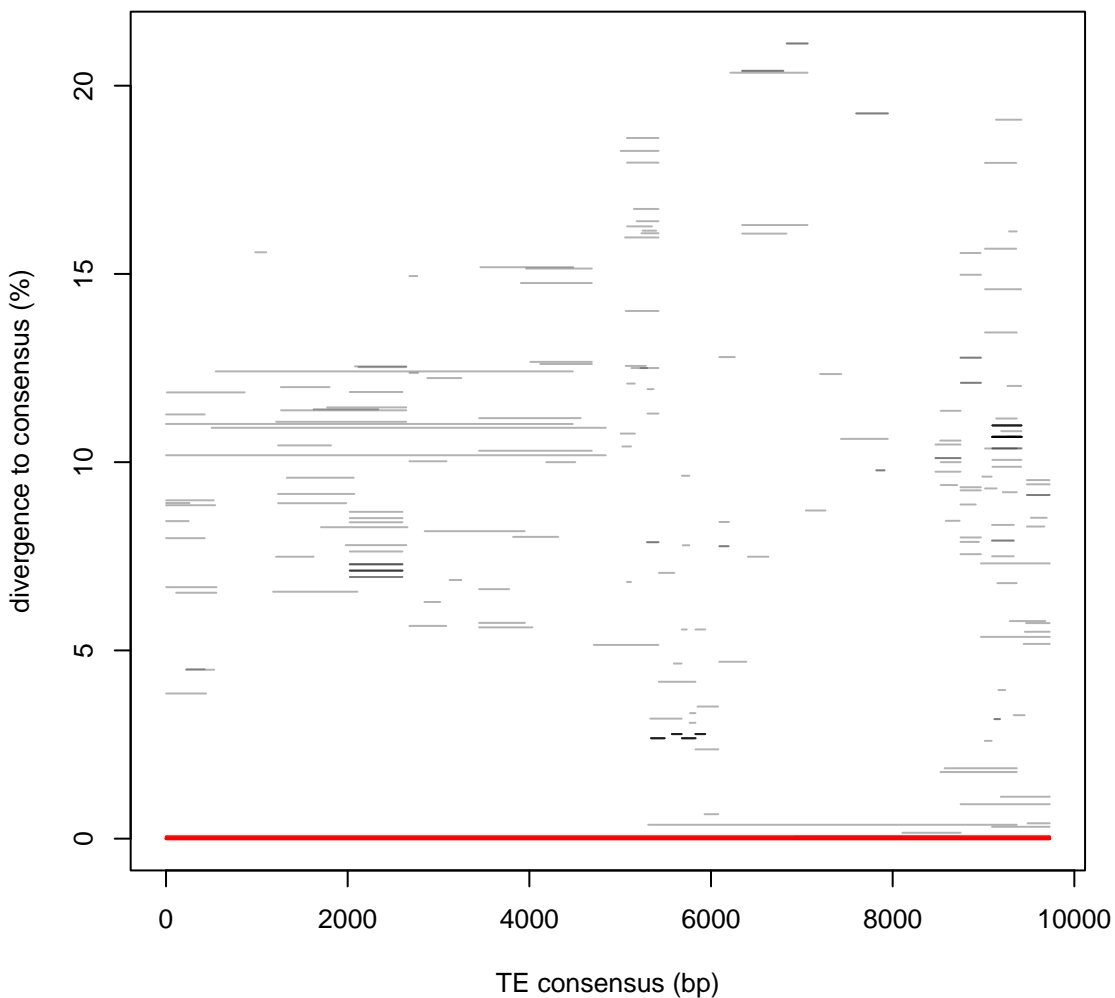
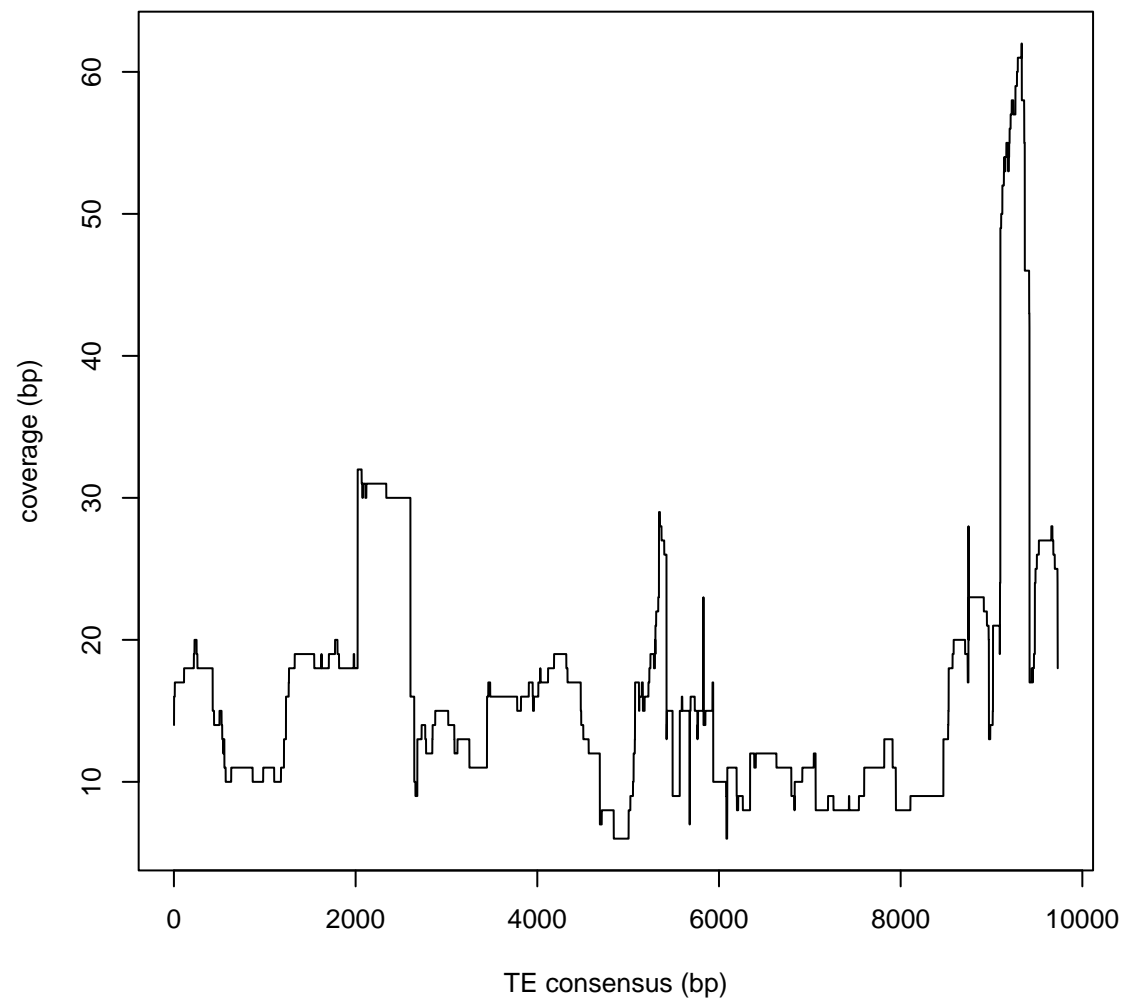


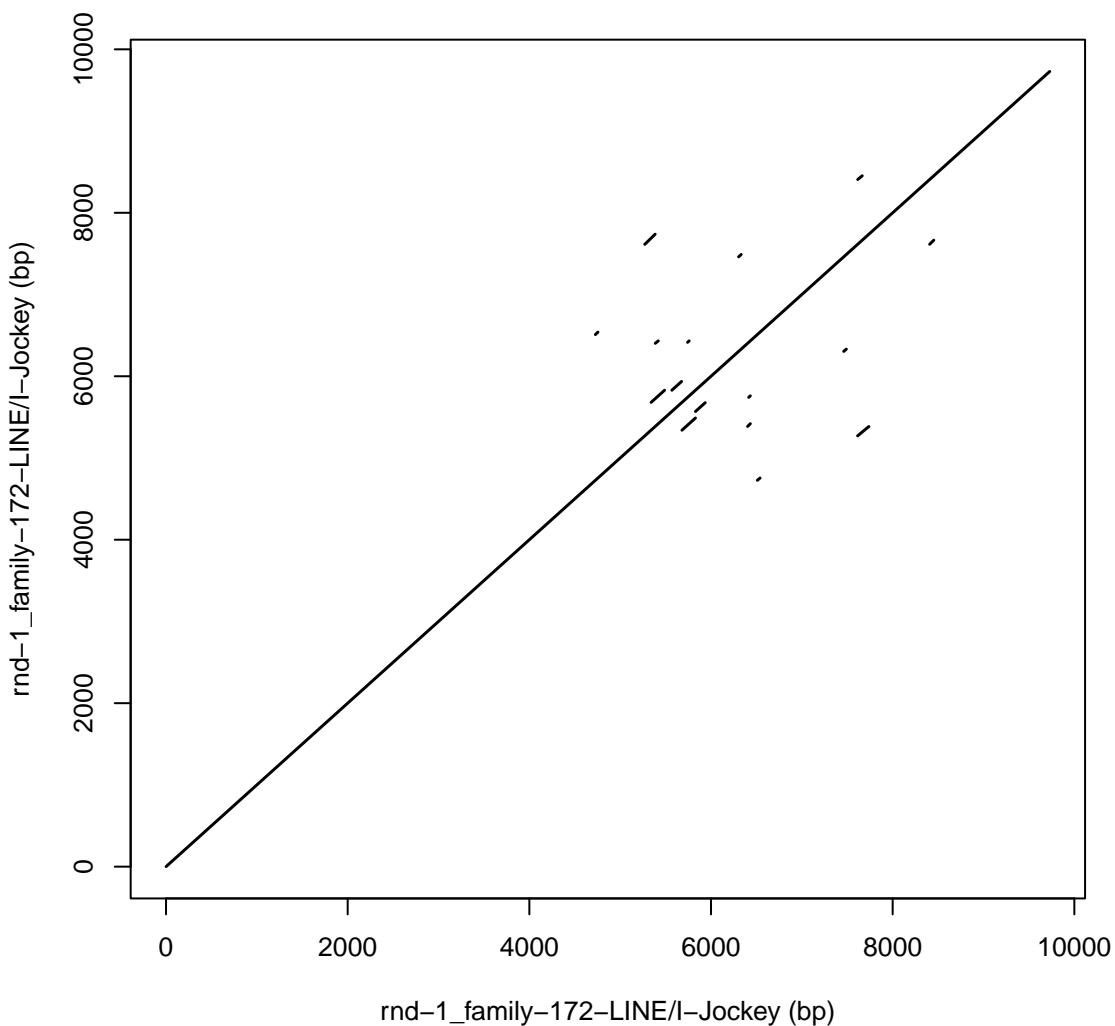
TE: rnd-1\_family-172-LINE/I-Jockey  
consensus size: 9729bp; fragments: 253; full length: 5 (>=8756.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

