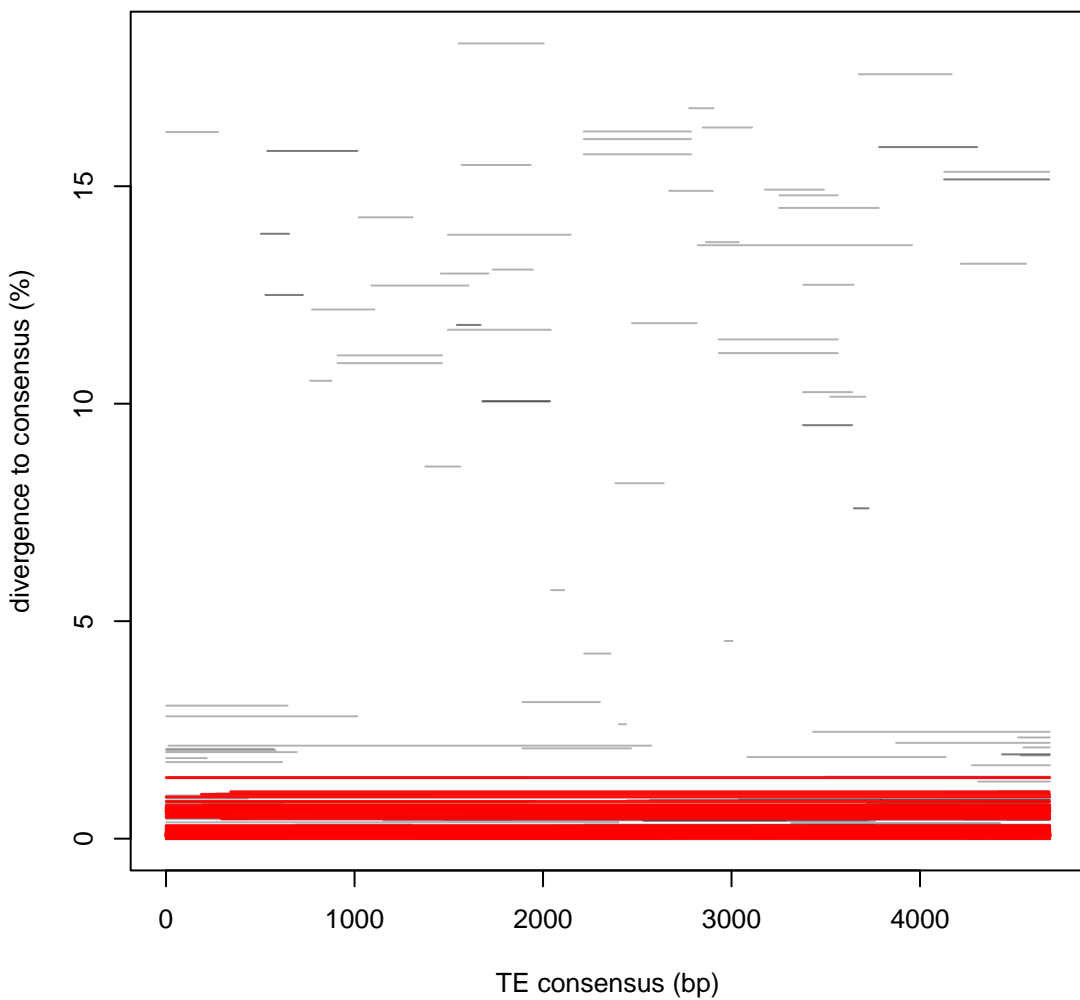
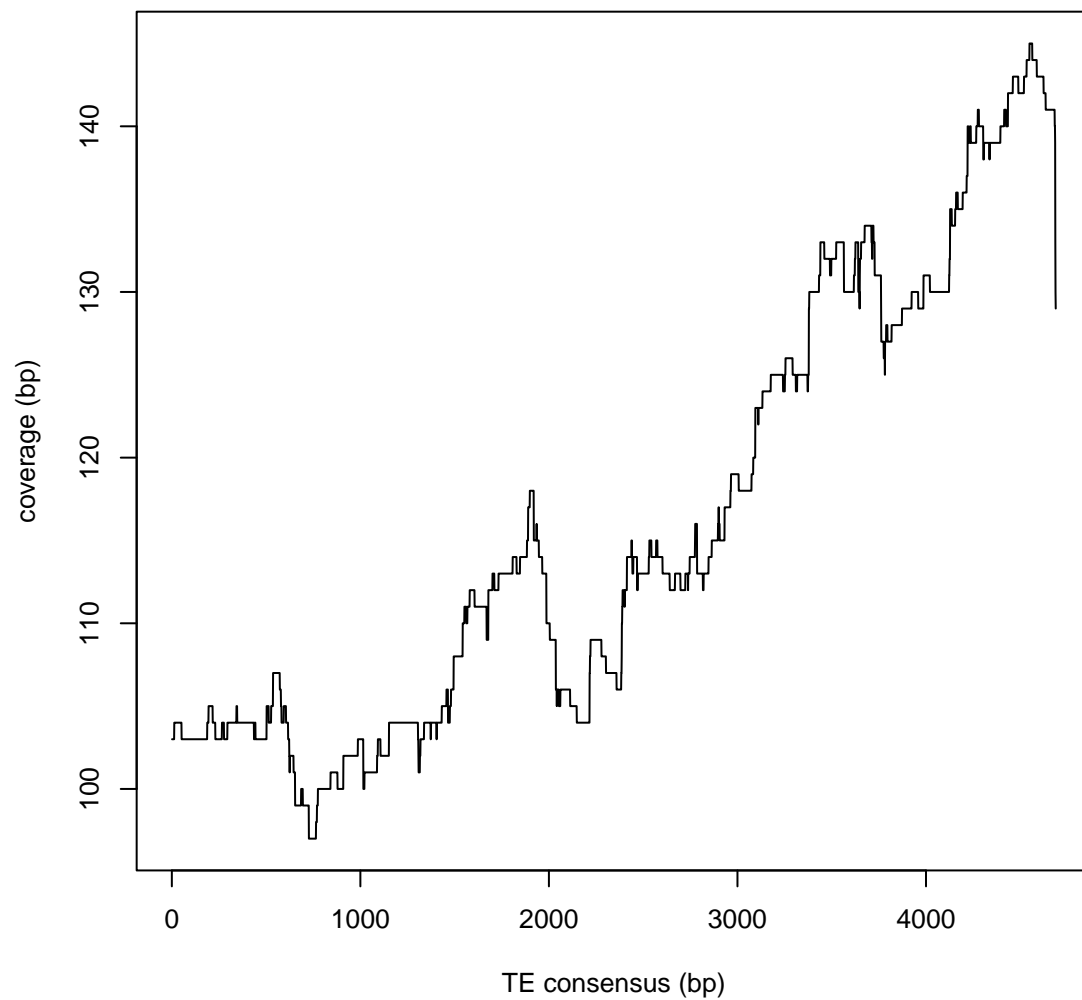


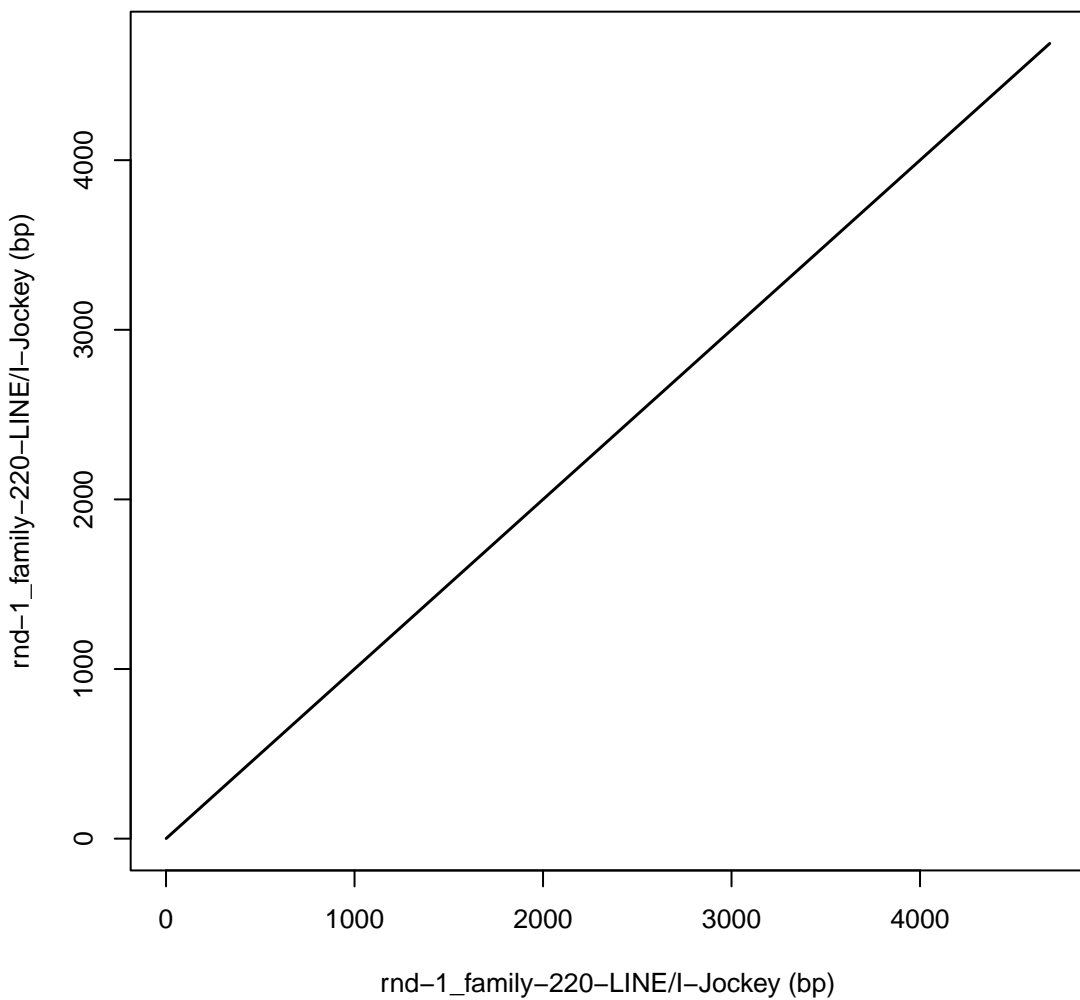
TE: rnd-1\_family-220-LINE/I-Jockey  
consensus size: 4688bp; fragments: 262; full length: 75 (>=4219.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

