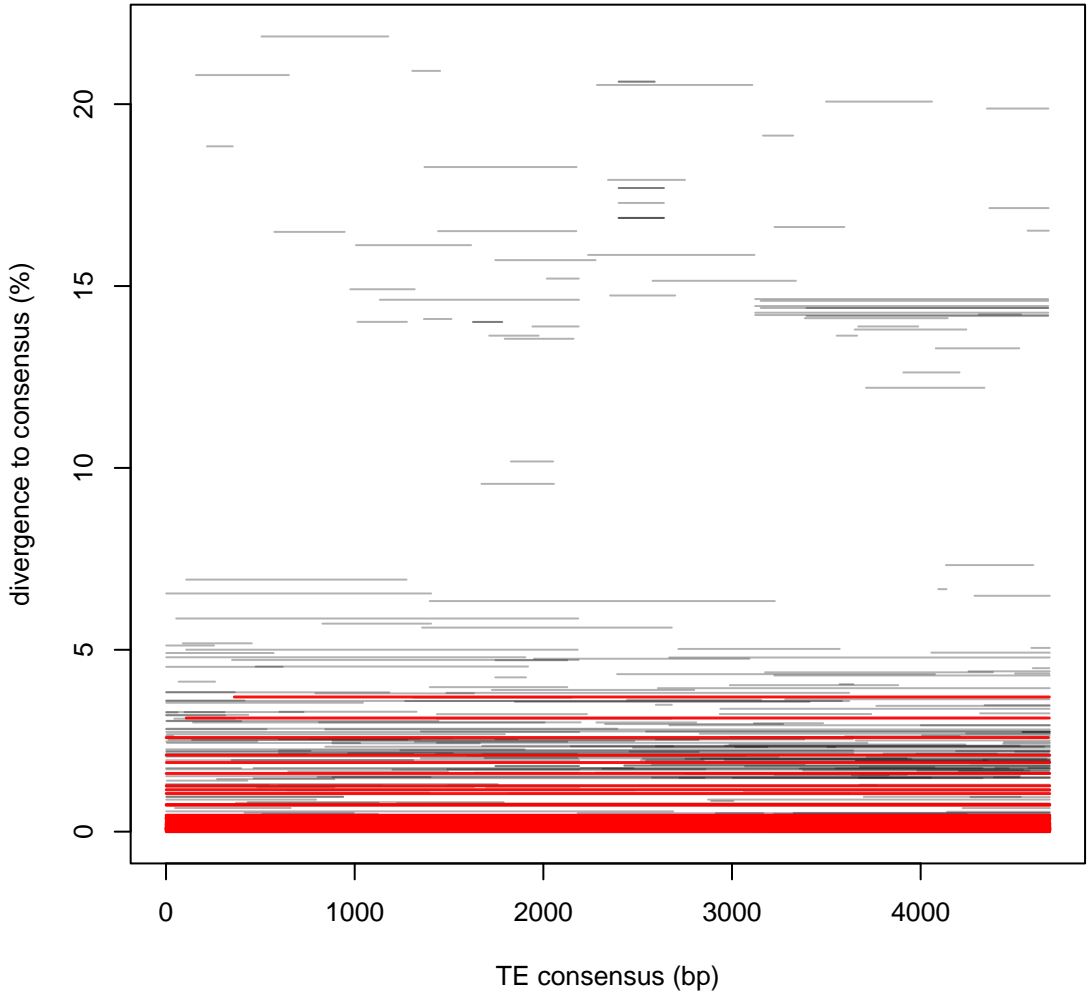
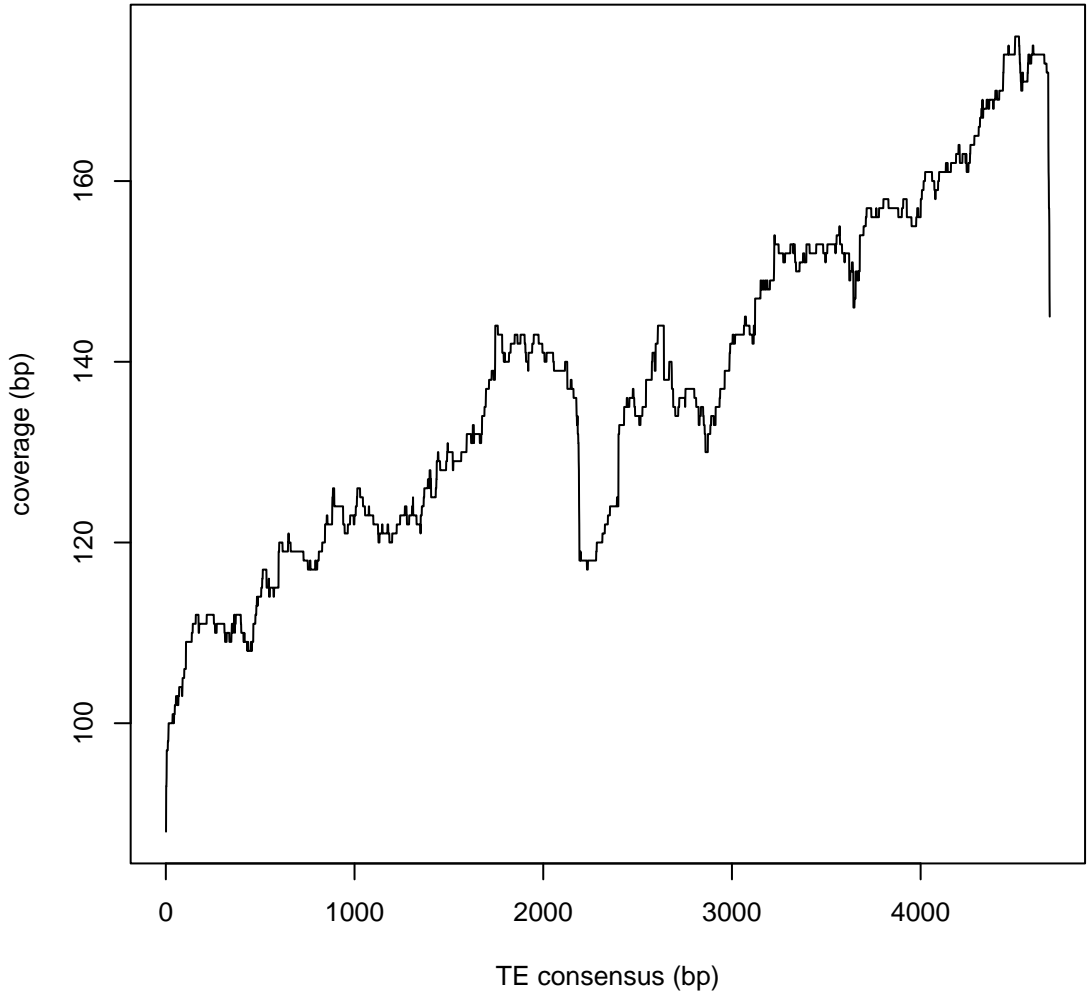


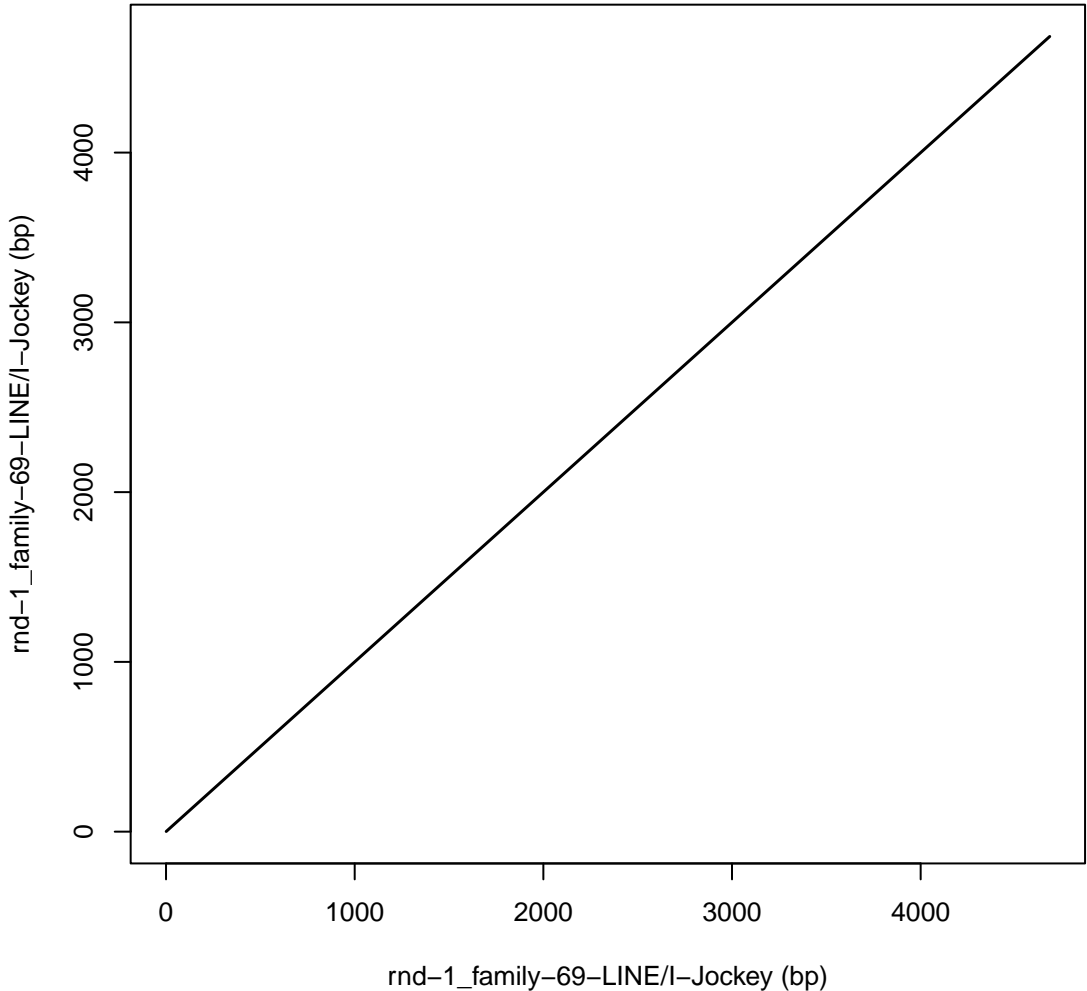
TE: rnd-1\_family-69-LINE/I-Jockey  
consensus size: 4684bp; fragments: 394; full length: 51 (>=4215.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

