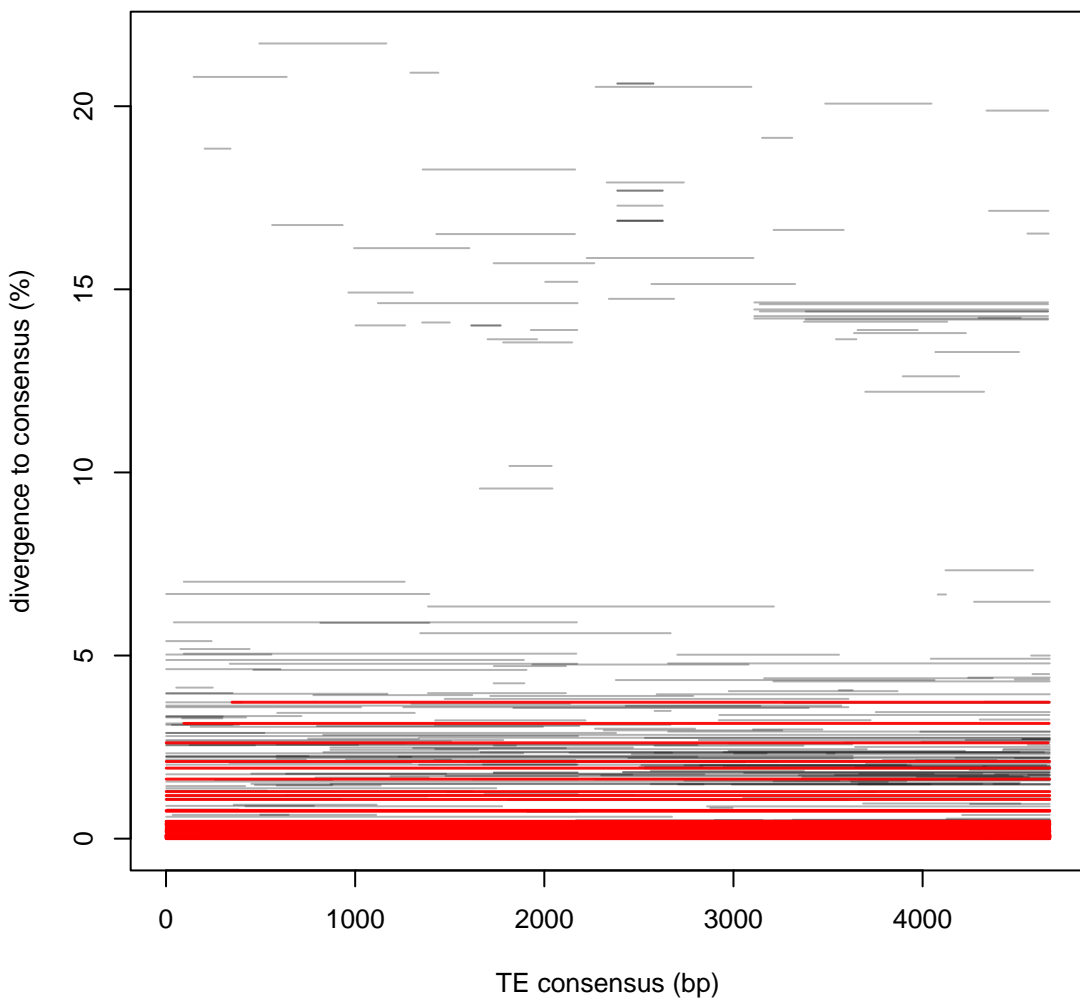
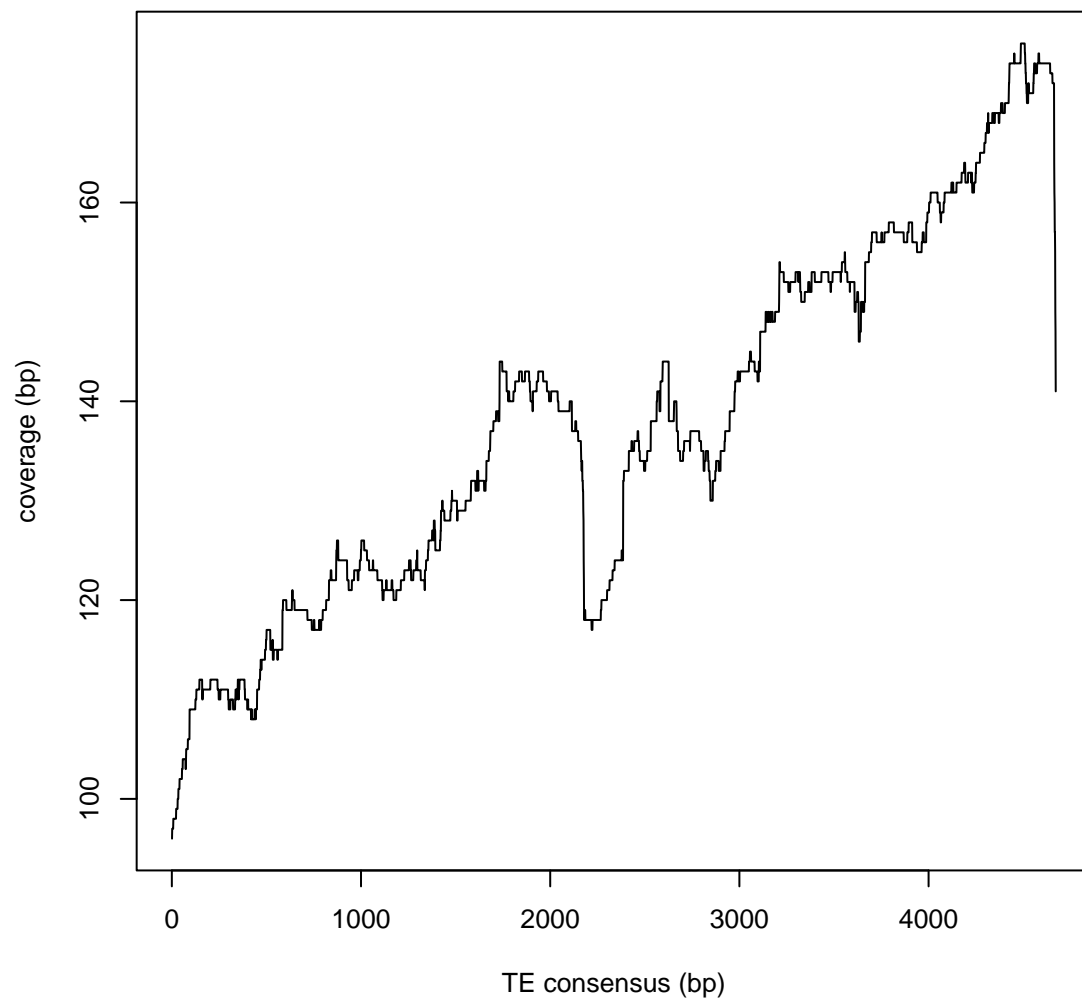


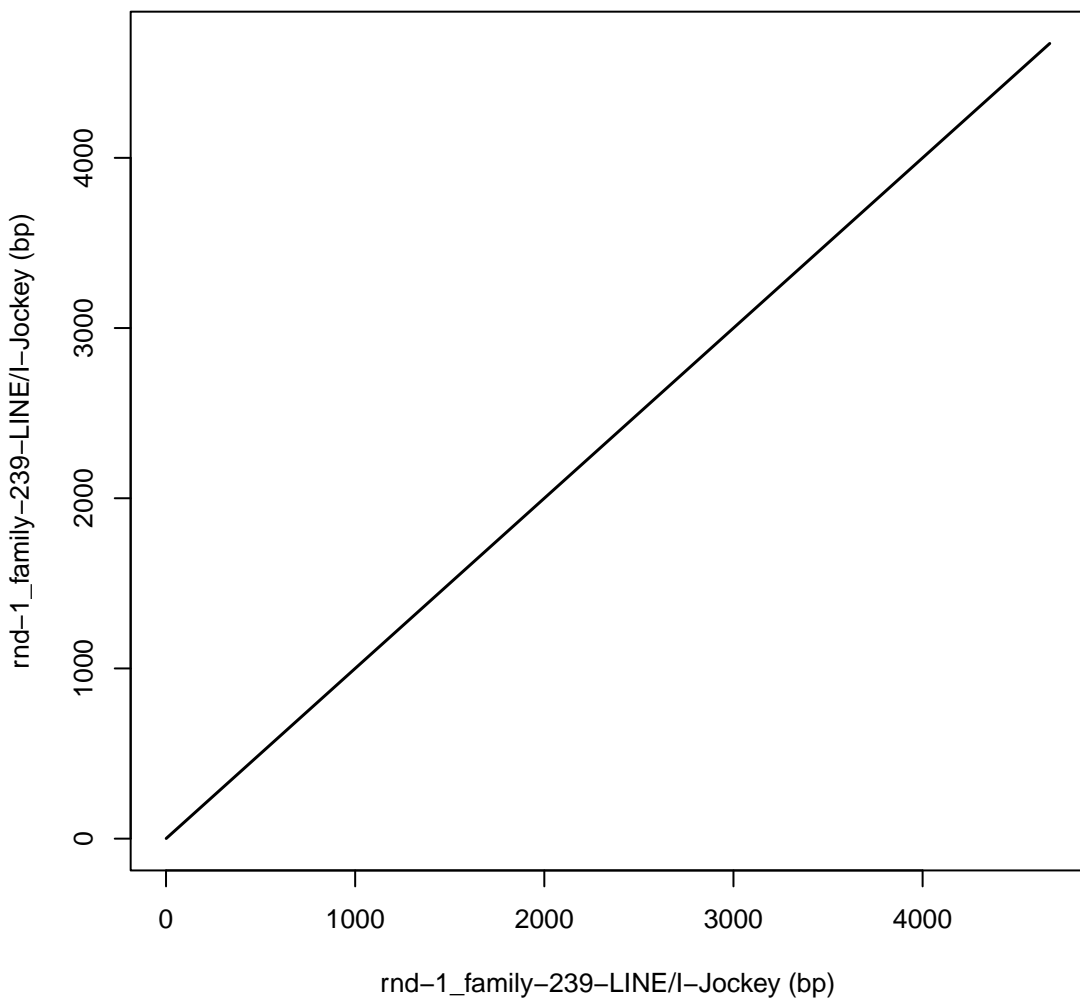
TE: rnd-1\_family-239-LINE/I-Jockey  
consensus size: 4672bp; fragments: 392; full length: 52 (>=4204.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

