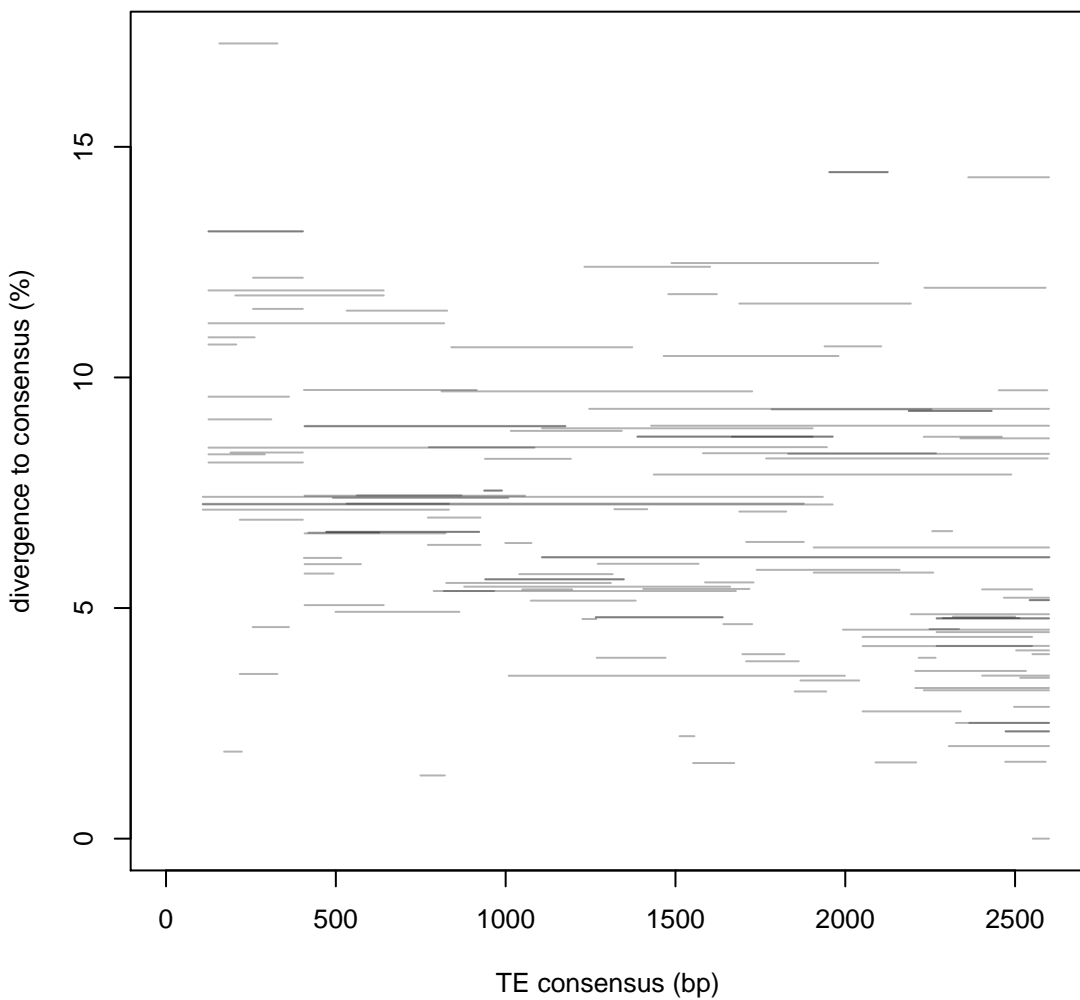
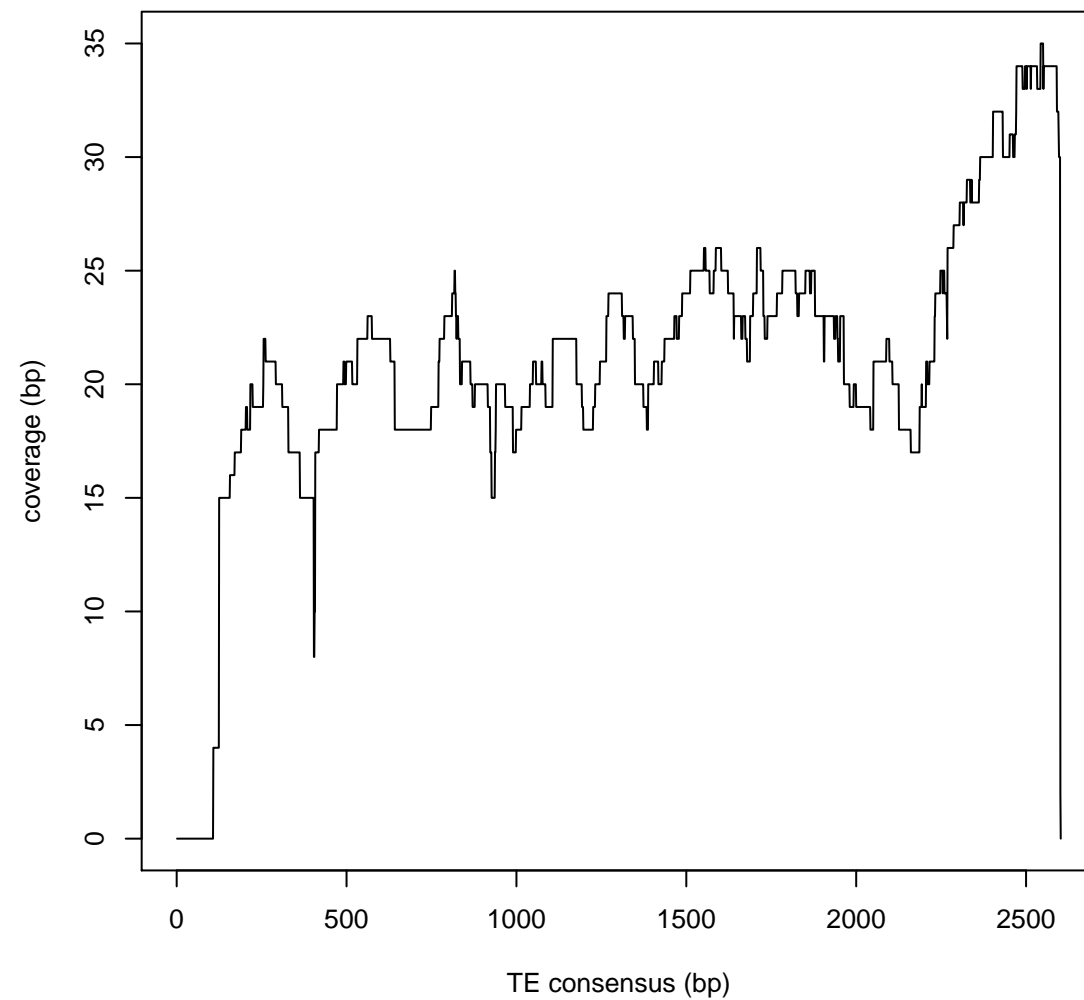


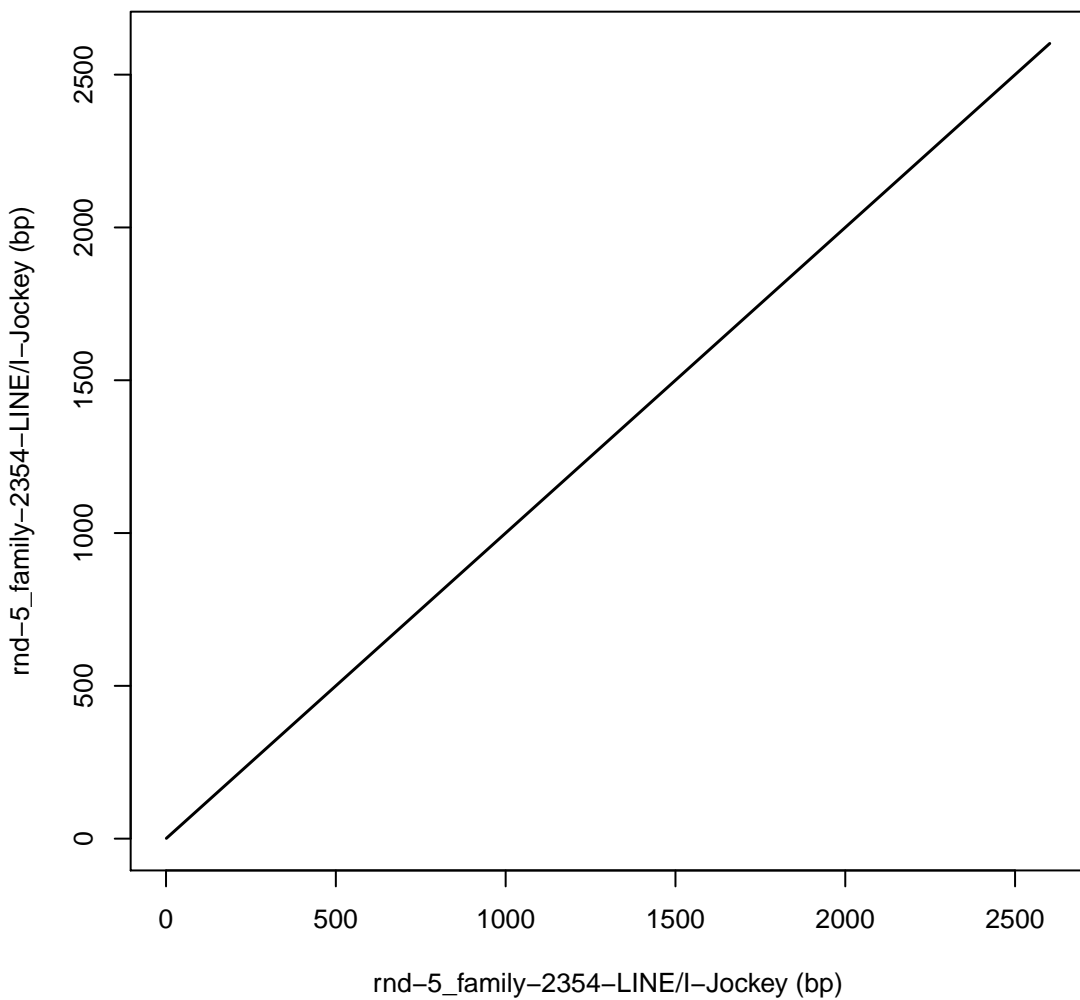
TE: rnd-5_family-2354-LINE/l-Jockey
consensus size: 2602bp; fragments: 144; full length: 0 (≥ 2341.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

