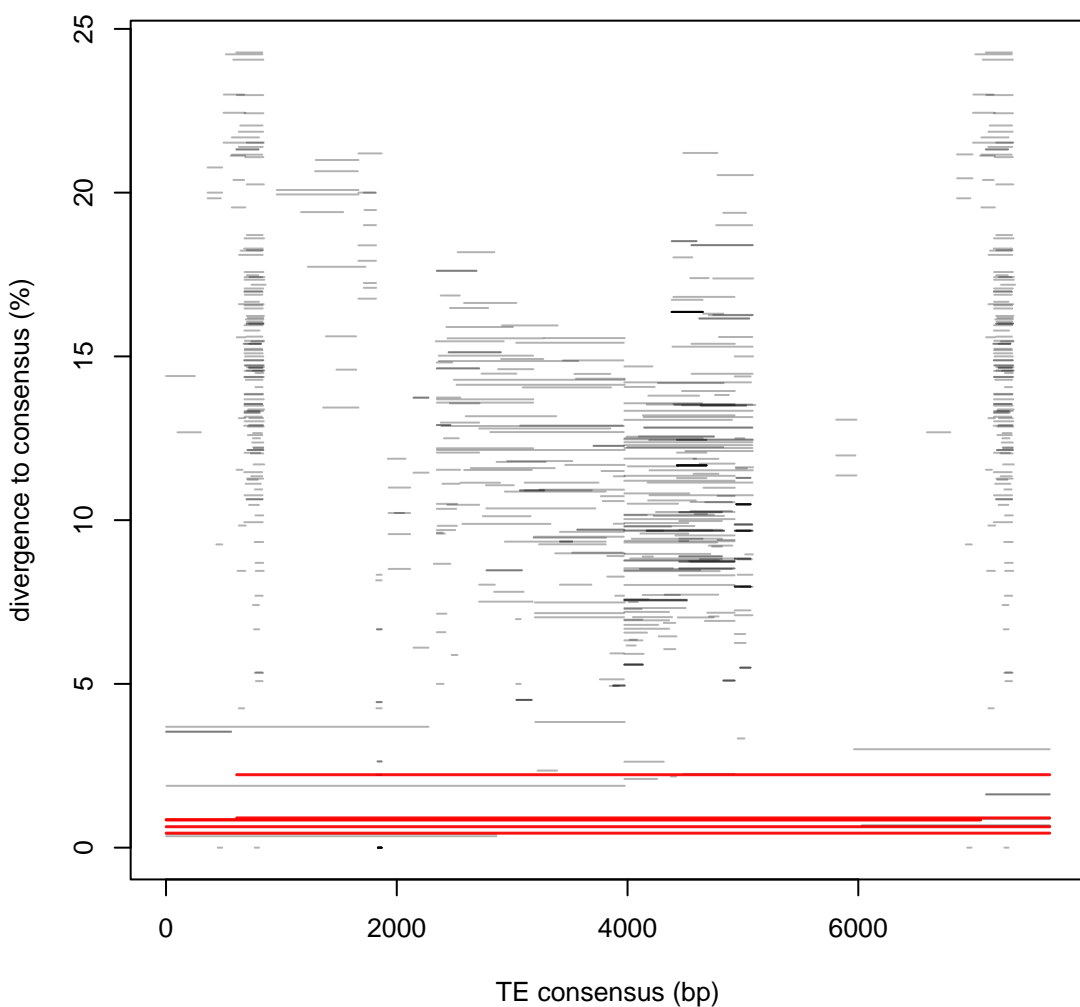
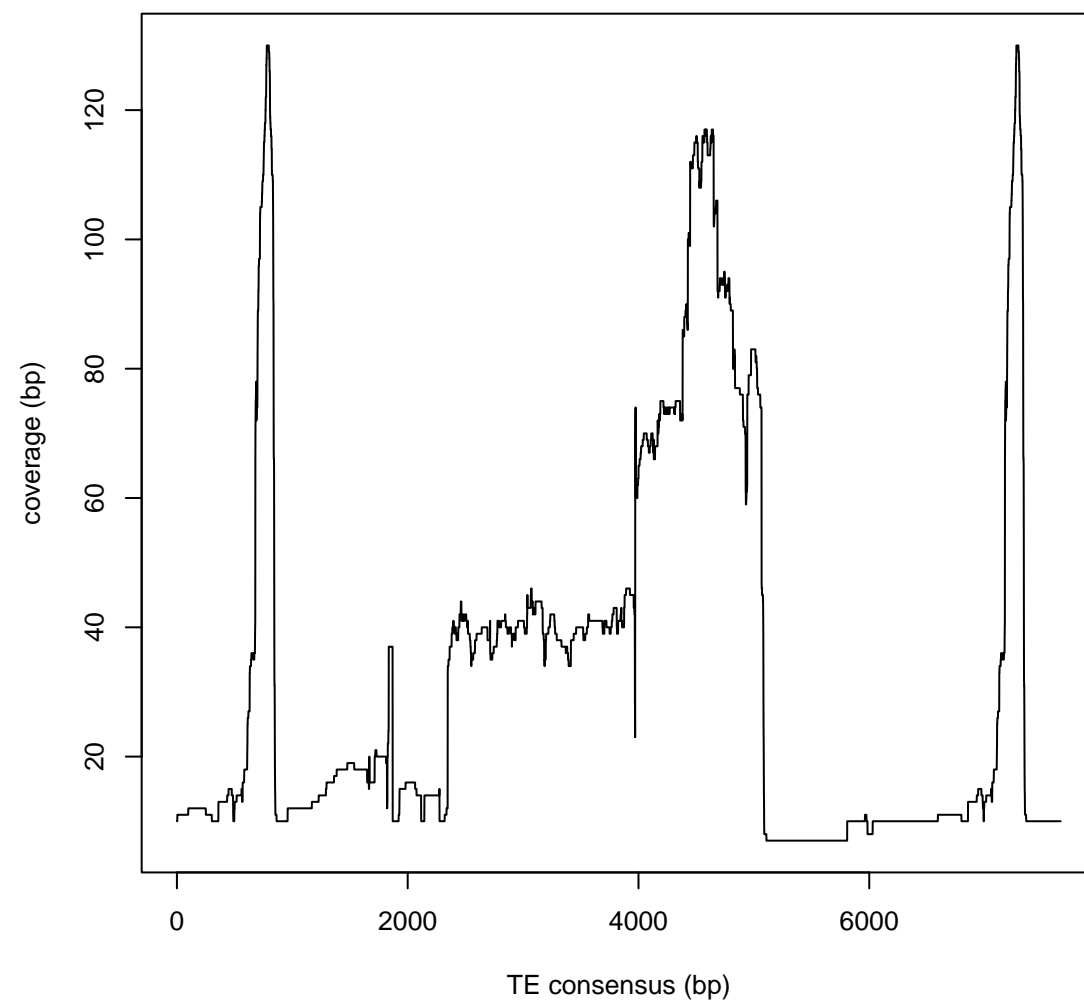


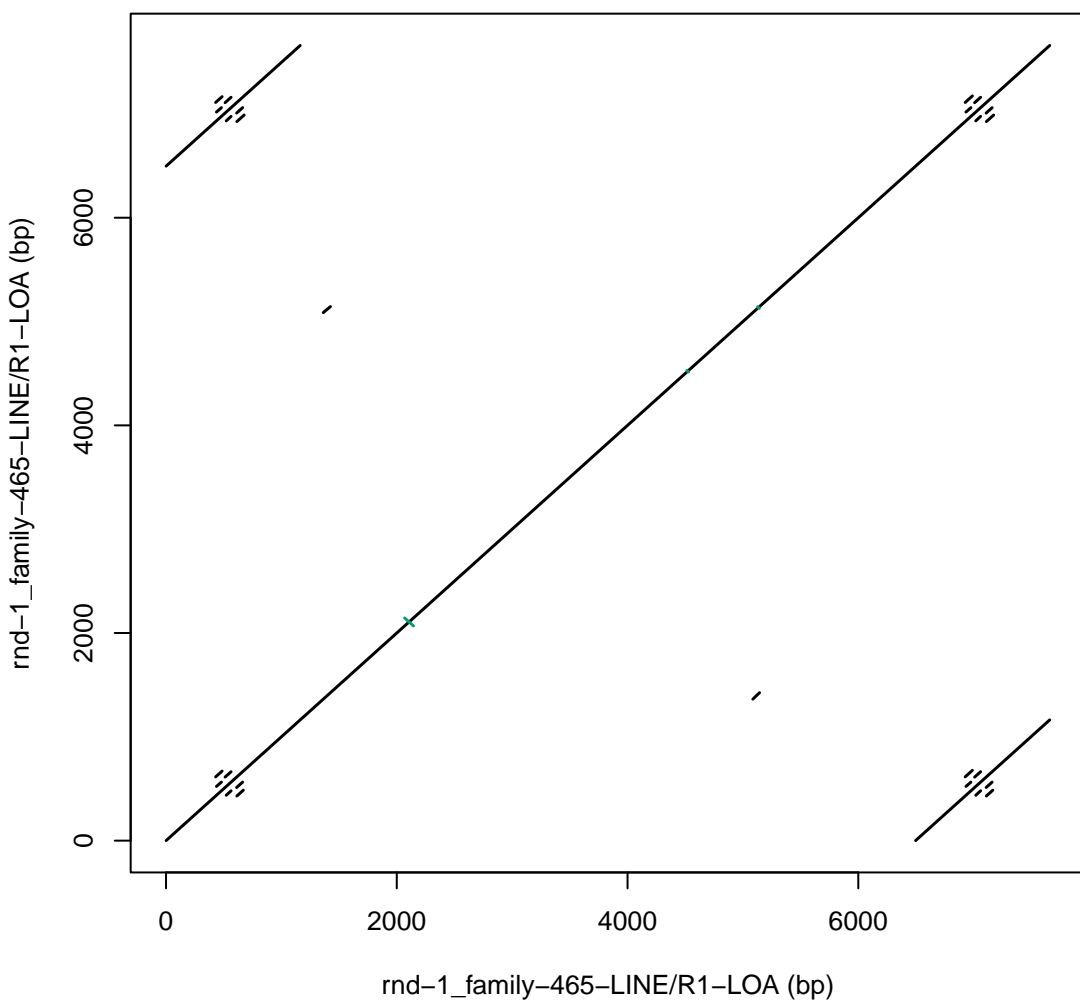
TE: rnd-1_family-465-LINE/R1-LOA
consensus size: 7659bp; fragments: 696; full length: 6 (≥ 6893.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

