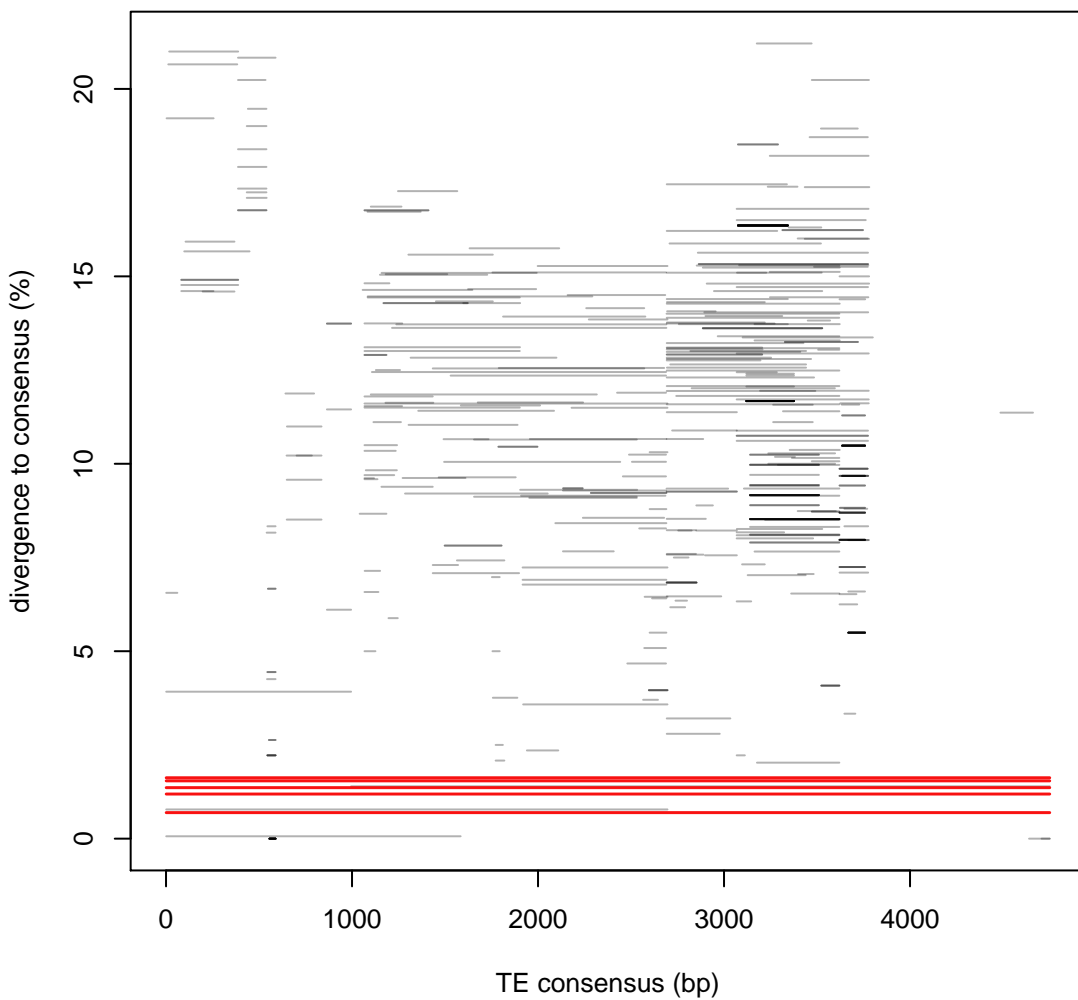
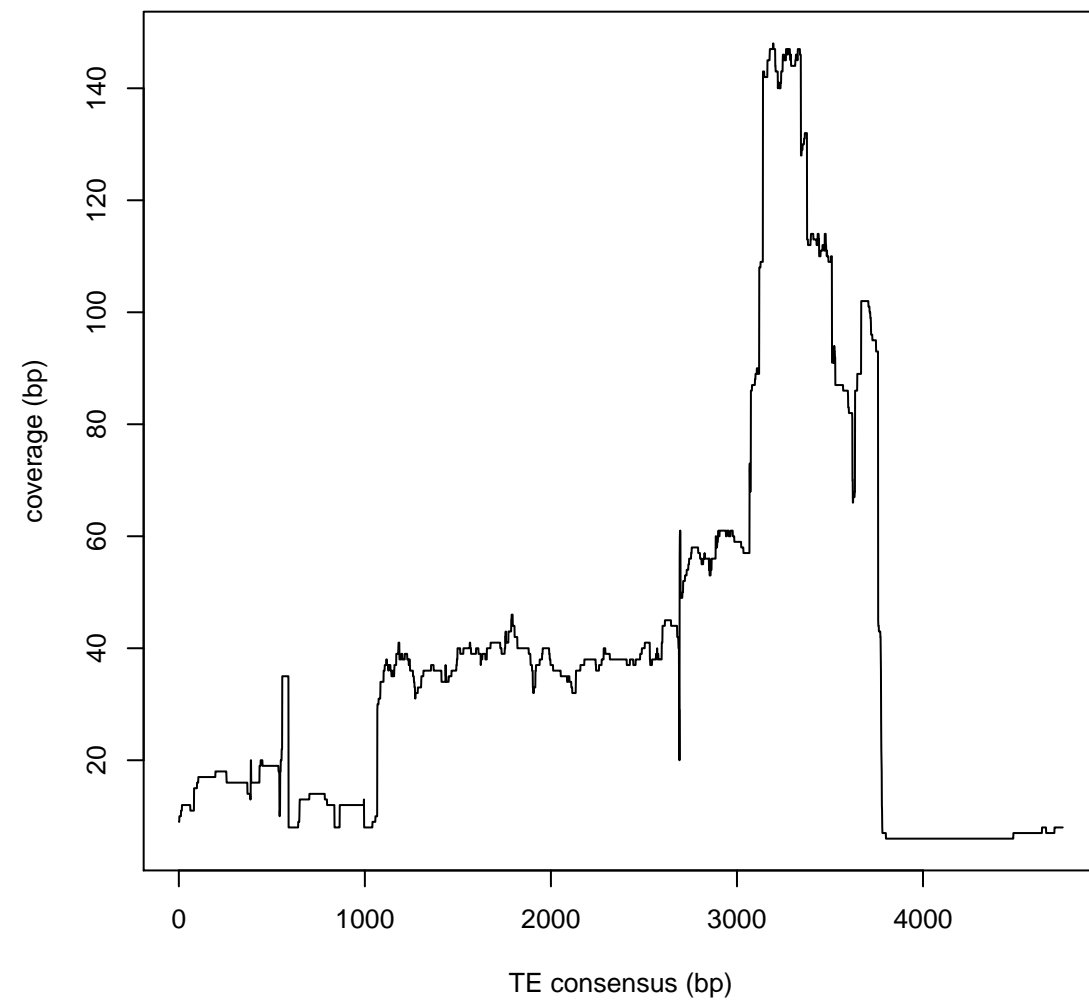


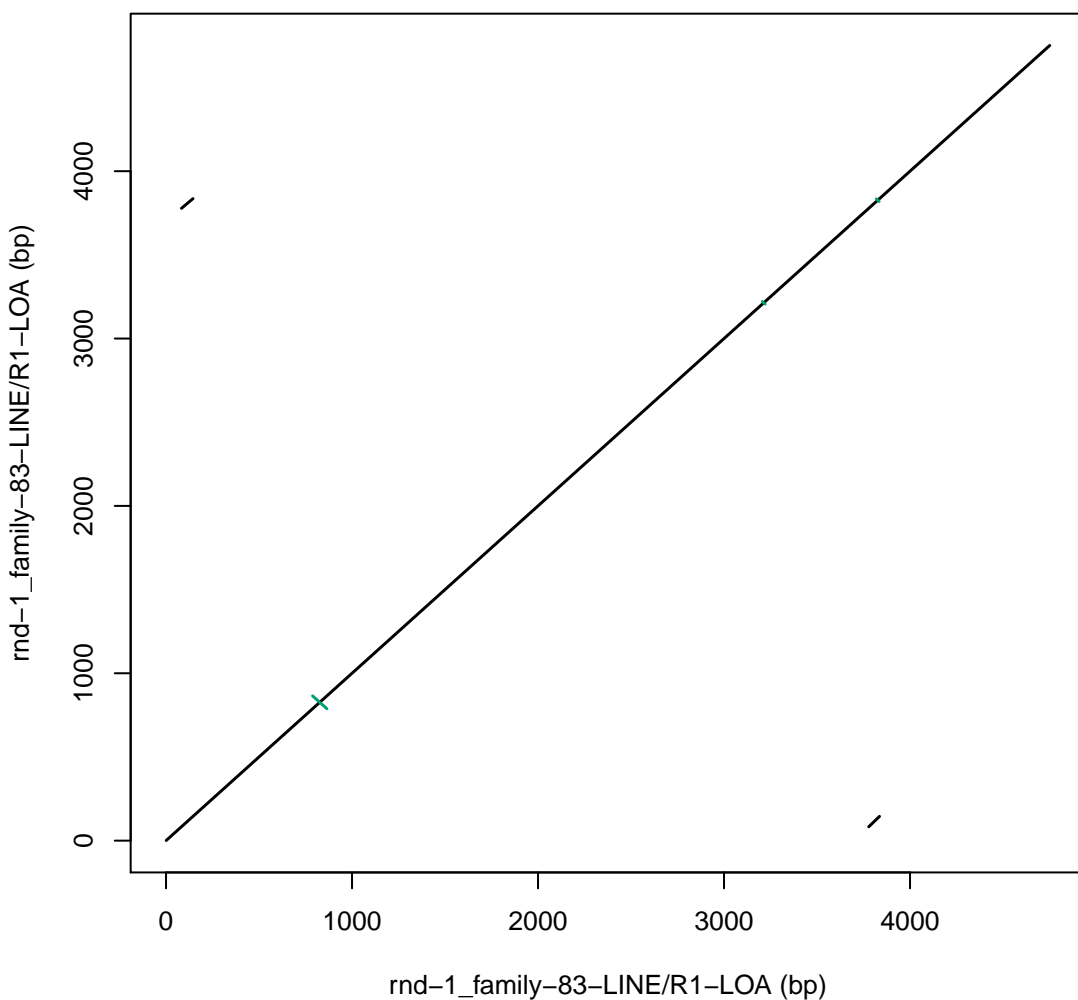
TE: rnd-1_family-83-LINE/R1-LOA
consensus size: 4751bp; fragments: 456; full length: 5 (≥ 4275.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

