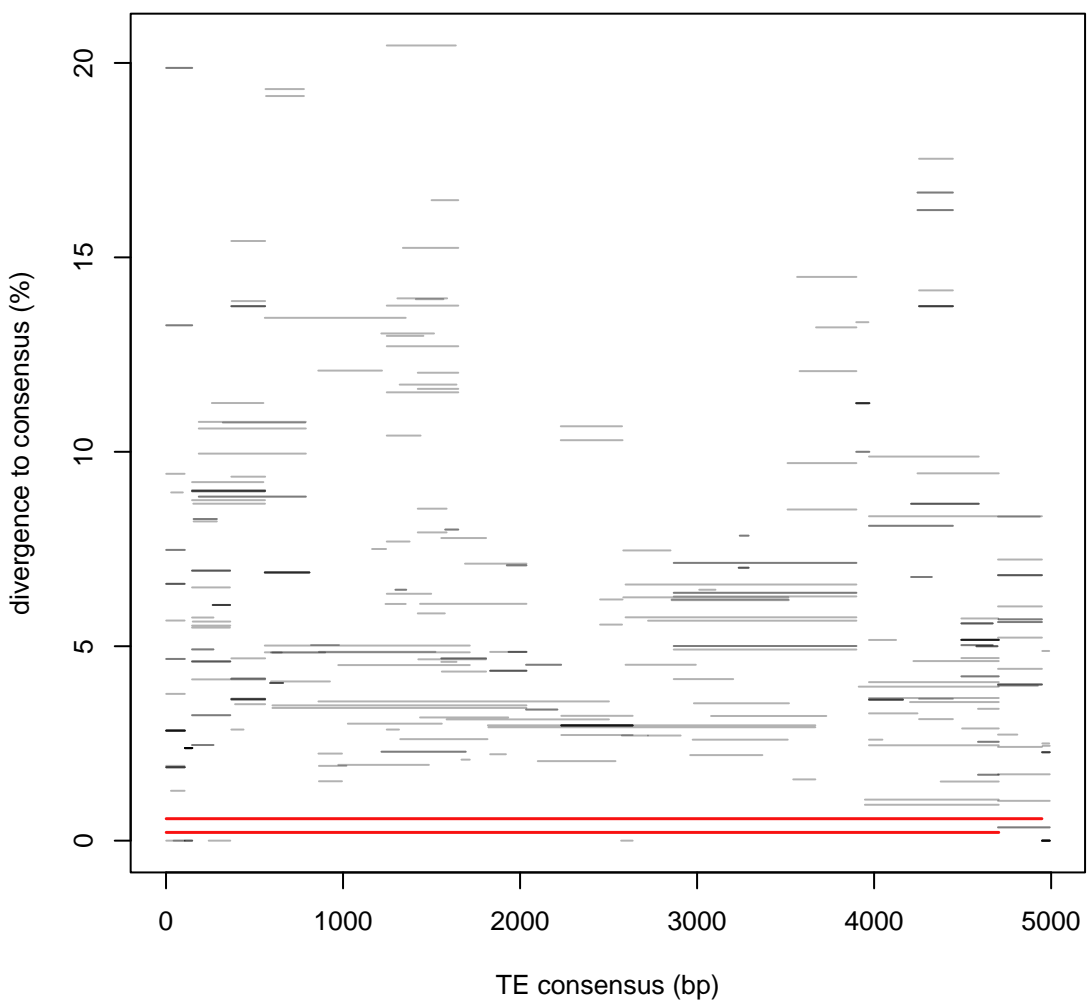
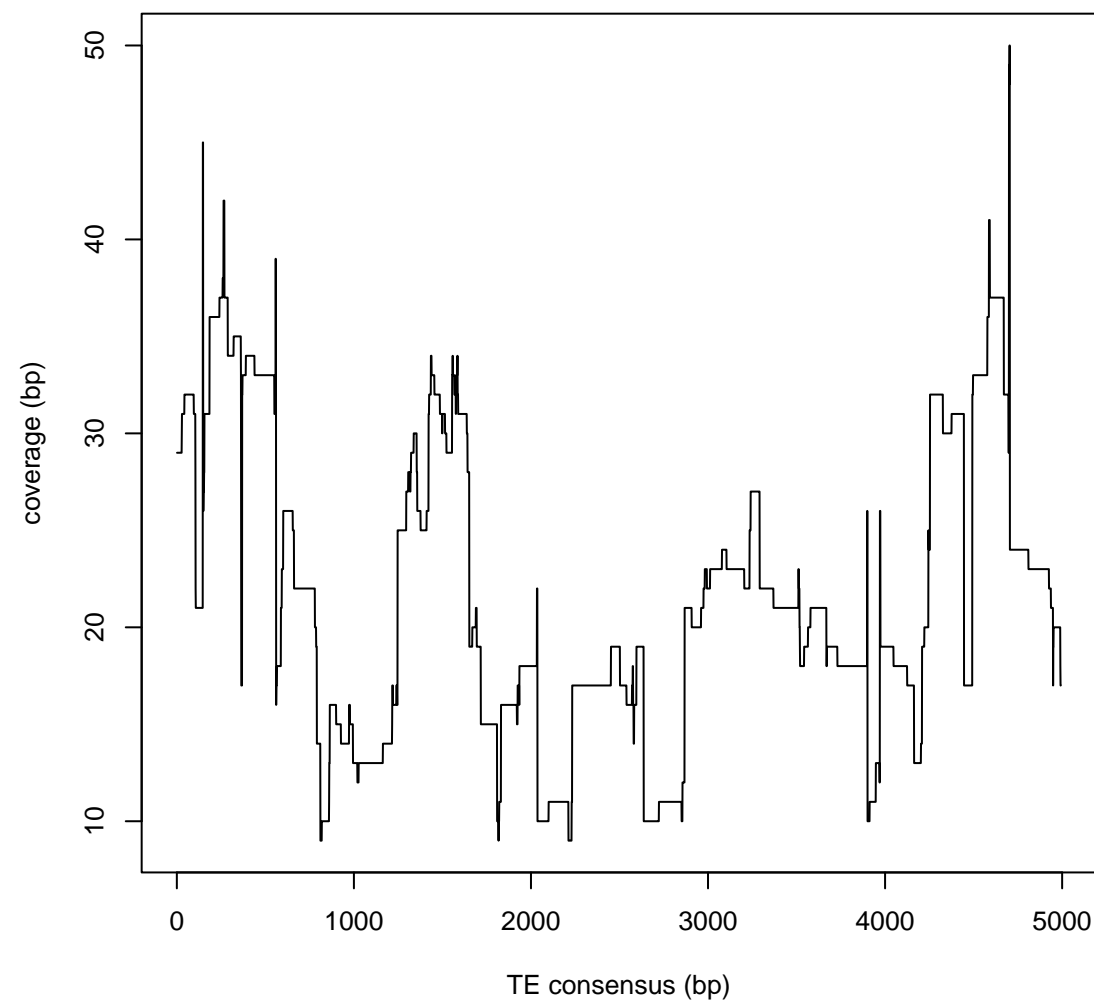


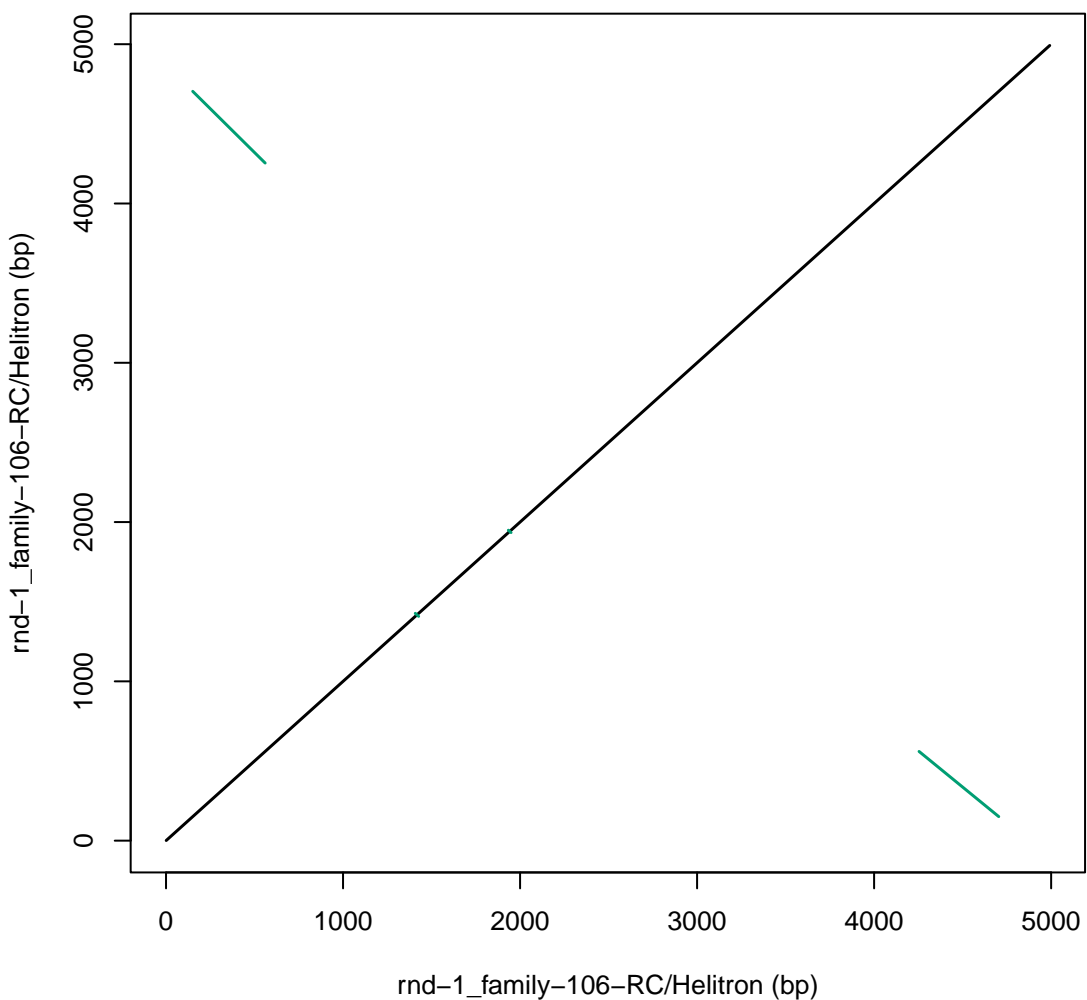
TE: rnd-1_family-106-RC/Helitron
consensus size: 4992bp; fragments: 345; full length: 2 (≥ 4492.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

