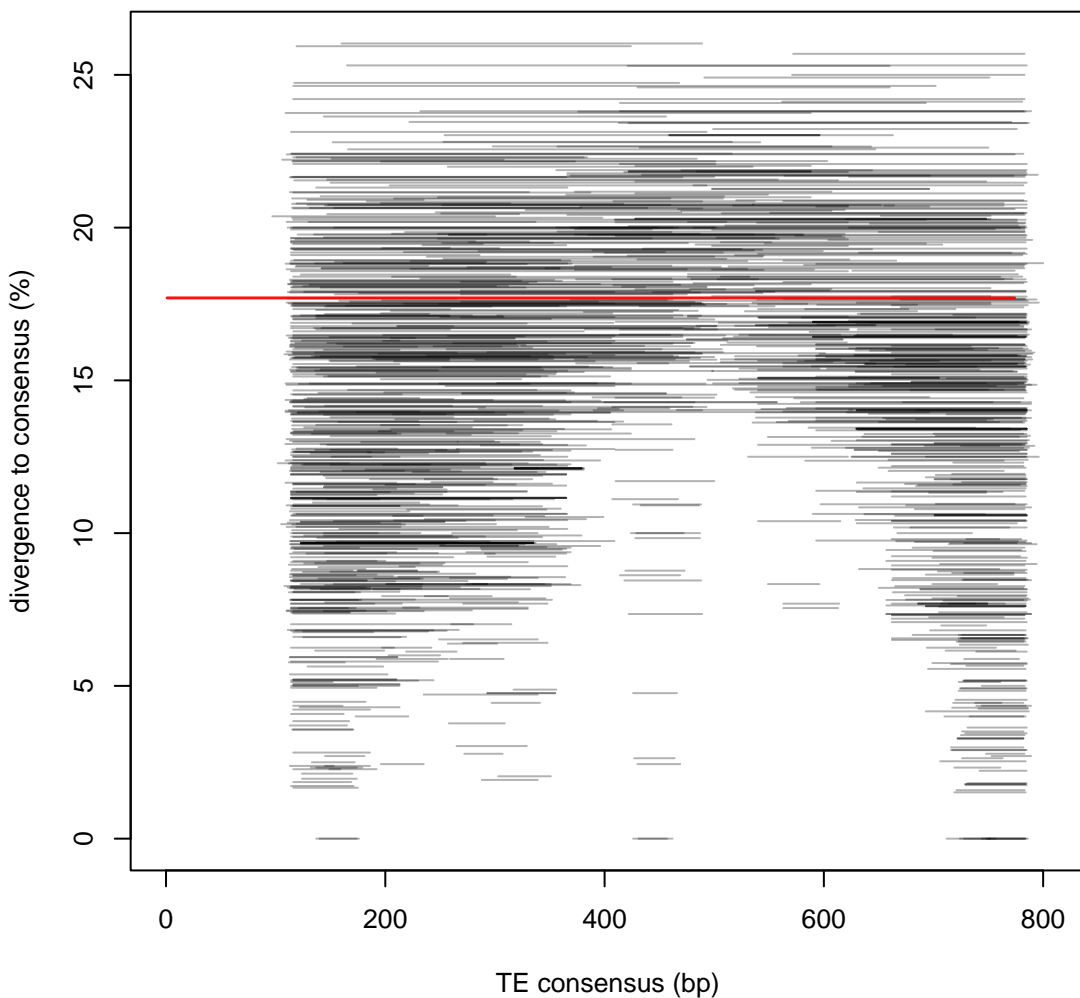
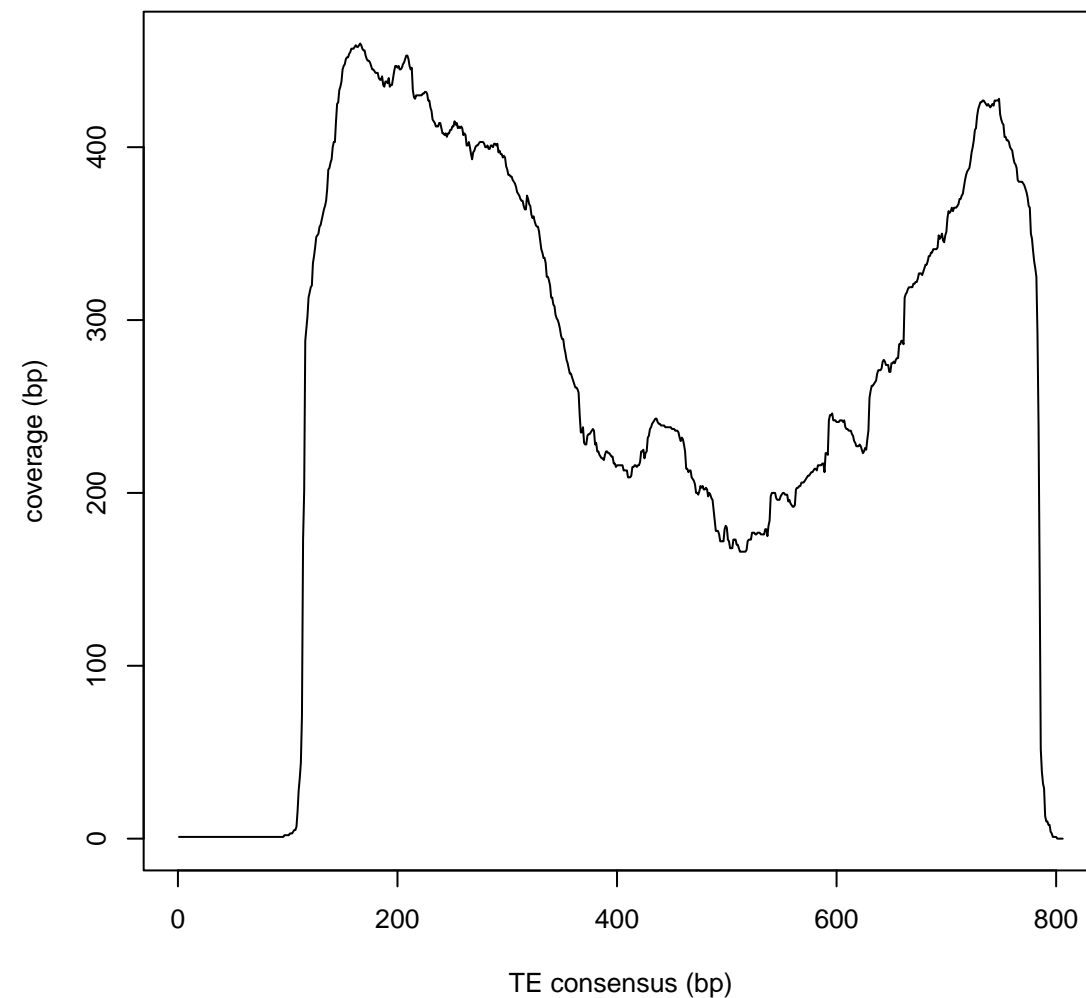


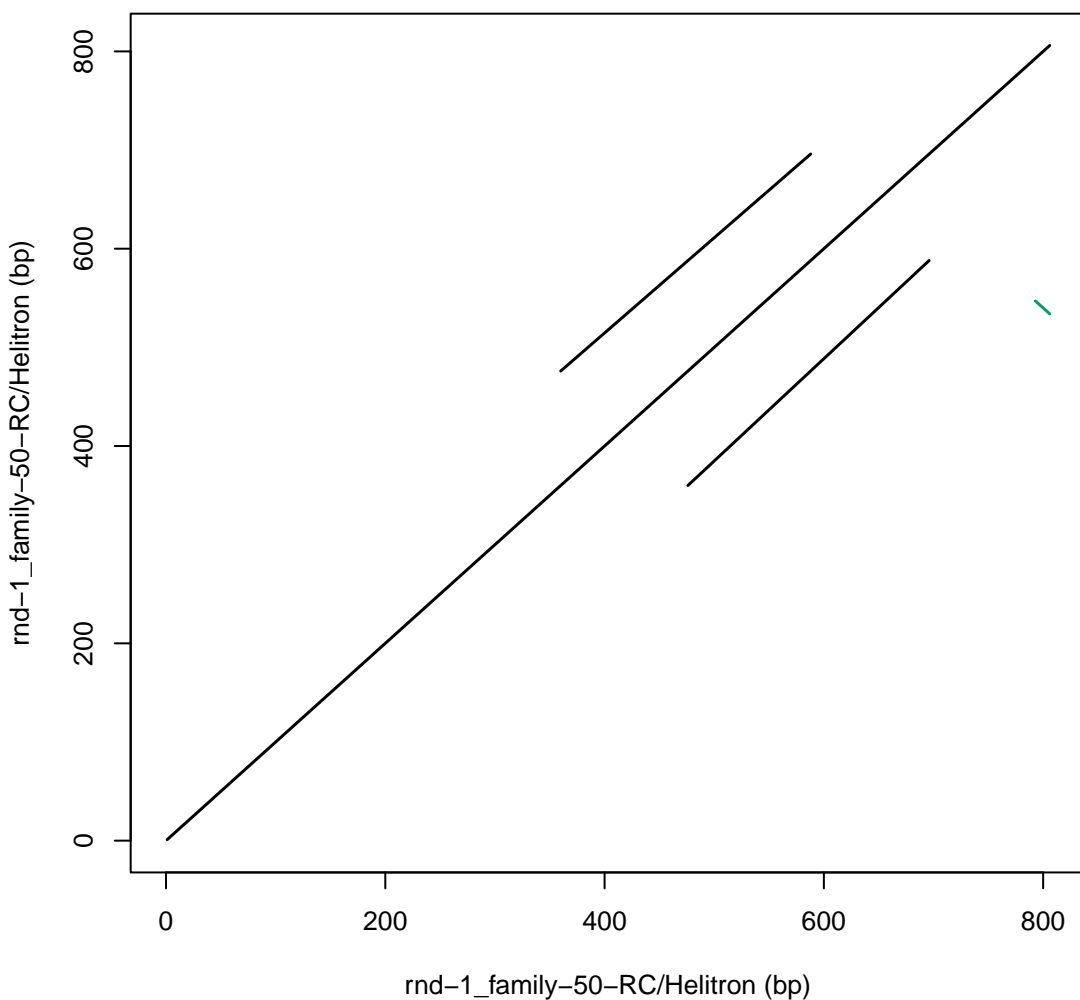
TE: rnd-1_family-50-RC/Helitron
consensus size: 806bp; fragments: 1157; full length: 1 (≥ 725.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

