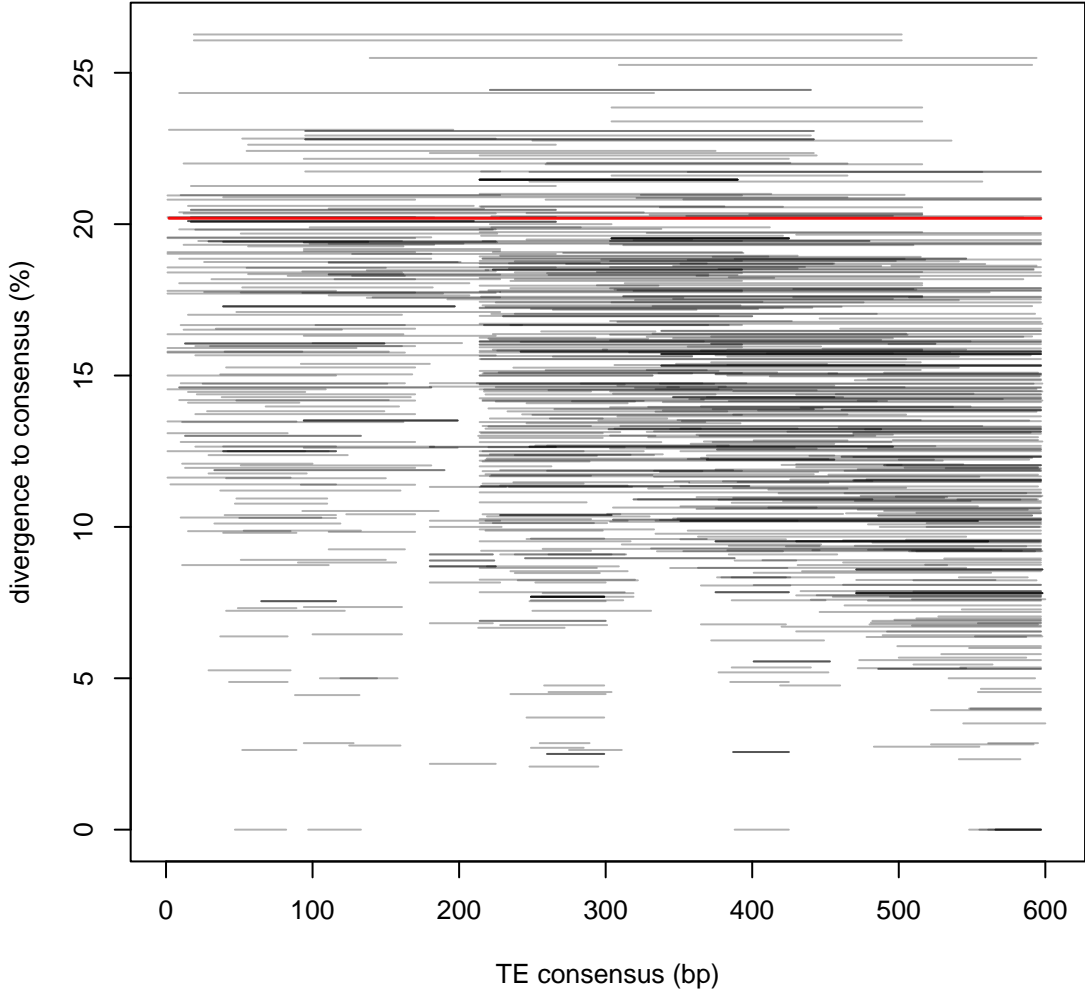
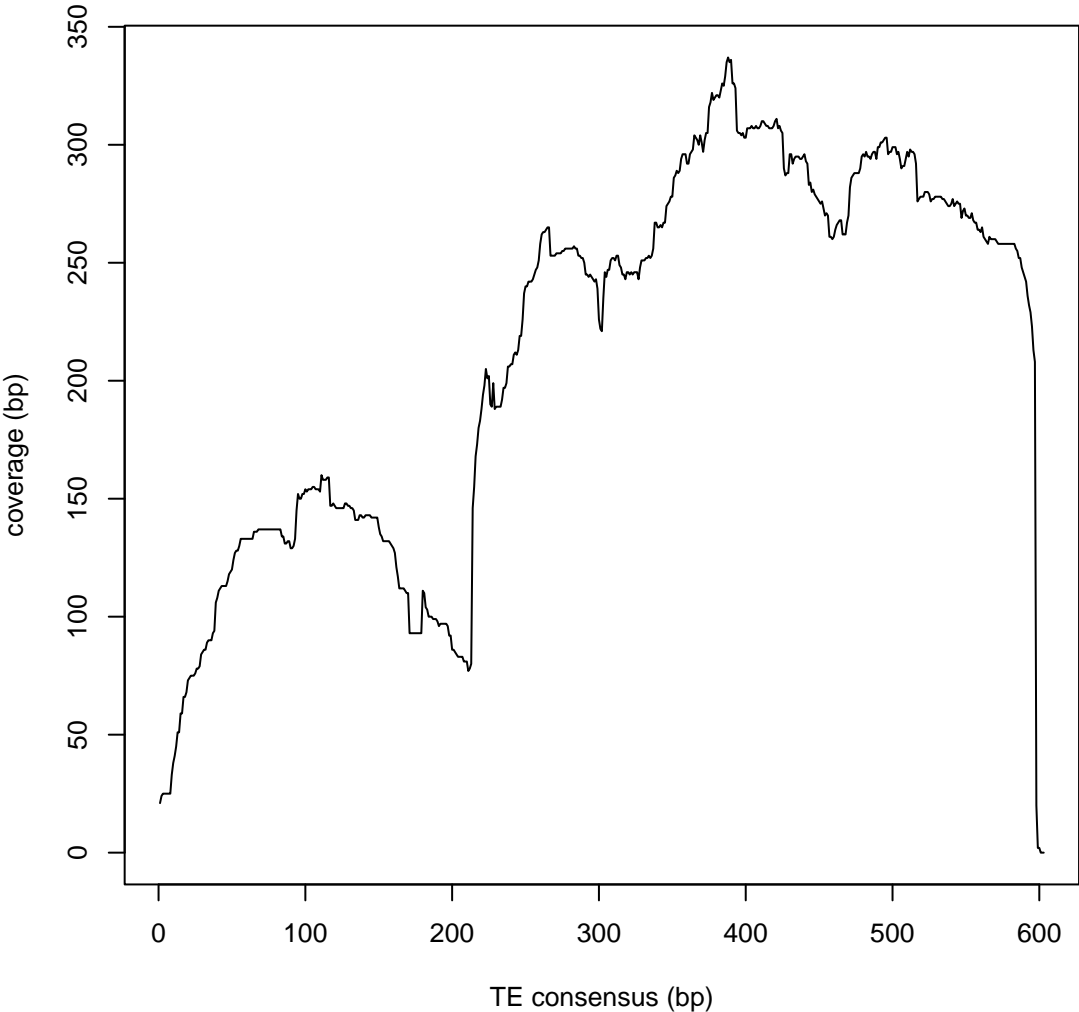


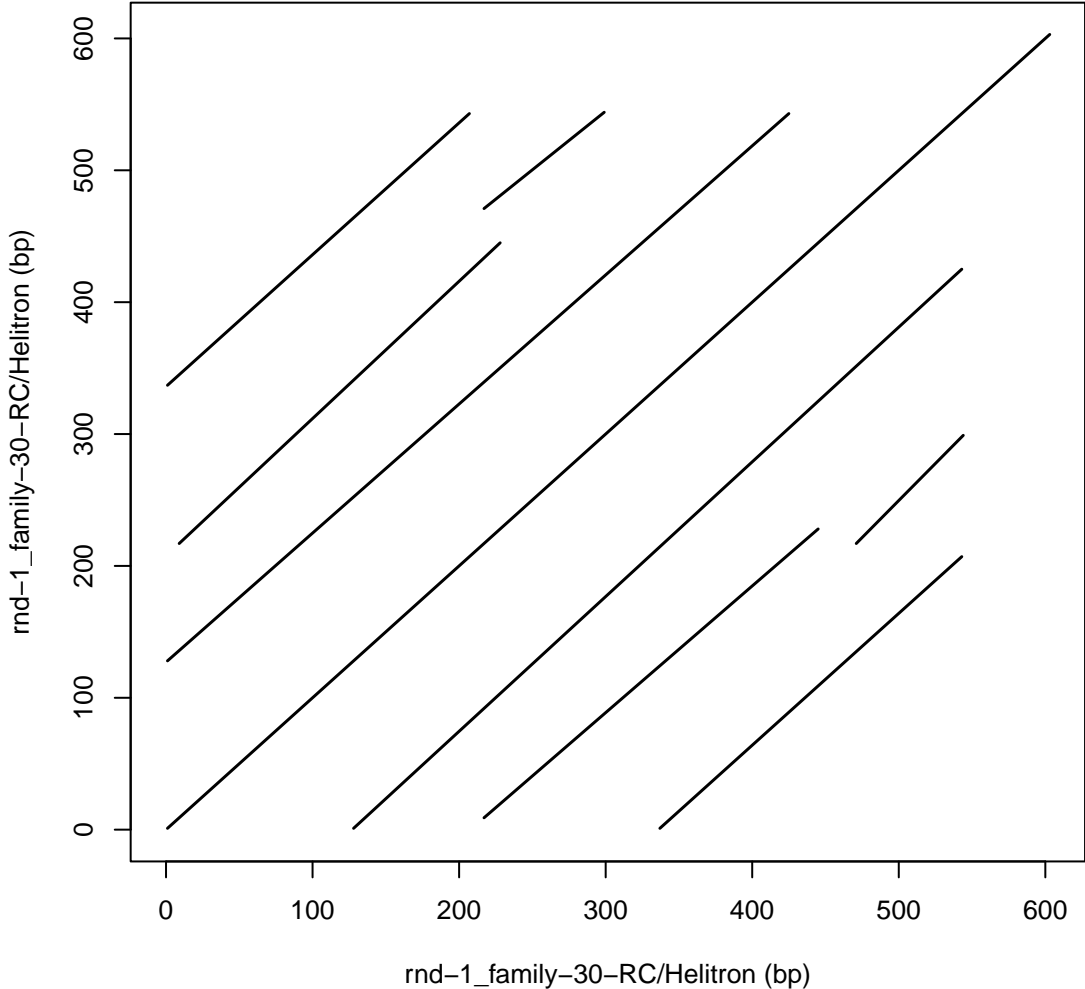
TE: rnd-1_family-30-RC/Helitron
consensus size: 603bp; fragments: 815; full length: 1 (>=542.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

