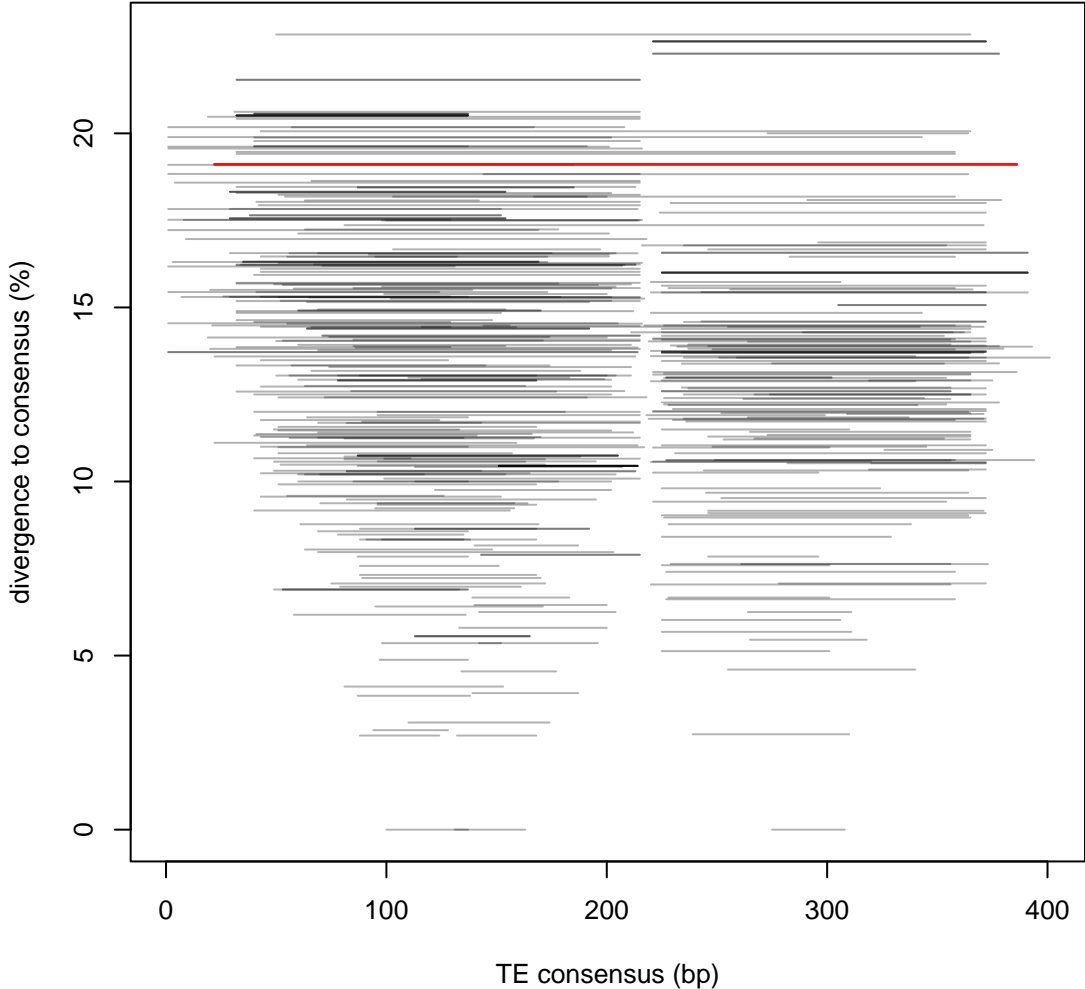
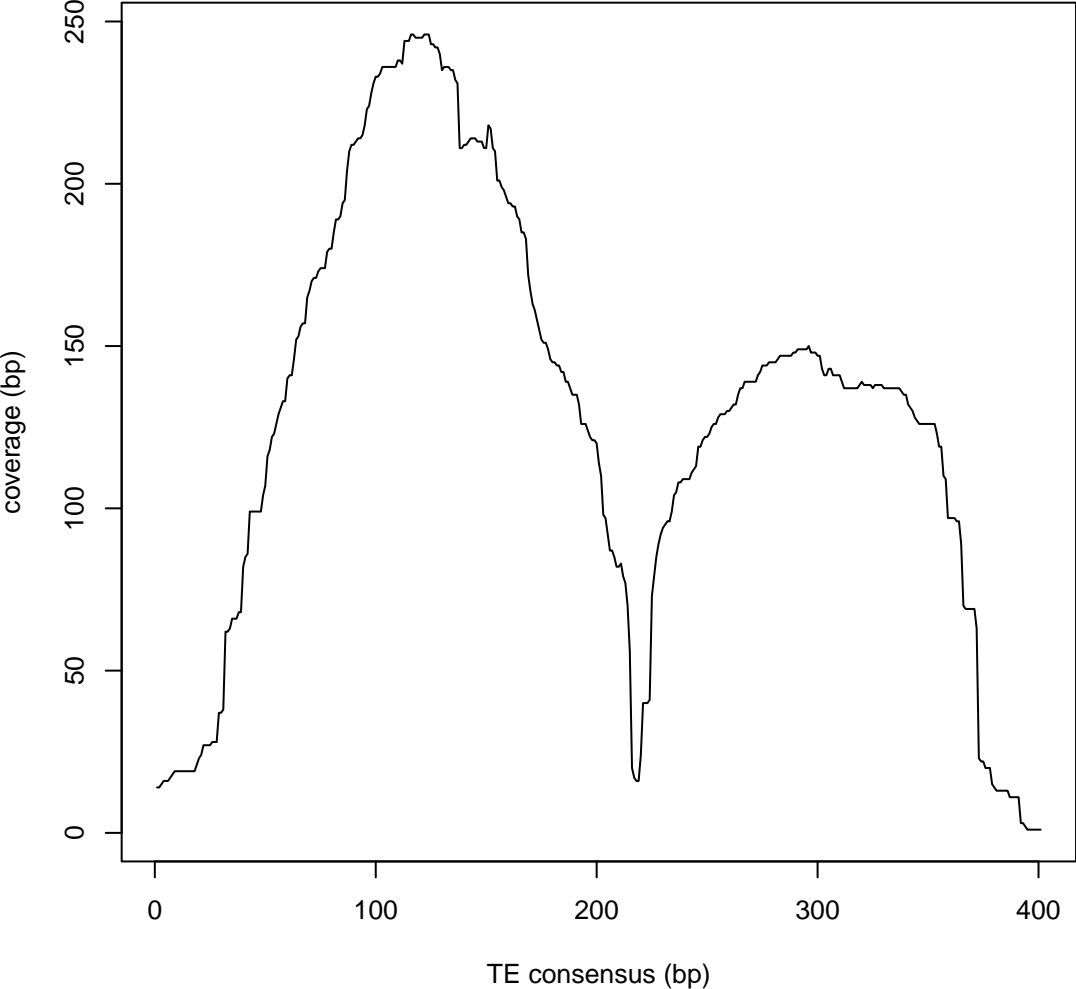


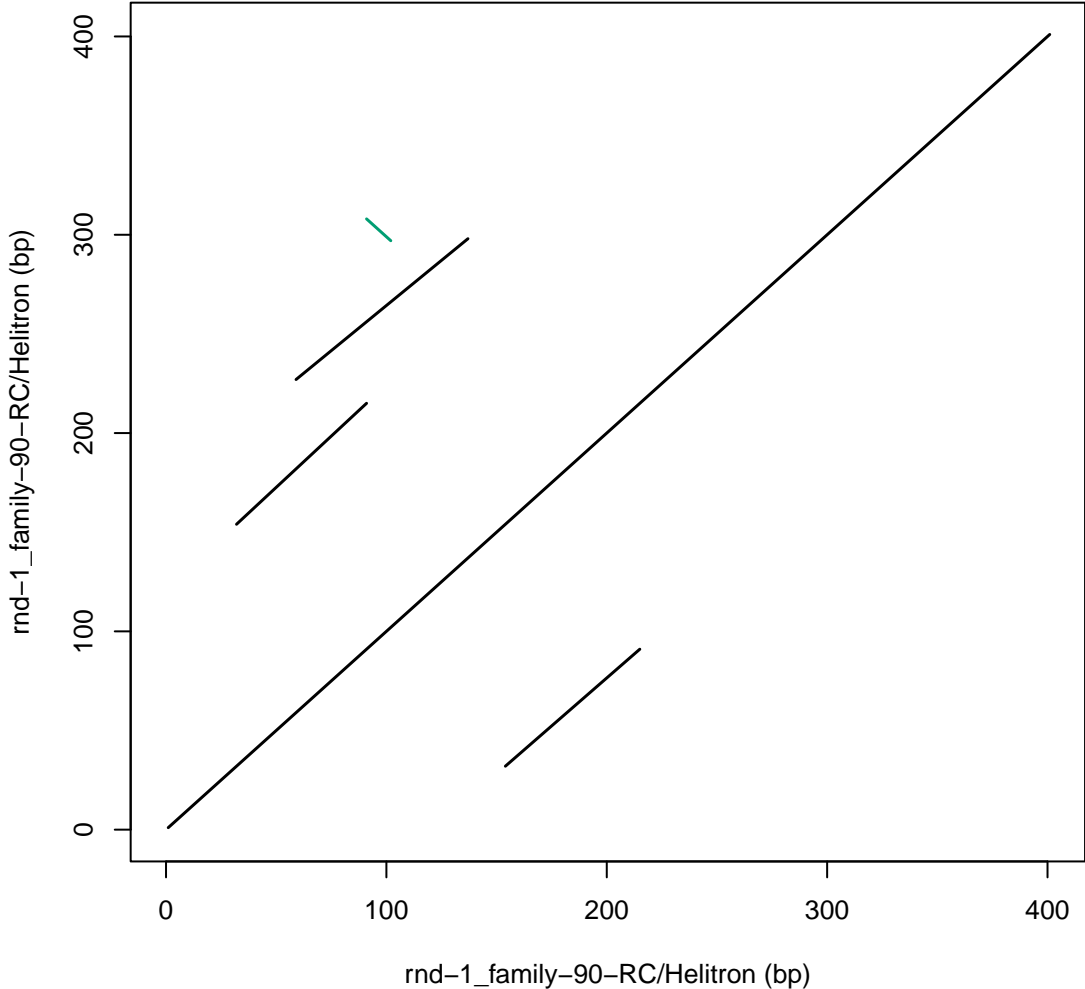
TE: rnd-1_family-90-RC/Helitron
consensus size: 401bp; fragments: 417; full length: 1 (>=360.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

