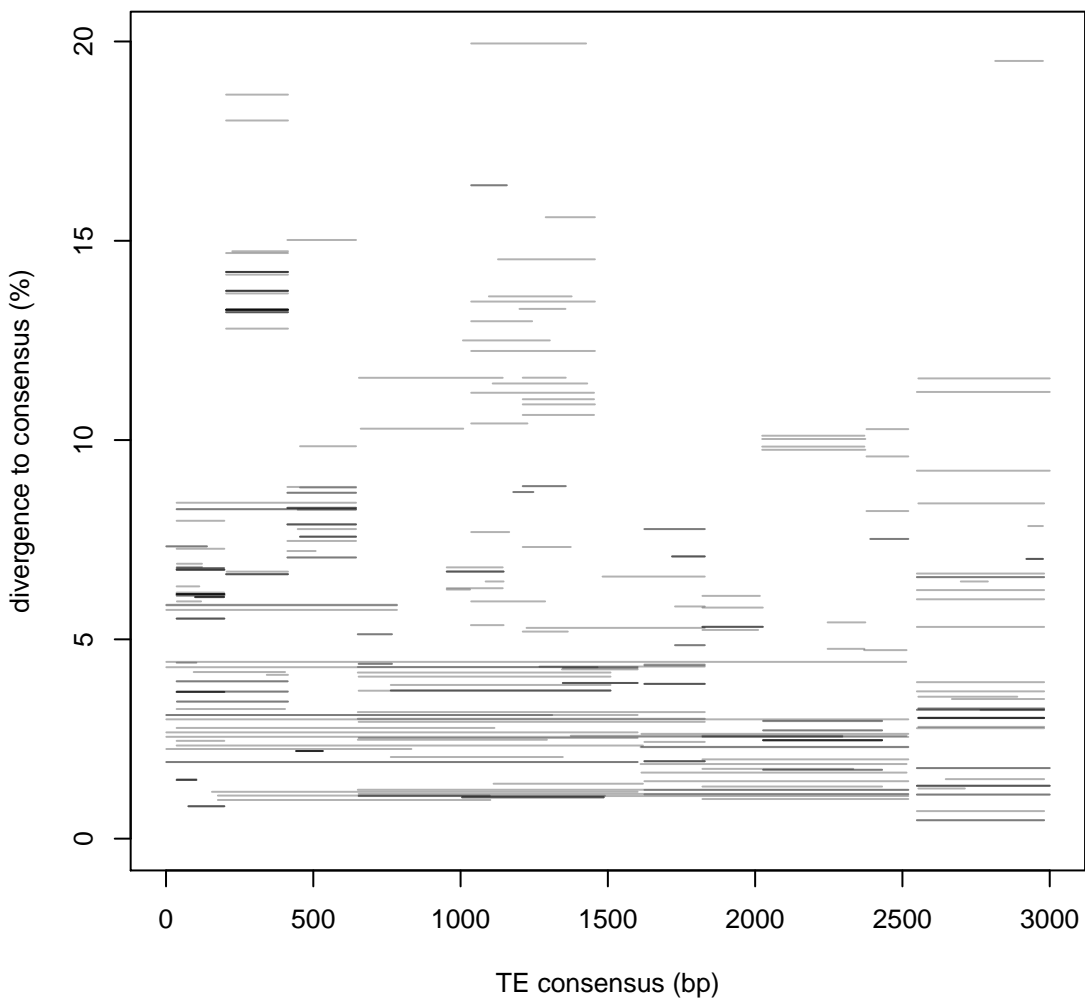
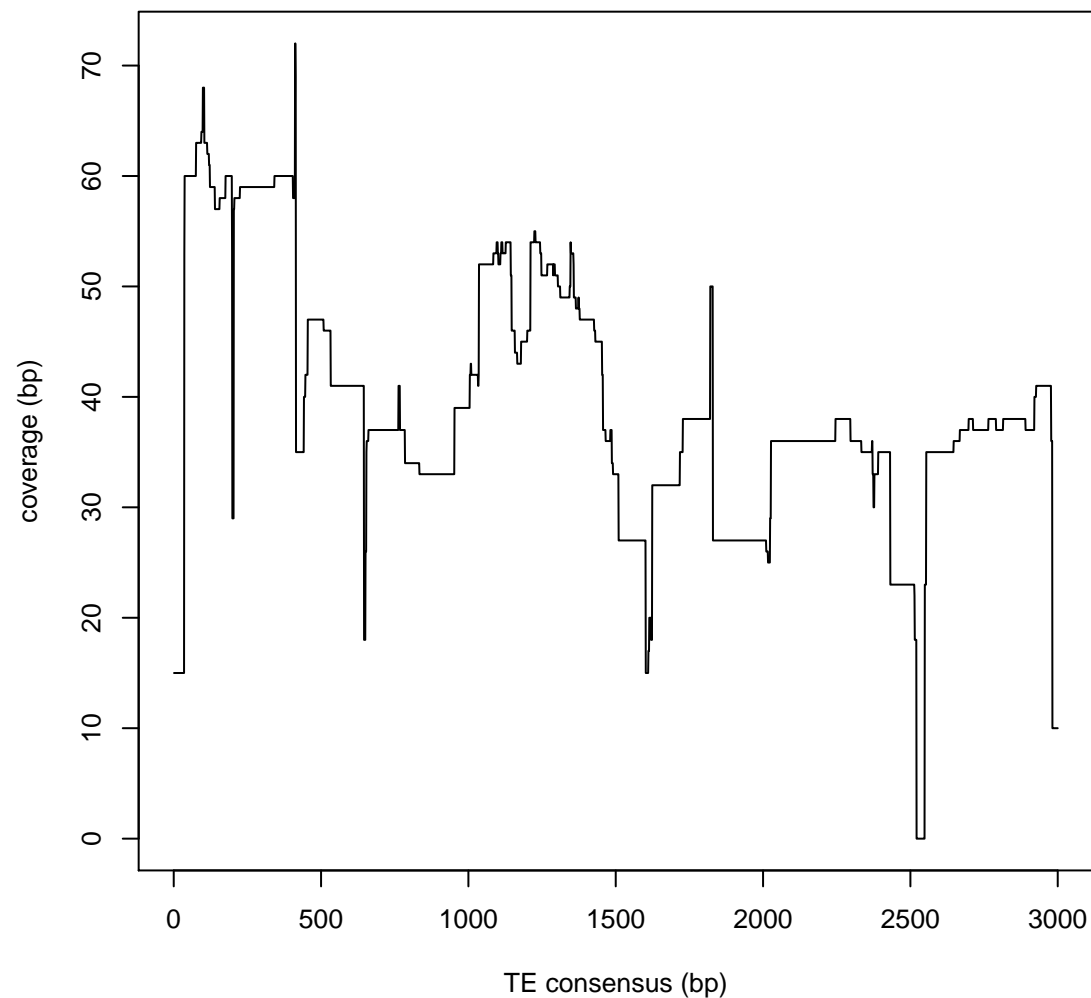


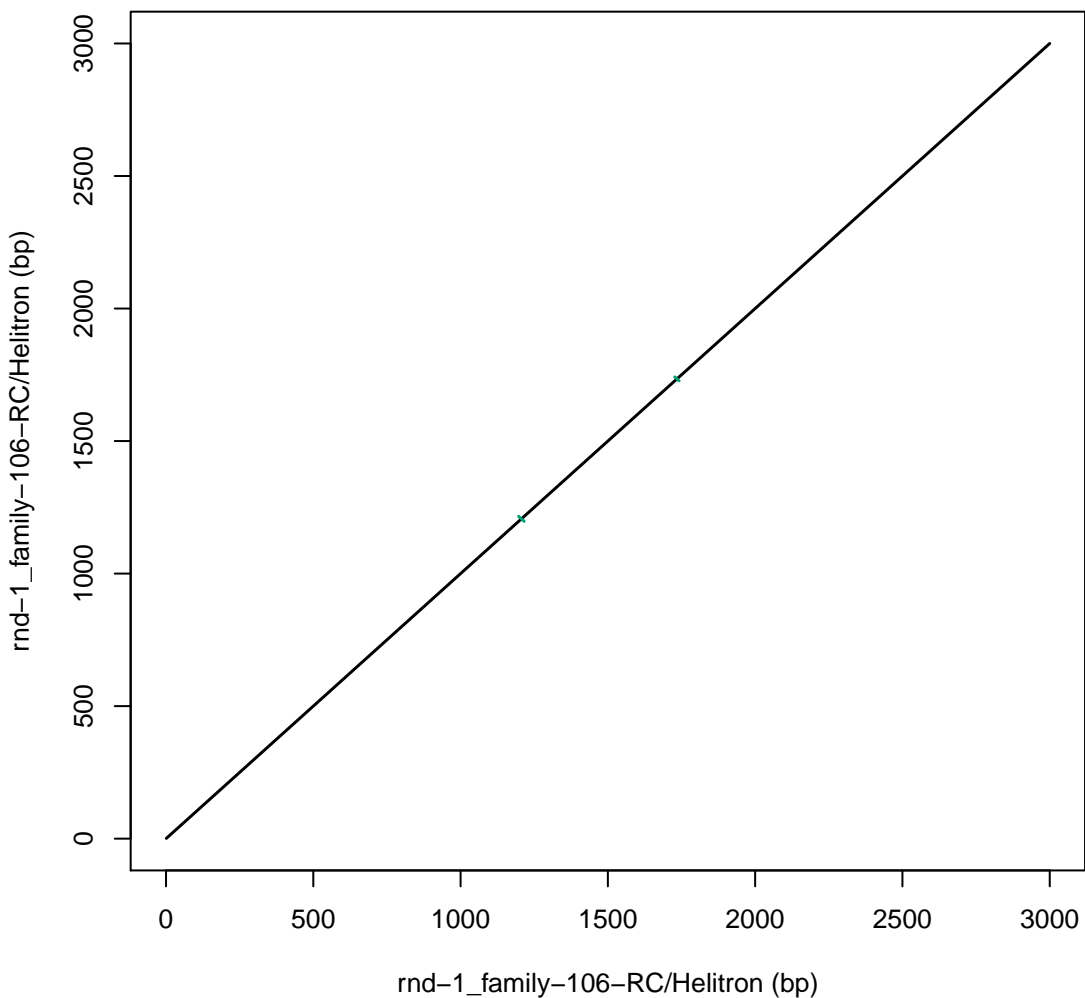
TE: rnd-1_family-106-RC/Helitron
consensus size: 3000bp; fragments: 302; full length: 0 (>=2700bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

