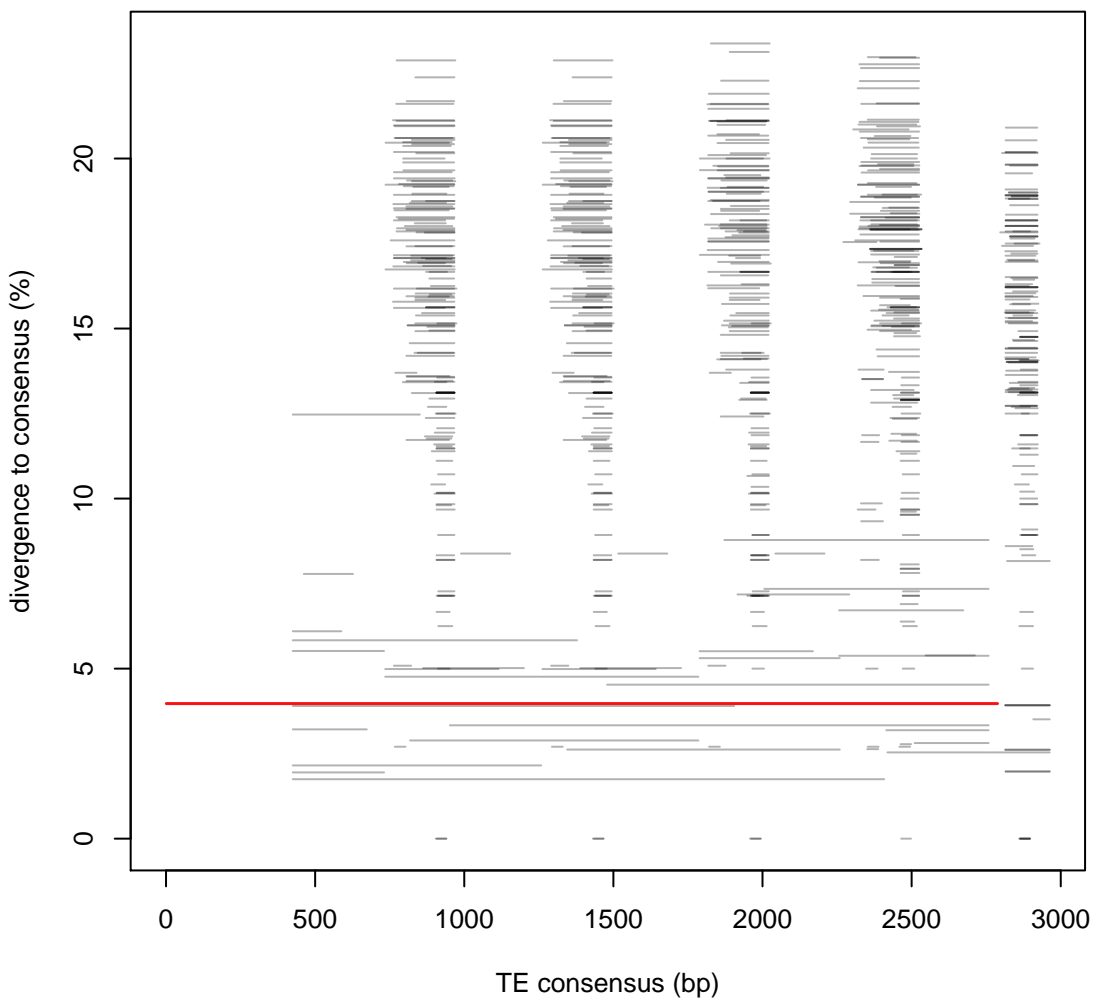
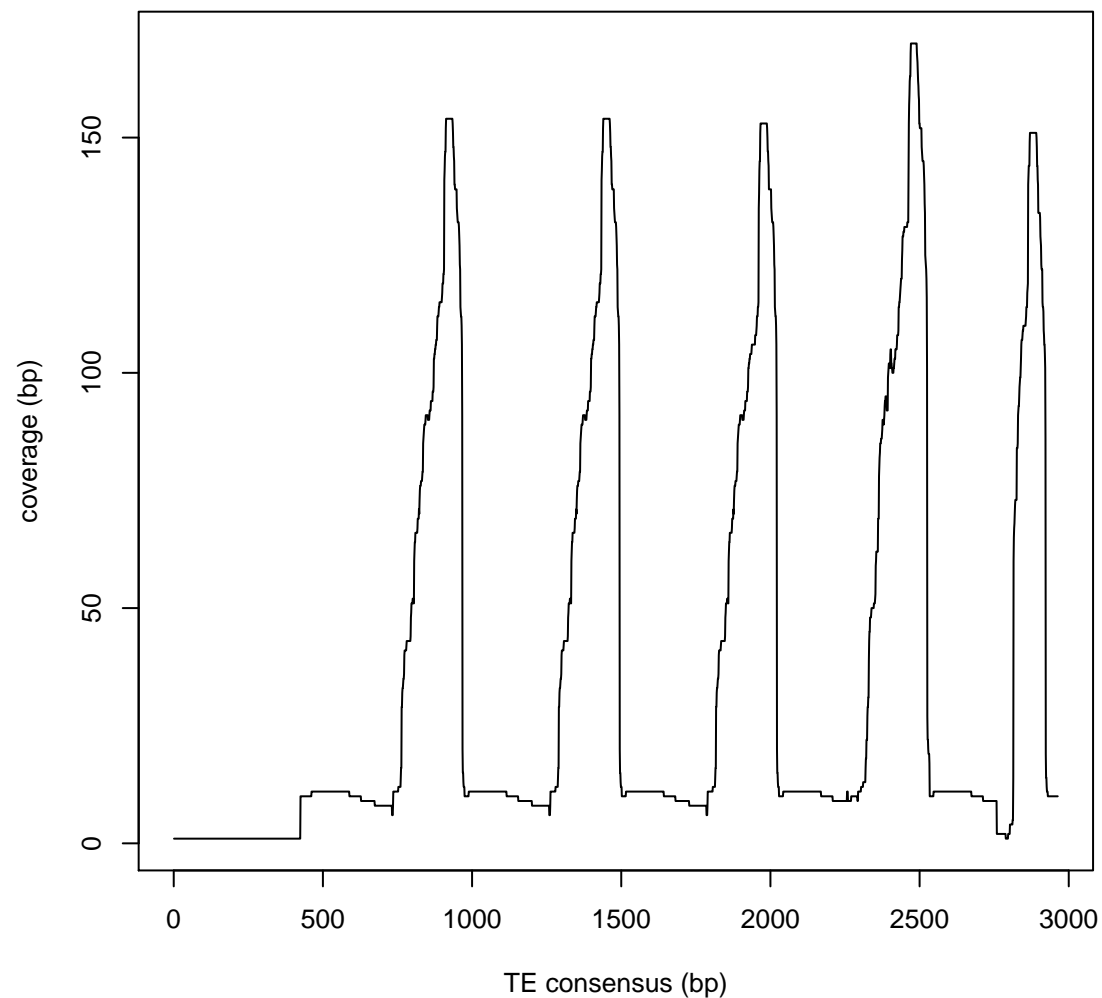


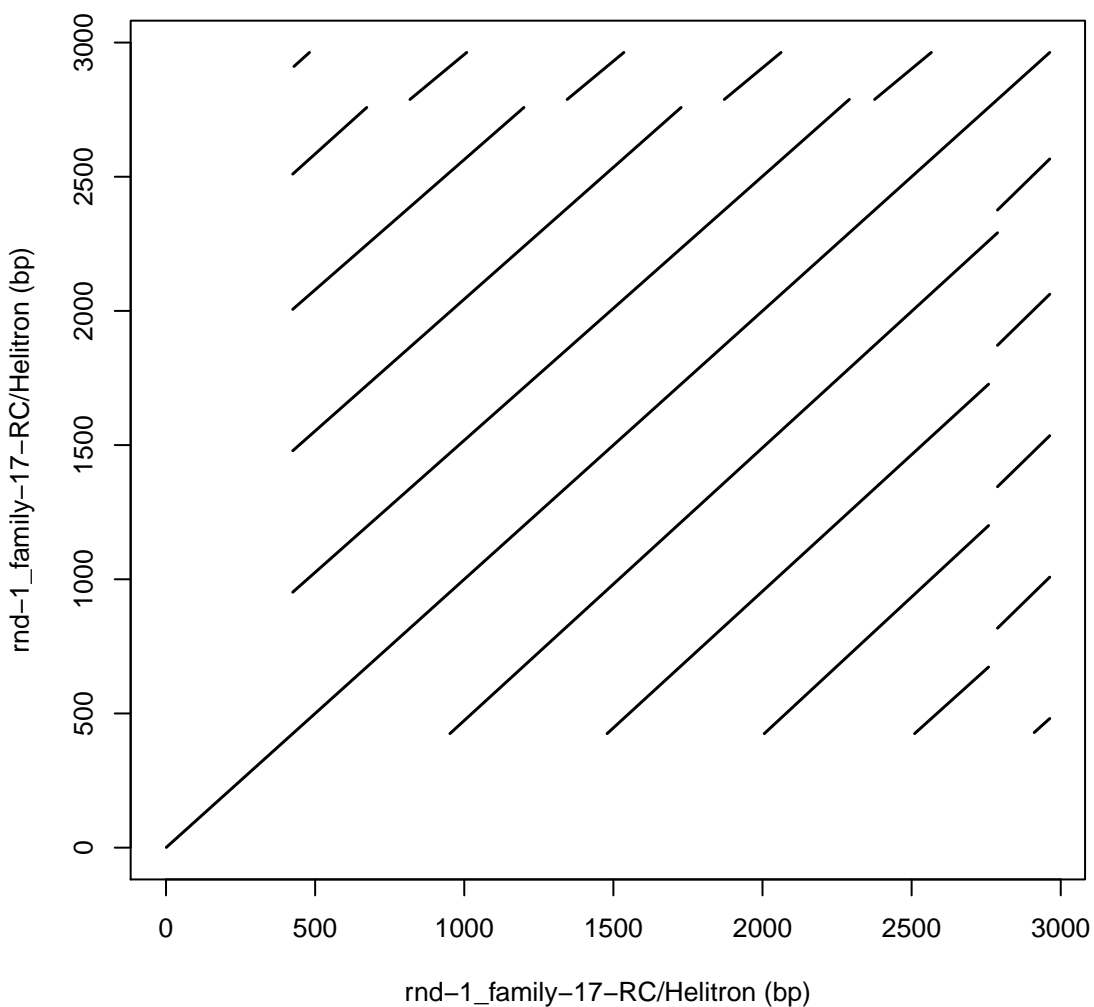
TE: rnd-1_family-17-RC/Helitron
consensus size: 2963bp; fragments: 803; full length: 1 (>=2666.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

