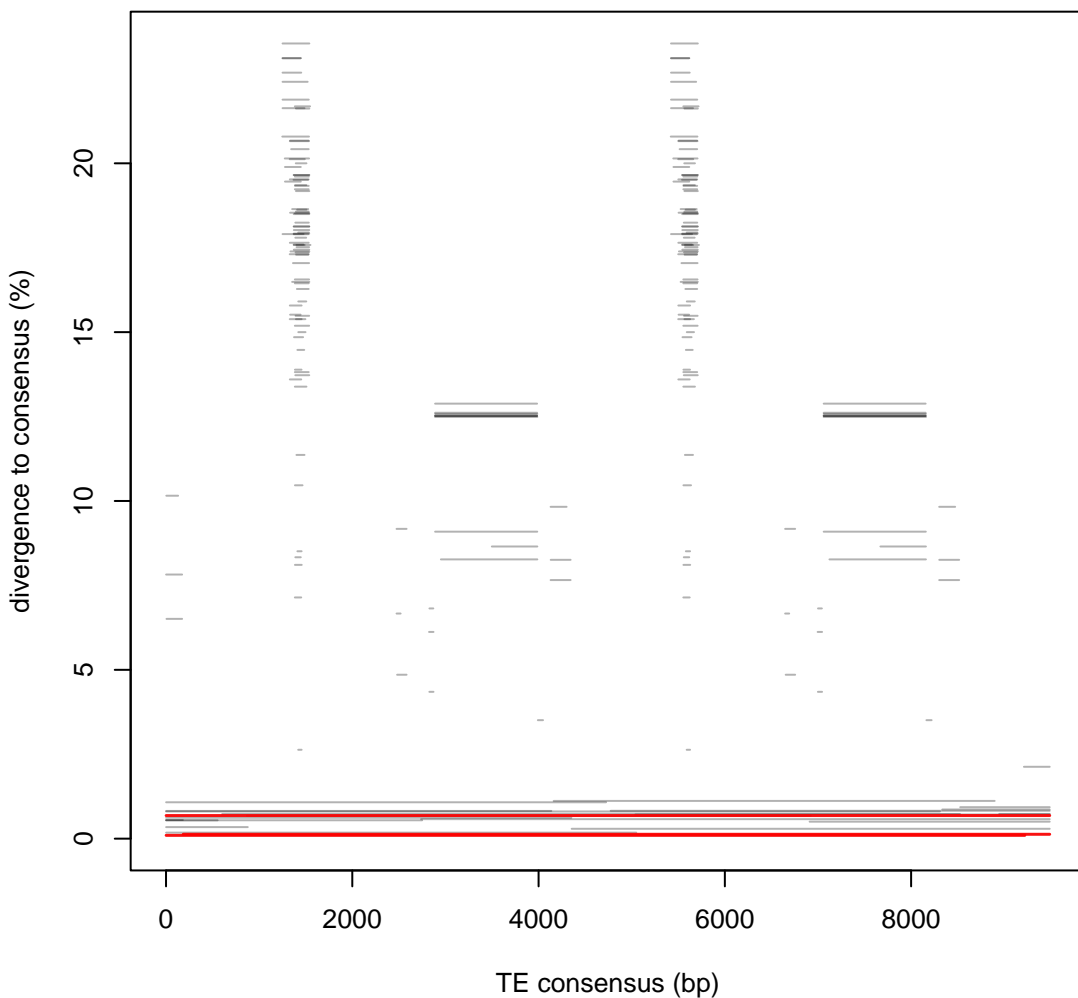
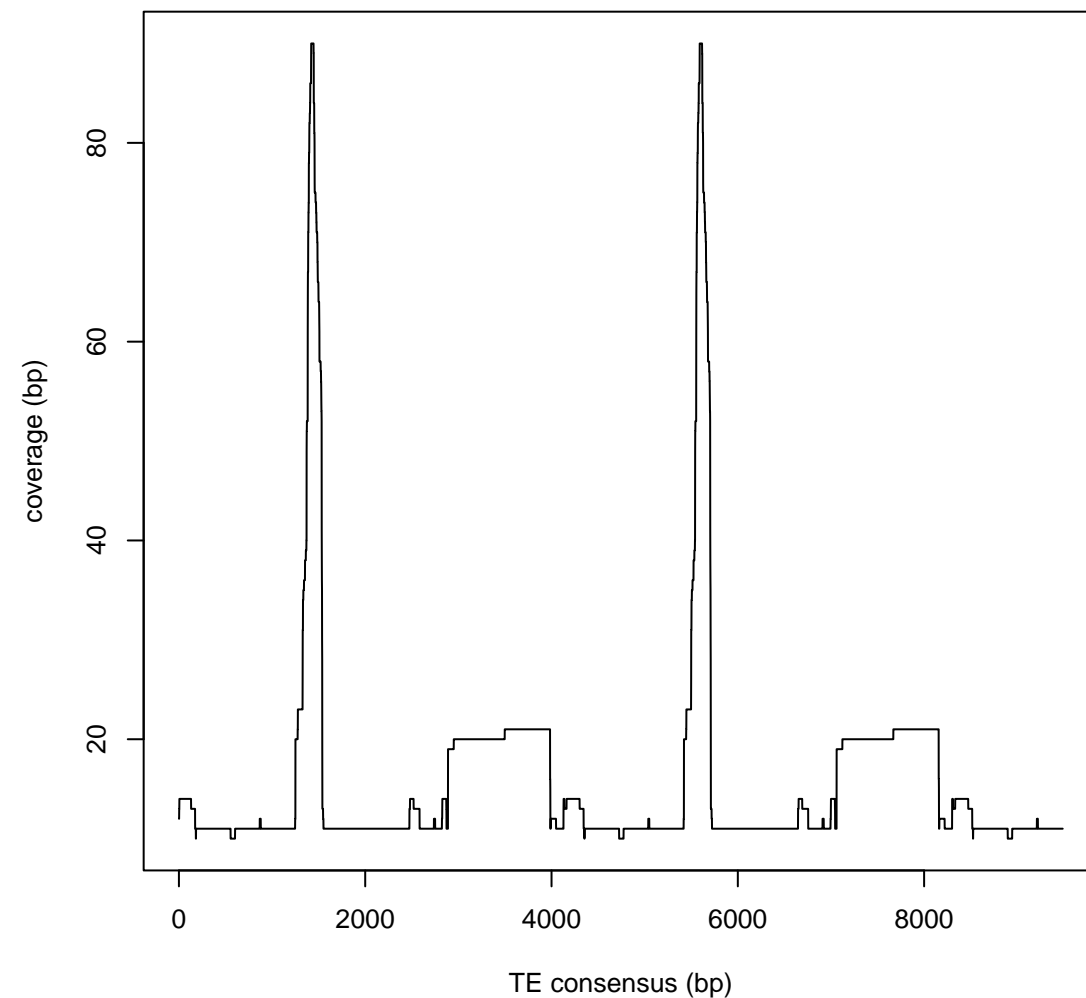


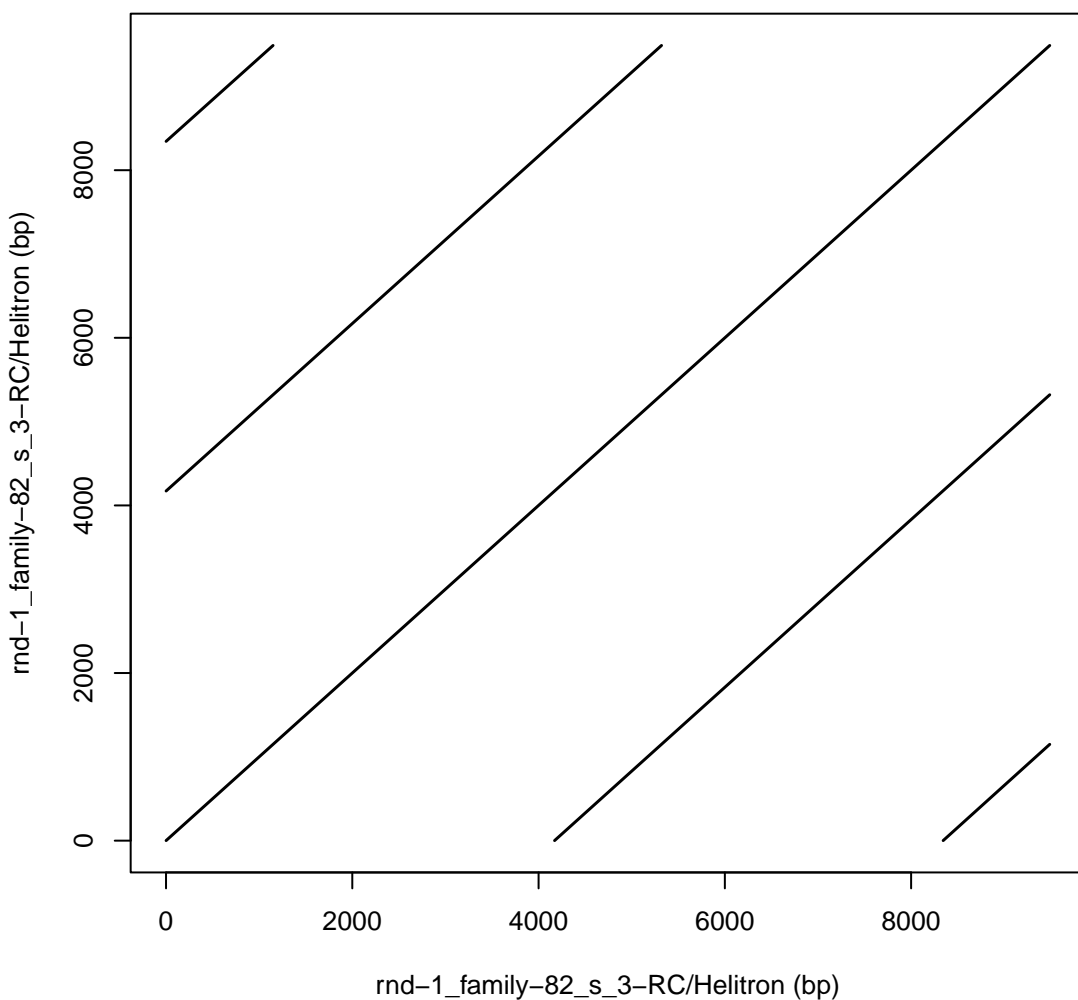
TE: rnd-1_family-82_s_3-RC/Helitron
consensus size: 9488bp; fragments: 225; full length: 3 (>=8539.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

