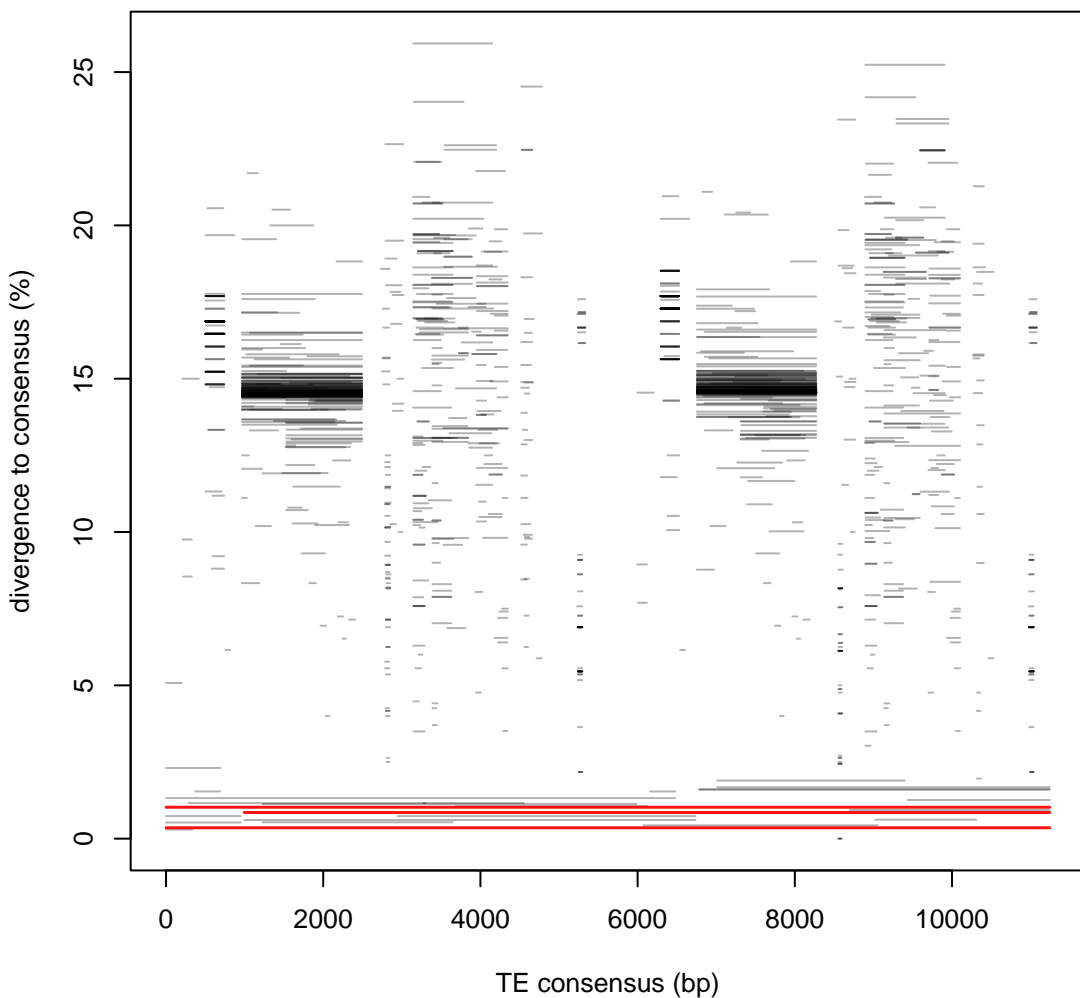
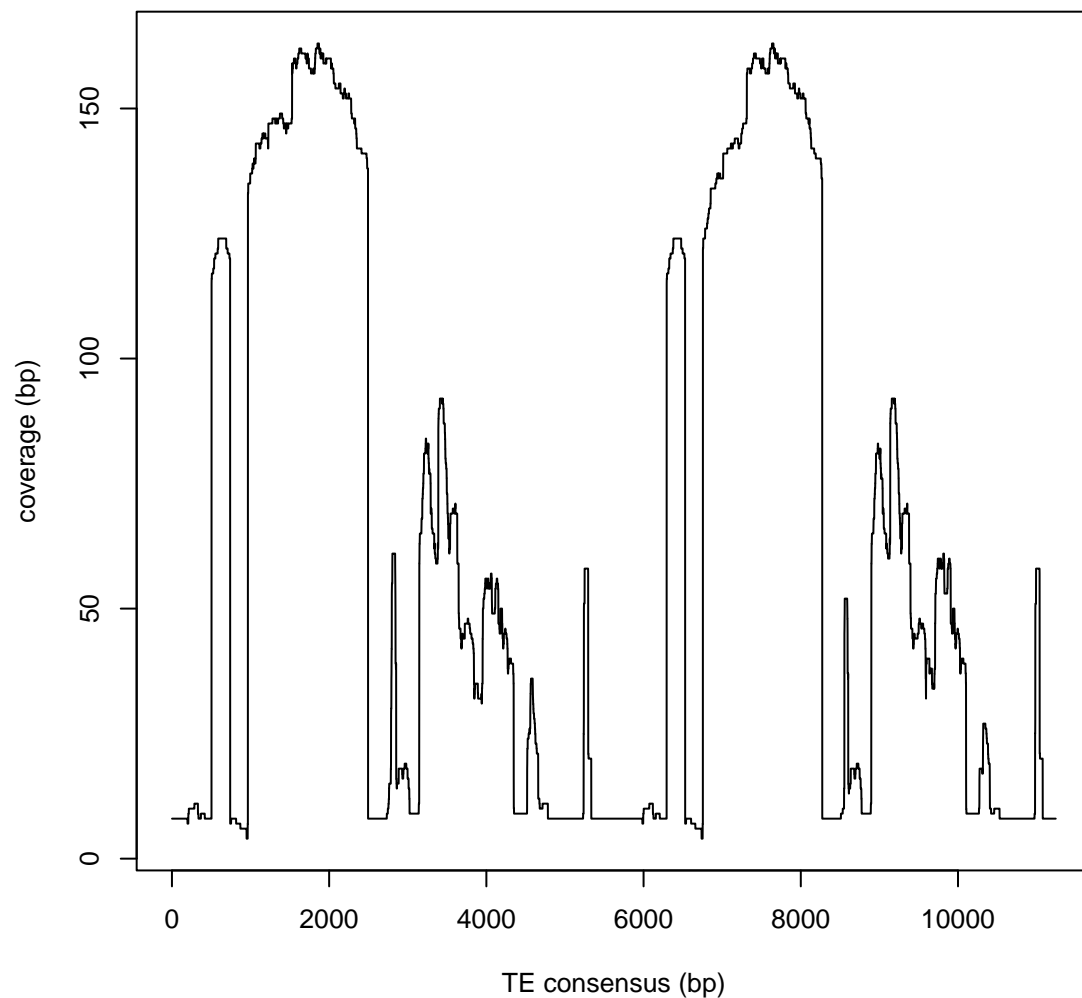


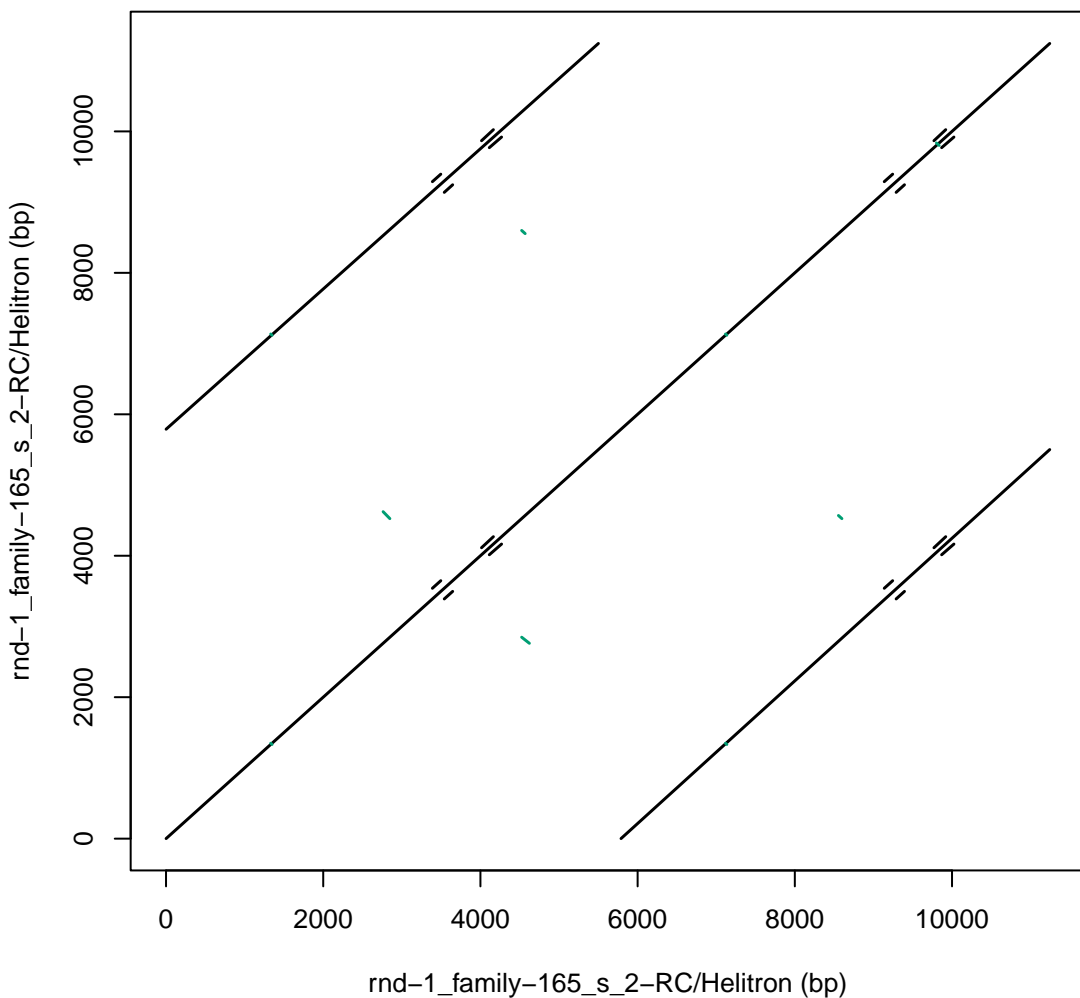
TE: rnd-1_family-165_s_2-RC/Helitron
consensus size: 11243bp; fragments: 1298; full length: 3 (>=10118.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

