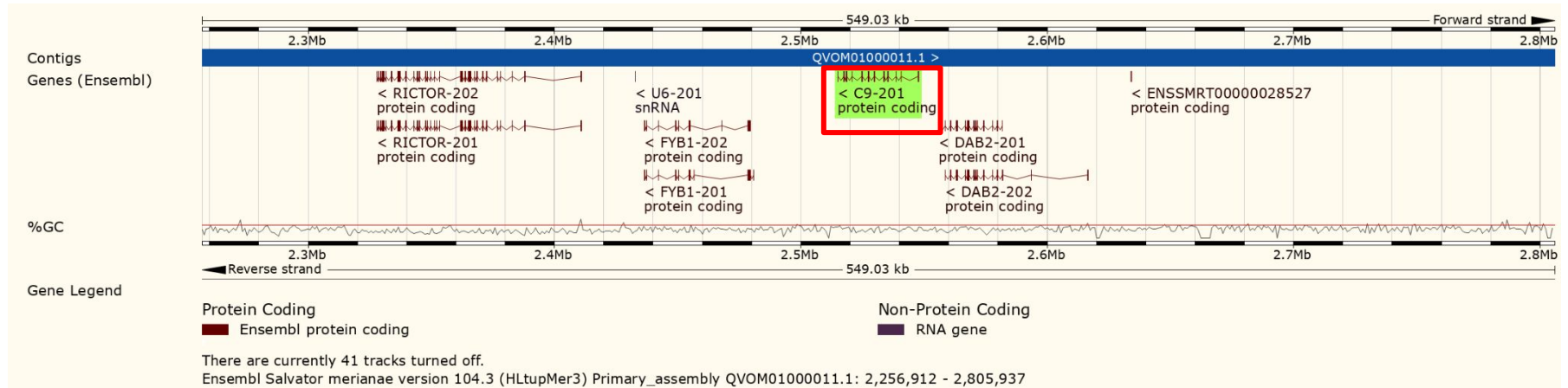
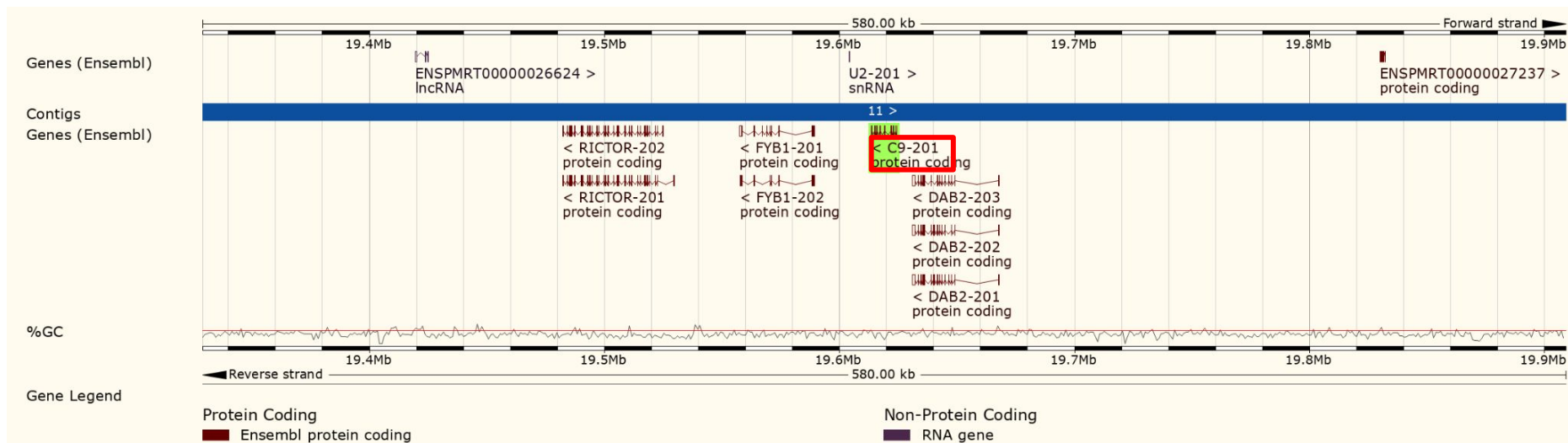


C9 Synteny in Squamata

Argentine Black and white Tegu



Common Wall Lizard



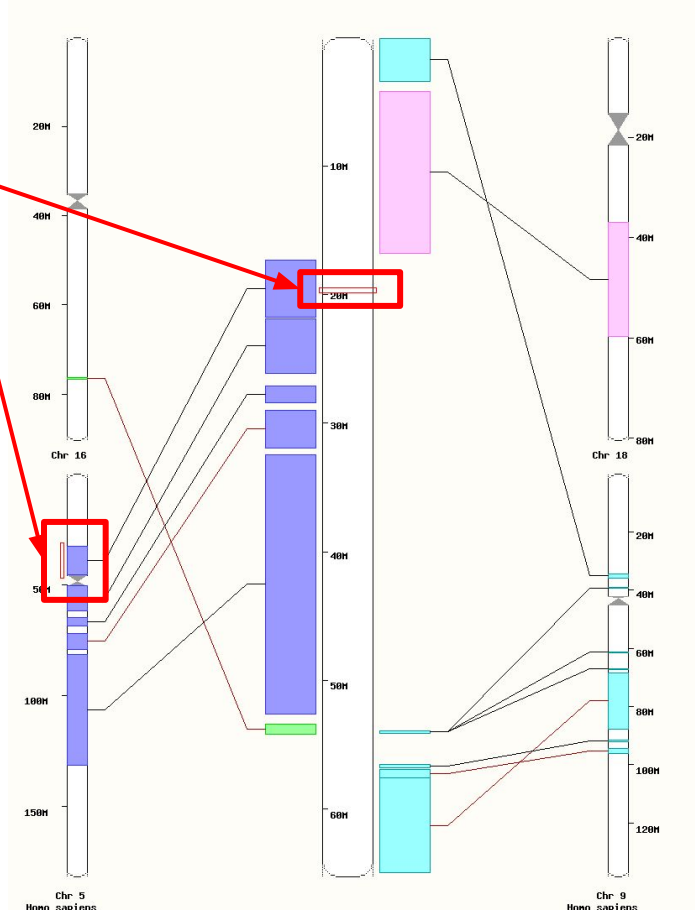
There are currently 32 tracks turned off.

Ensembl Podarcis muralis version 104.1 (PodMur_1.0) Primary_assembly 11: 19,328,948 - 19,908,951

Common Wall Lizard Chromosome 11

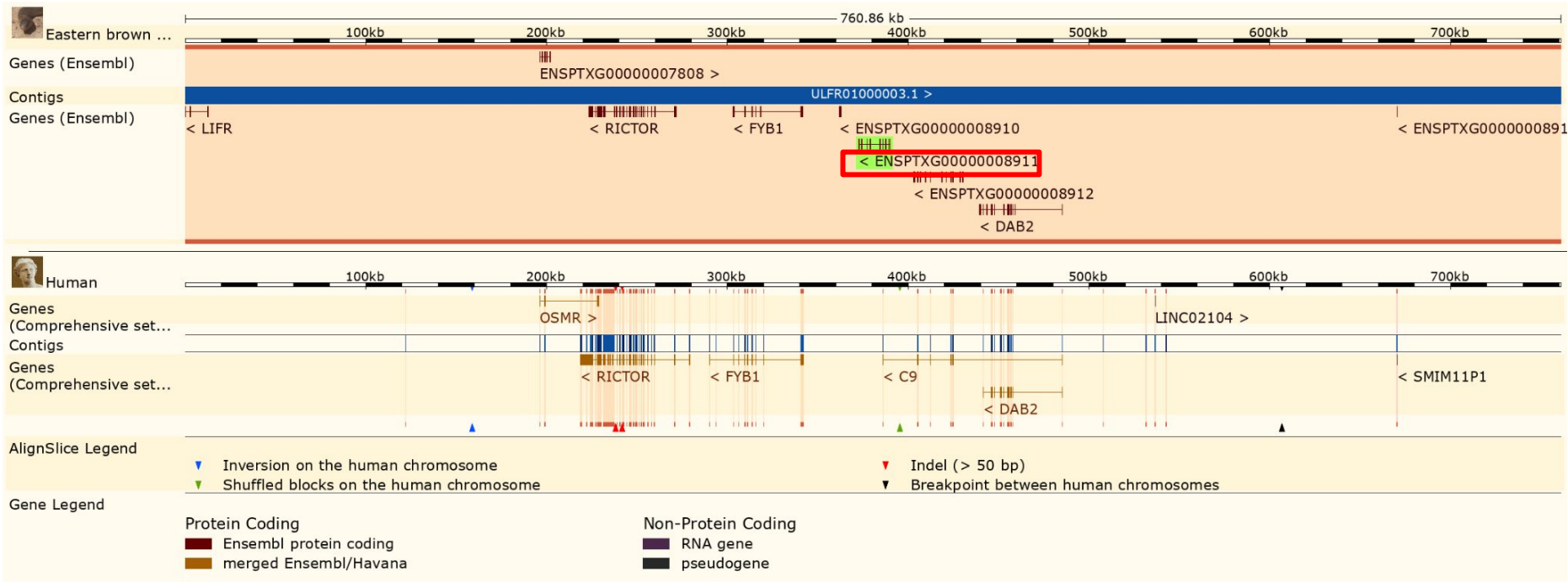
Regions of C9 gene

Human chromosome 5

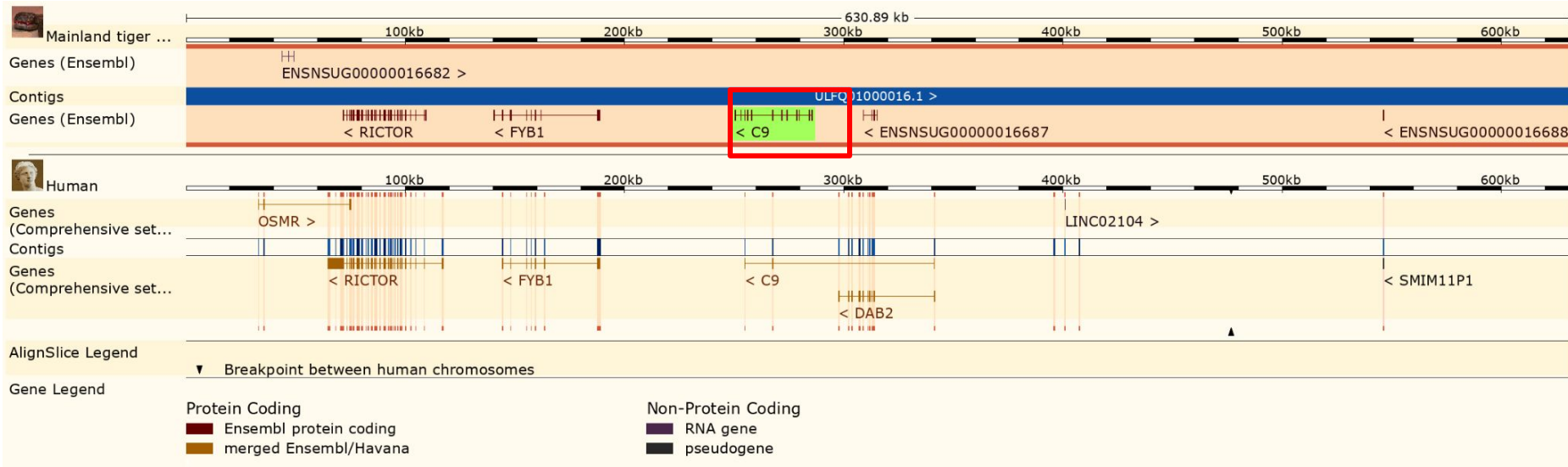


Human chromosome 9

Eastern Brown Snake

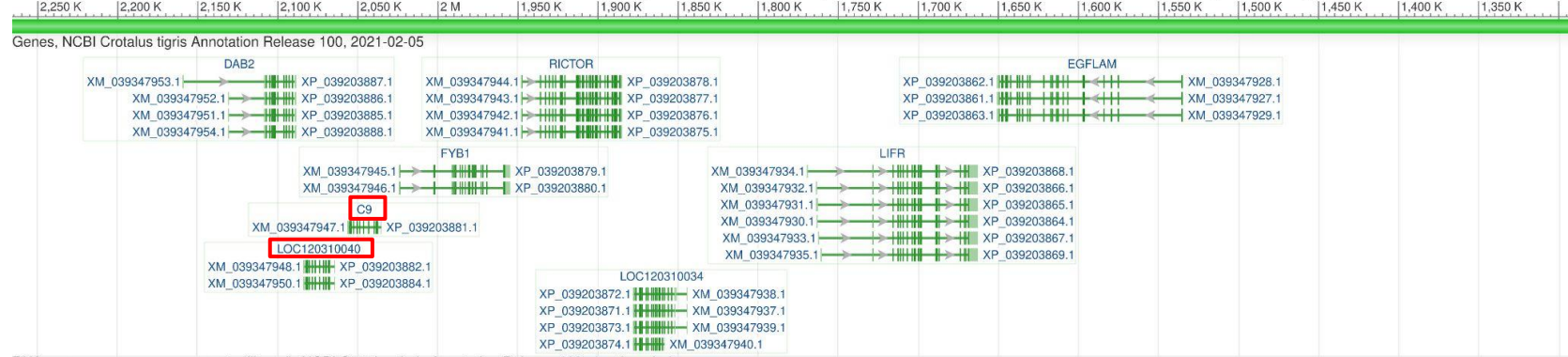


Mainland tiger snake

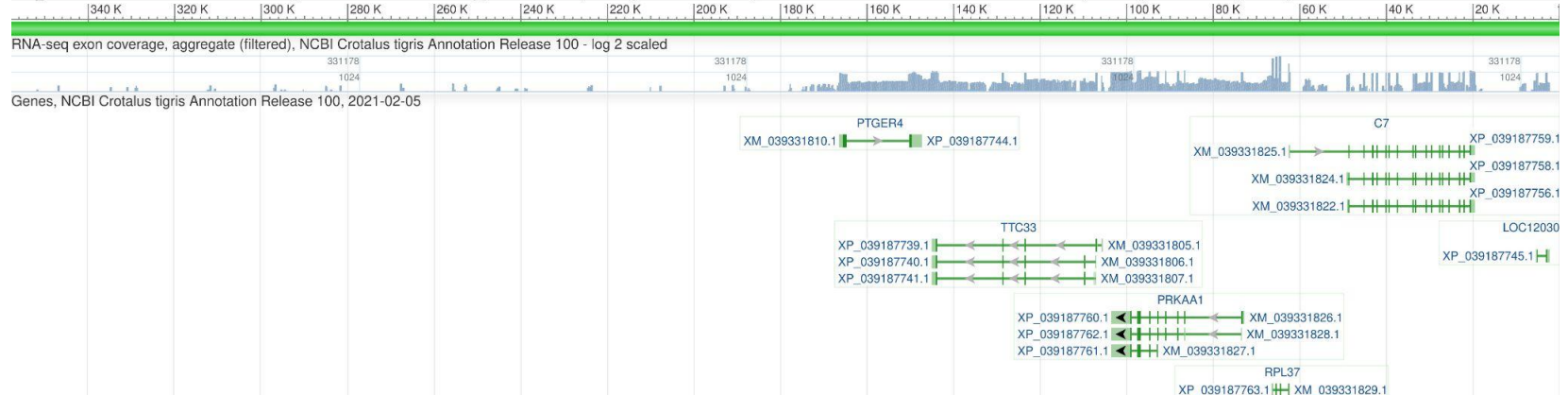


Tiger rattlesnake

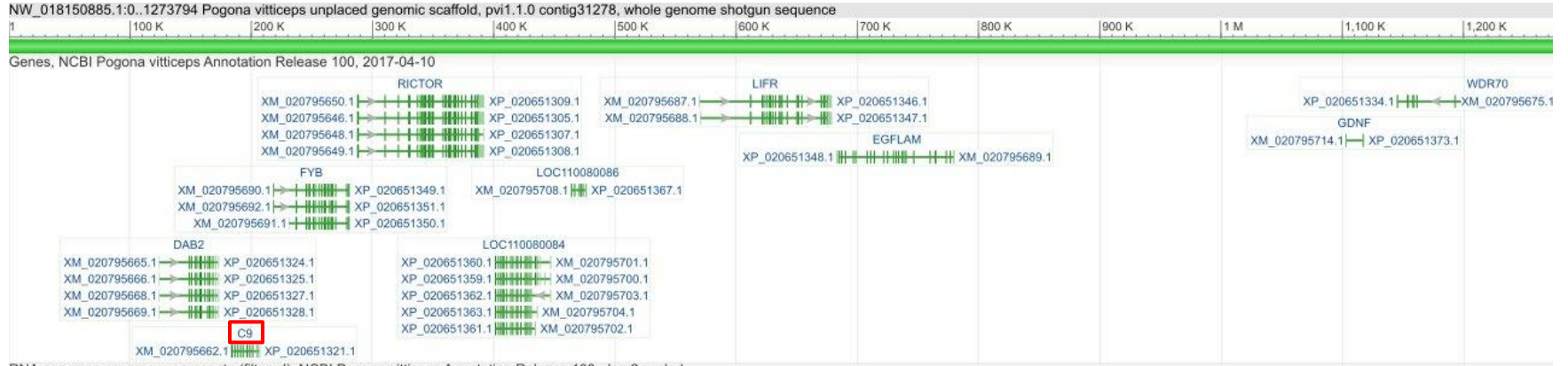
NW_024096824.1:1294201..2265977 *Crotalus tigris* isolate CLP2741 ecotype Arizona unplaced genomic scaffold, ASM1654583v1 scf7180000018103.2265978, whole genome shotgun sequence



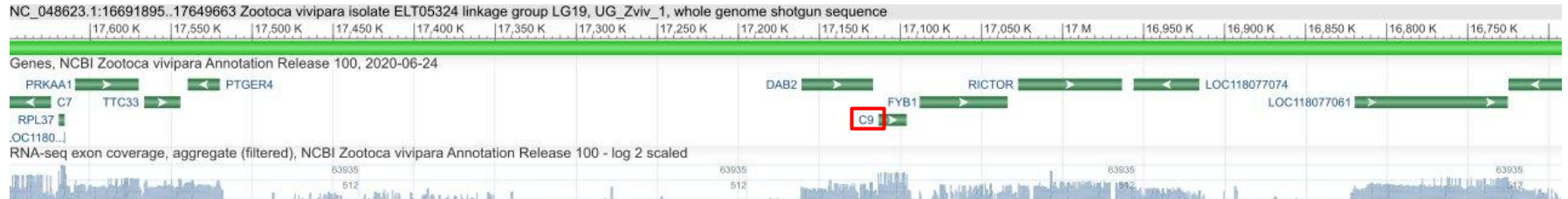
NW_024100607.1:0..357746 *Crotalus tigris* isolate CLP2741 ecotype Arizona unplaced genomic scaffold, ASM1654583v1 scf7180000024438.1716836, whole genome shotgun sequence



Central bearded dragon

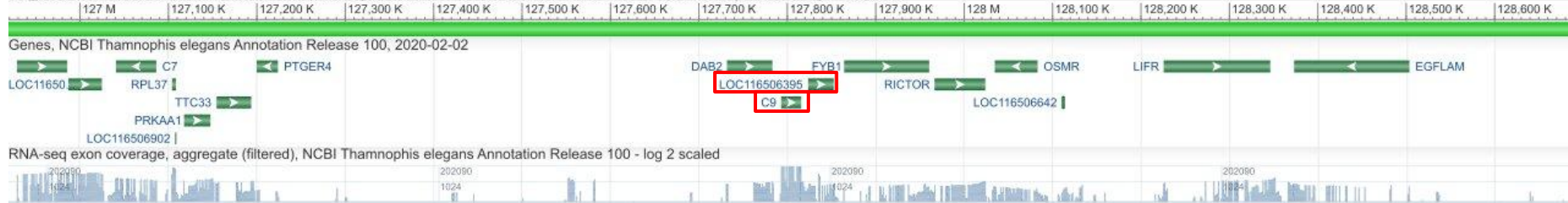


Common lizard



Western terrestrial garter snake

NC_045543.1:126918989..128689645 *Thamnophis elegans* isolate rThaEle1 chromosome 3, rThaEle1.pri, whole genome shotgun sequence



Sand lizard

NC_046322.1:17262049..18200051 *Lacerta agilis* isolate rLacAgi1 chromosome 11, rLacAgi1.pri, whole genome shotgun sequence

