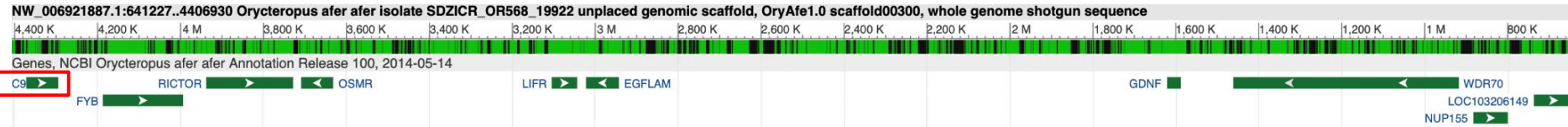
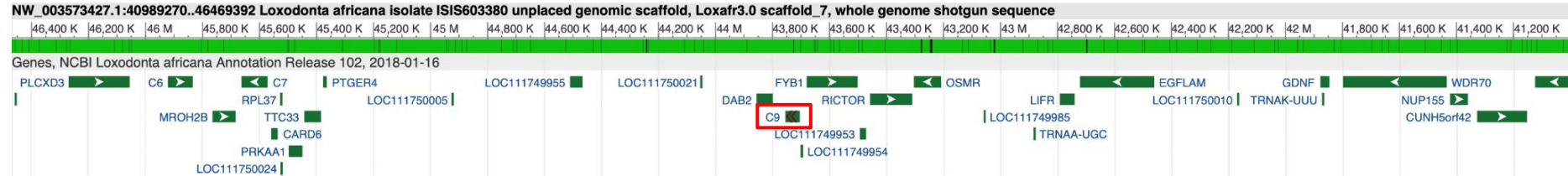


C9 Synteny in Afrotheria

Aardvark



African savanna elephant

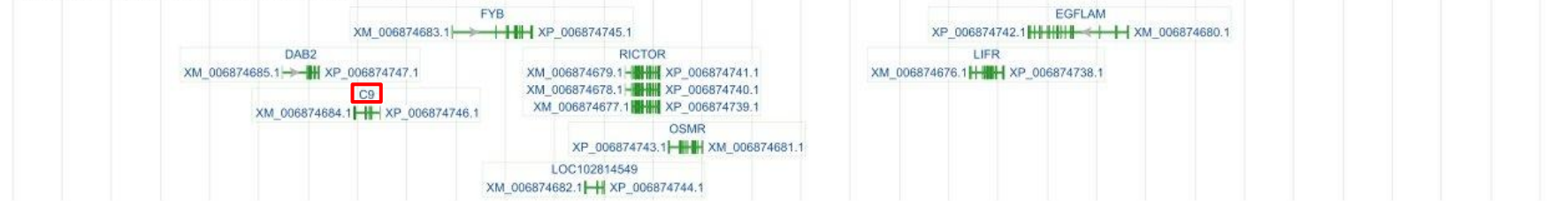


Cape golden mole

NW_006408811.1:265251..3505276 *Chrysochloris asiatica* unplaced genomic scaffold, ChrAsi1.0 scaffold00293, whole genome shotgun sequence

3,505,276 | 3,300 K | 3,200 K | 3,100 K | 3 M | 2,900 K | 2,800 K | 2,700 K | 2,600 K | 2,500 K | 2,400 K | 2,300 K | 2,200 K | 2,100 K | 2 M | 1,900 K | 1,800 K | 1,700 K | 1,600 K | 1,500 K | 1,400 K | 1,300 K | 1,200 K | 1,100 K | 1 M | 900 K | 800 K | 700 K | 600 K | 500 K | 400 K

Genes, NCBI *Chrysochloris asiatica* Annotation Release 100, 2014-02-19



Florida manatee

NW_004444133.1:1949714..3651968 *Trichechus manatus latirostris* isolate Lorelei unplaced genomic scaffold, TriManLat1.0 scaffold00197, whole genome shotgun sequence

3,600 K | 3,500 K | 3,400 K | 3,300 K | 3,200 K | 3,100 K | 3 M | 2,900 K | 2,800 K | 2,700 K | 2,600 K | 2,500 K | 2,400 K | 2,300 K | 2,200 K | 2,100 K | 2 M

Genes, NCBI *Trichechus manatus latirostris* Annotation Release 102, 2018-01-25



Cape elephant shrew

NW_006399829.1:3144477..4317201 Elephantulus edwardii isolate Elephantulus edwardii_28Jun2011 unplaced genomic scaffold, EleEdw1.0 scaffold00079, whole genome shotgun sequence



small Madagascar hedgehog

NW_022111373.1:111063724..112748637 Echinops telfairi isolate 266 unplaced genomic scaffold, ASM31398v2 ScPeMD7_7469:HRSCAF=8507_1, whole genome shotgun sequence

