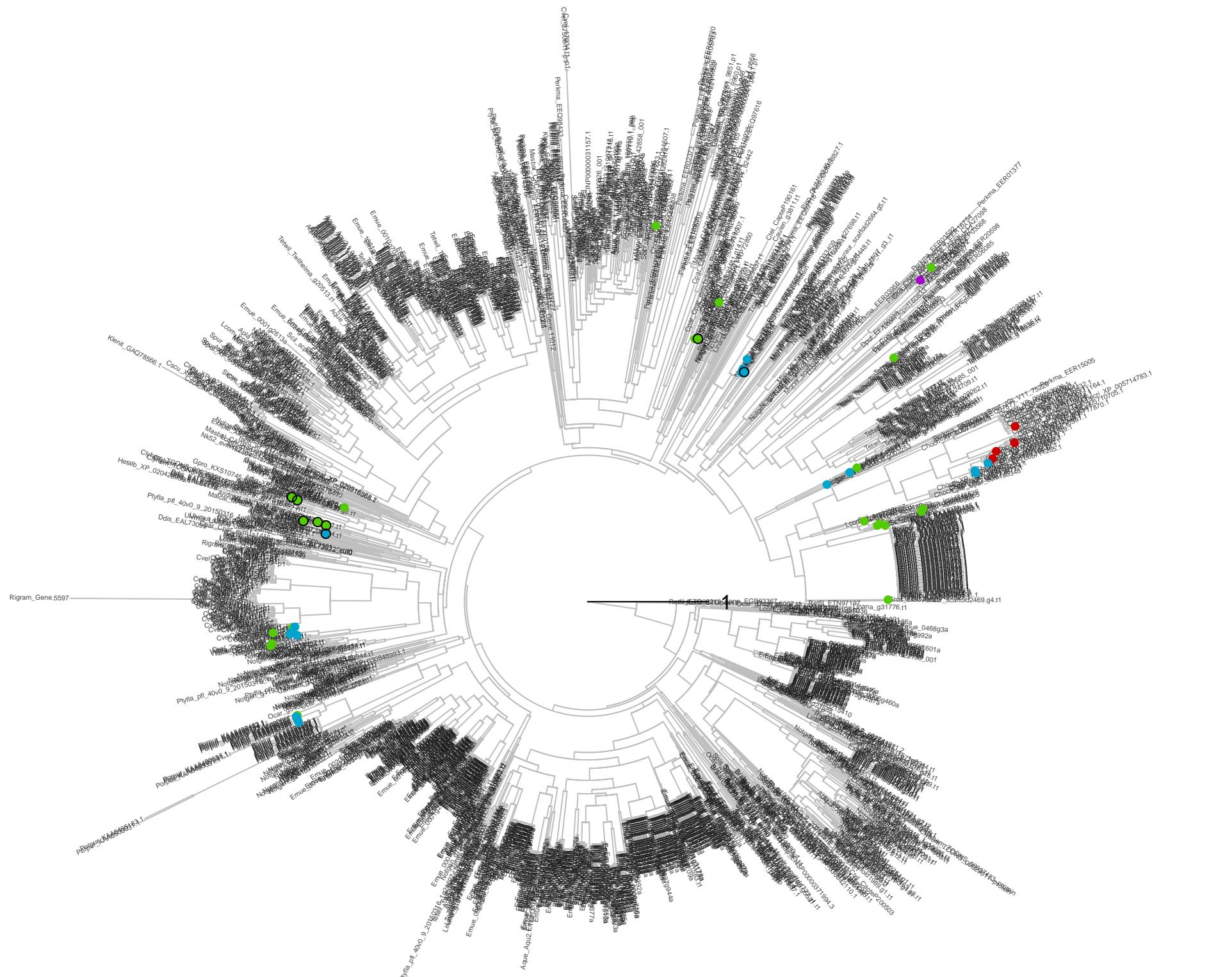
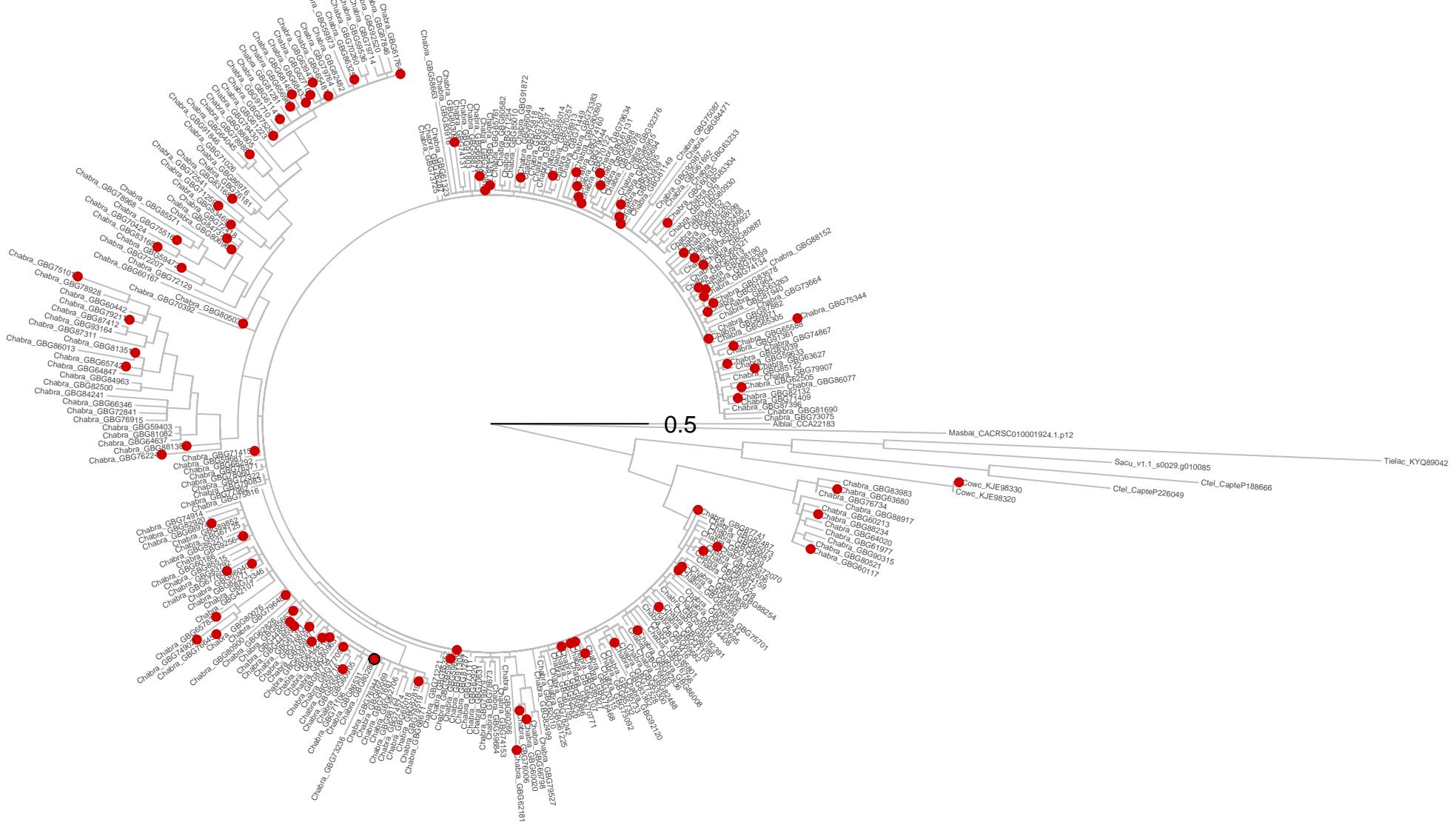


## **Integrase\_H2C2.HG1**



1906 genes  
n=51 fusion(s) with Chromo domains  
(8 pass filters)

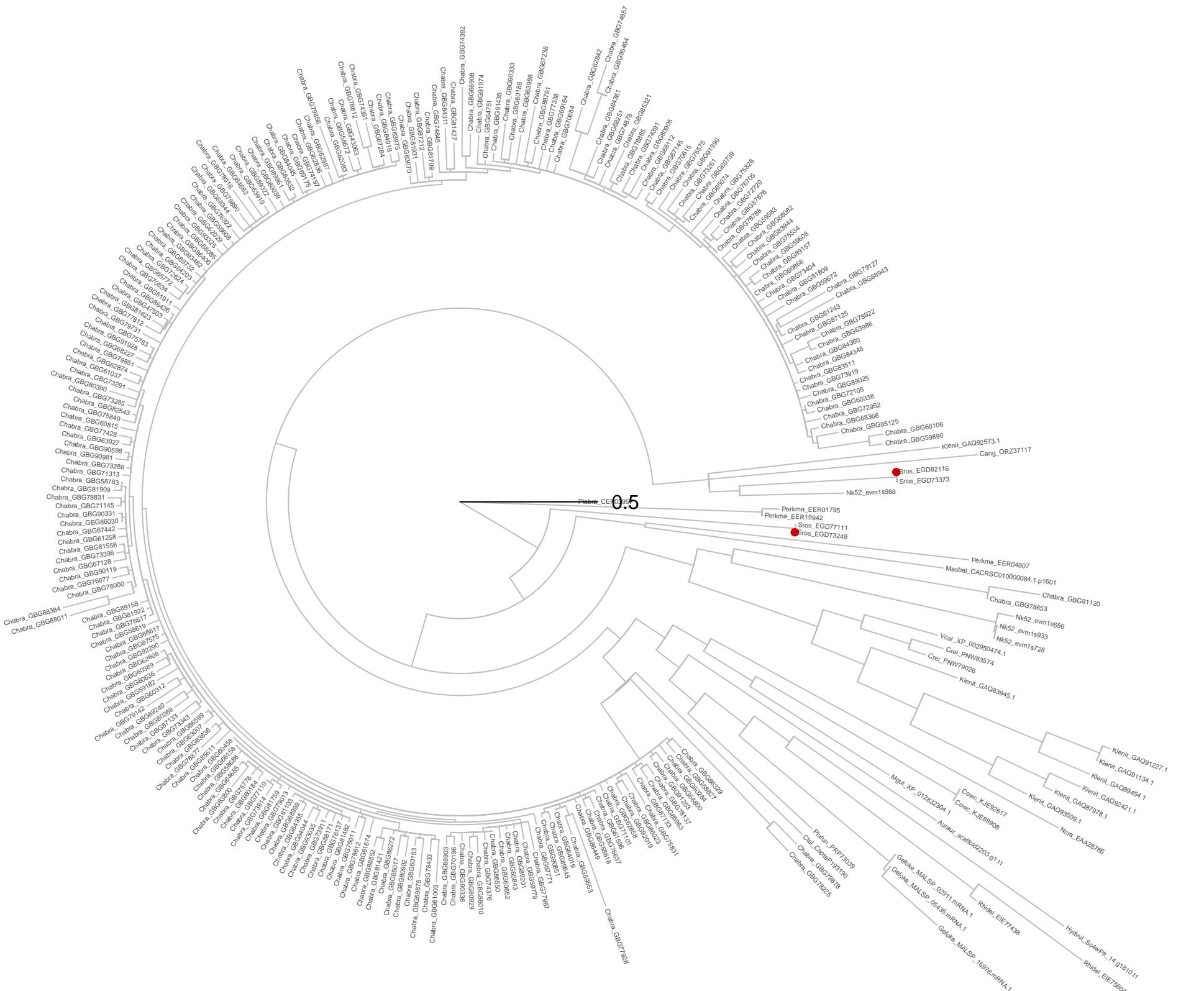
# Integrase\_H2C2.HG10



● Chromo.HG2.1:NA n = 98

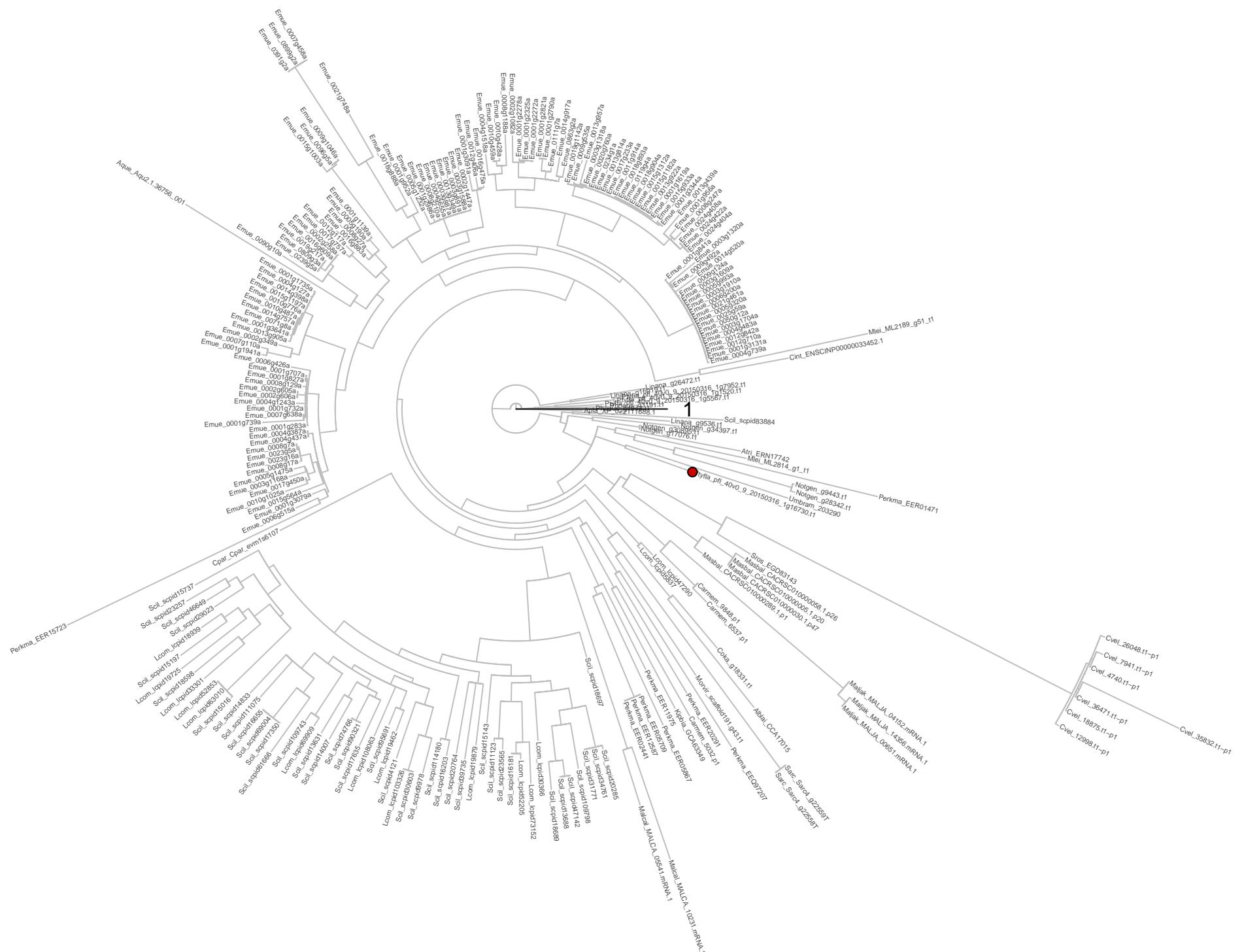
303 genes  
n=98 fusion(s) with Chromo domains  
(1 pass filters)

# Integrase\_H2C2.HG11



● Chromo.HG2.1:NA n = 2

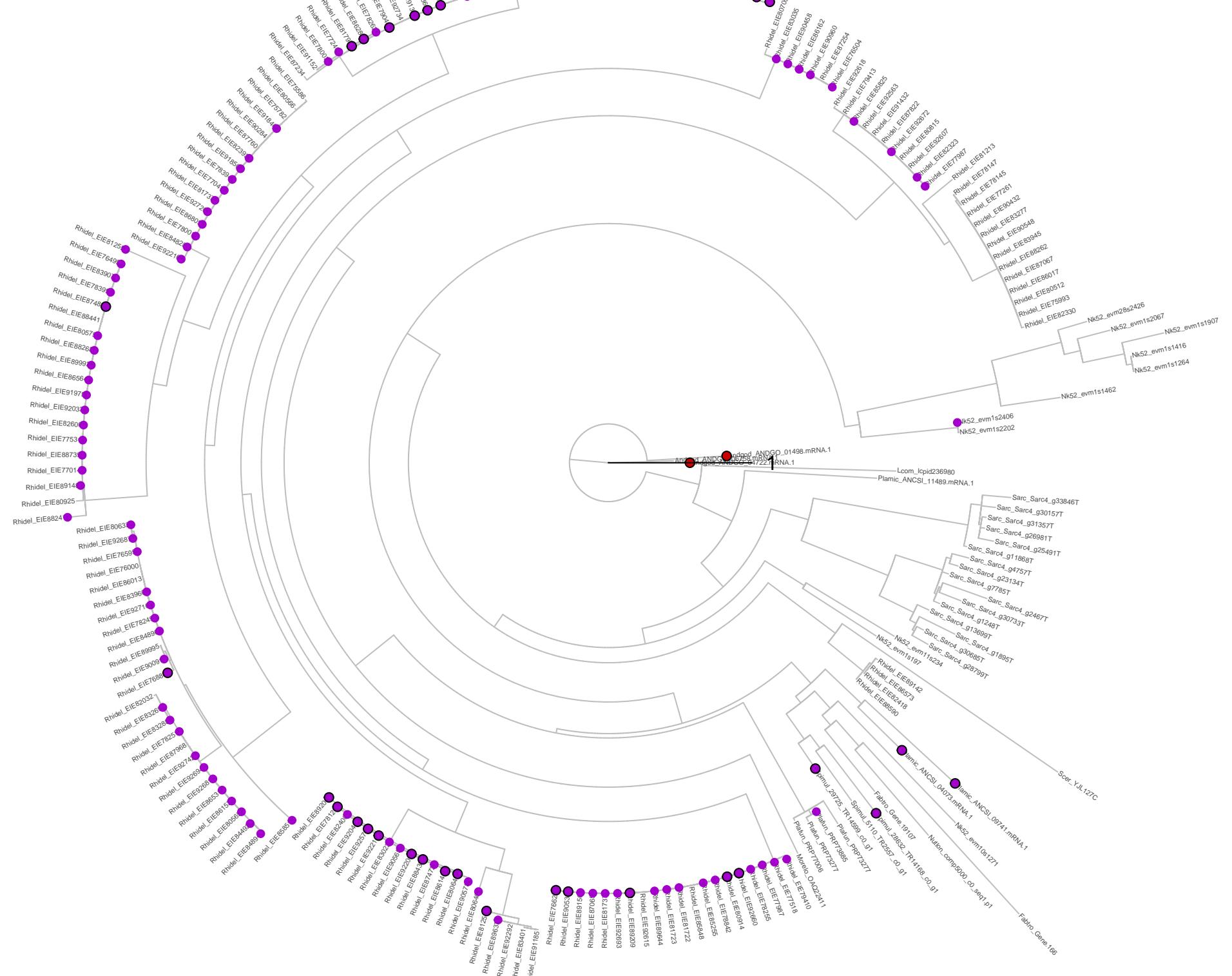
# Integrase\_H2C2.HG13



238 genes  
n=1 fusion(s) with Chromo domains  
(1 pass filters)

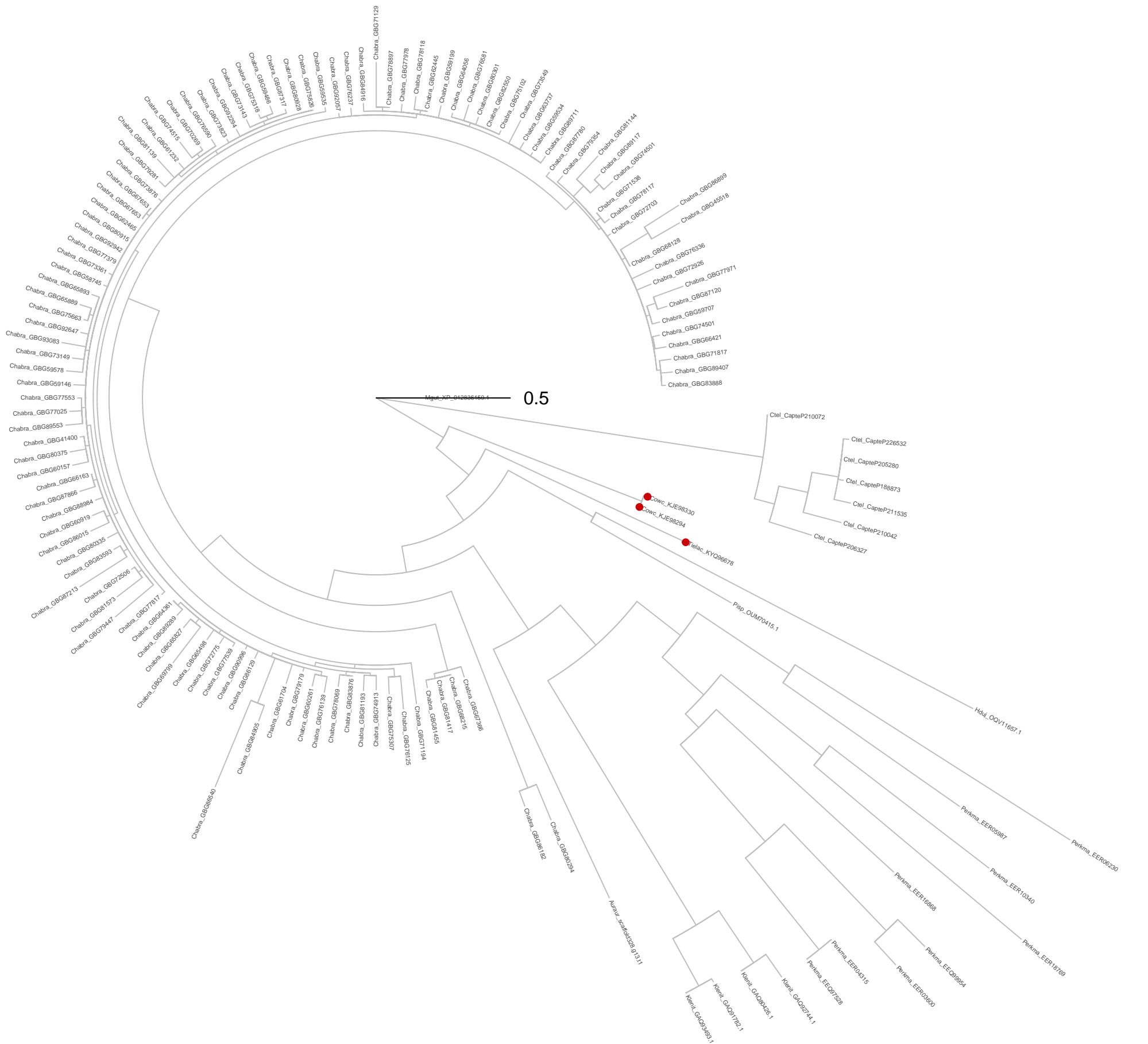
● Chromo.HG3.2:NA n = 1

## **Integrase\_H2C2.HG14**



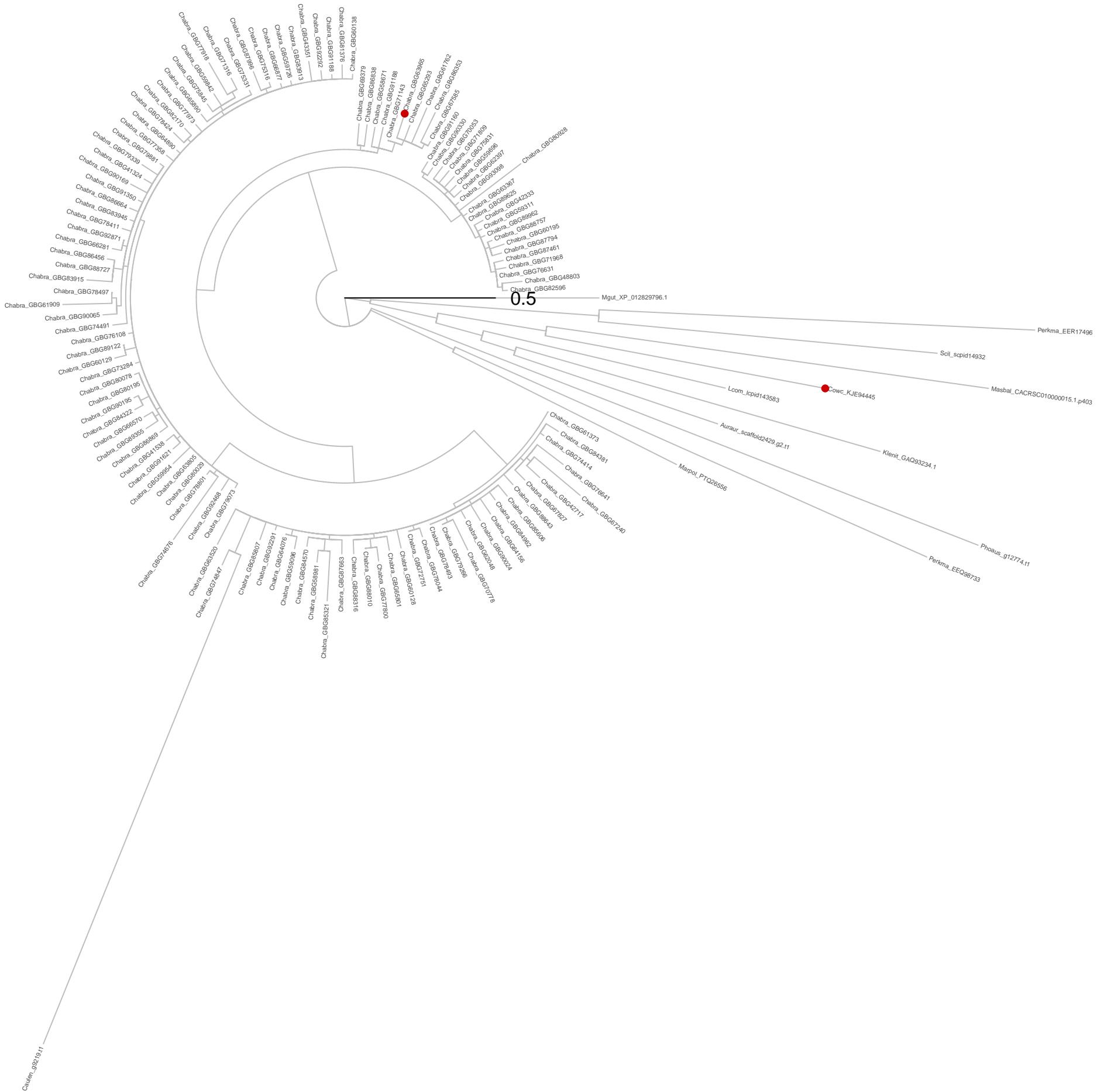
219 genes  
n=128 fusion(s) with Chromo domains  
(40 pass filters)

## **Integrase\_H2C2.HG16**



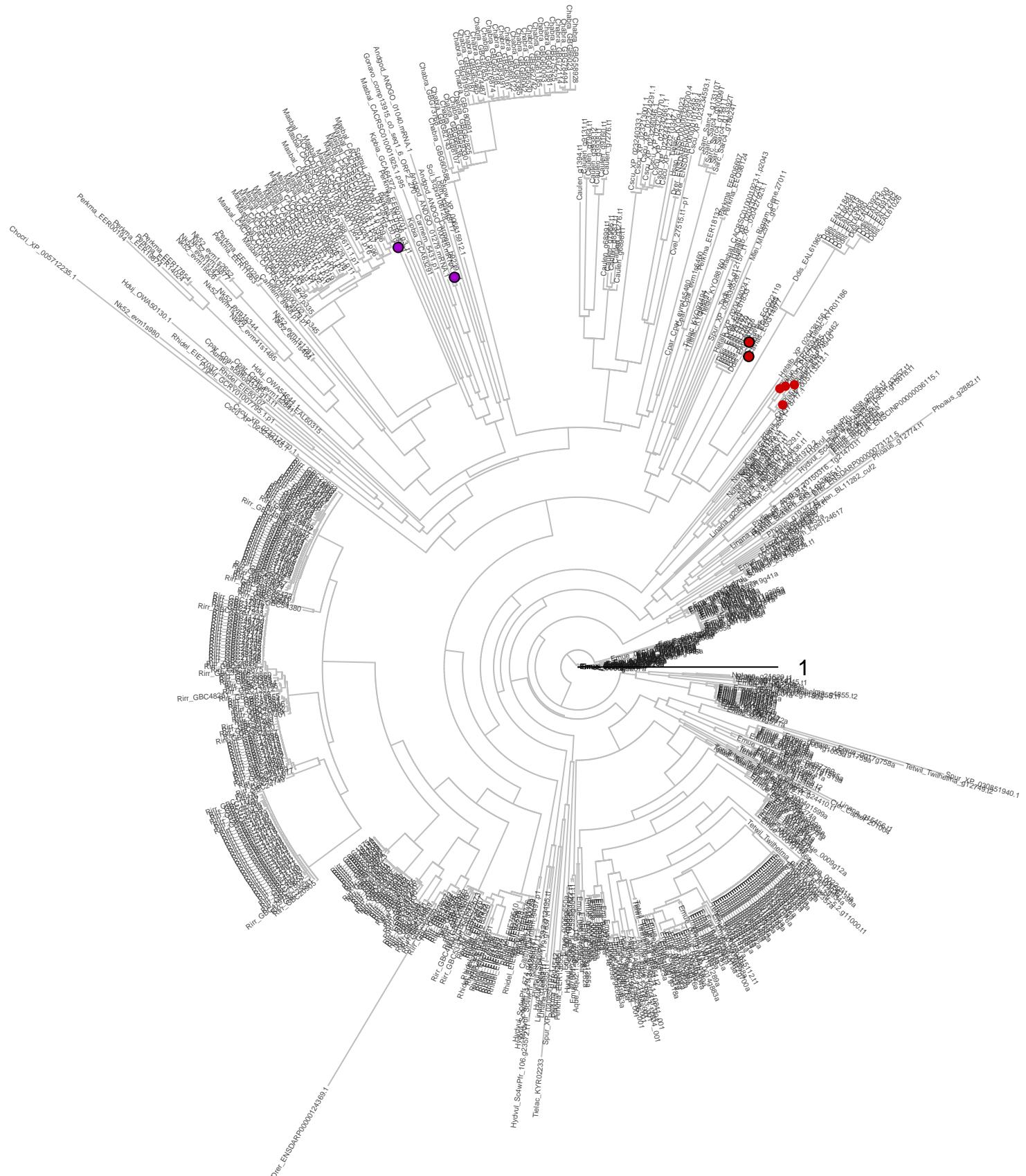
144 genes  
n=3 fusion(s) with Chromo domains  
(0 pass filters)

# Integrase\_H2C2.HG18



135 genes  
n=2 fusion(s) with Chromo domains  
(0 pass filters)

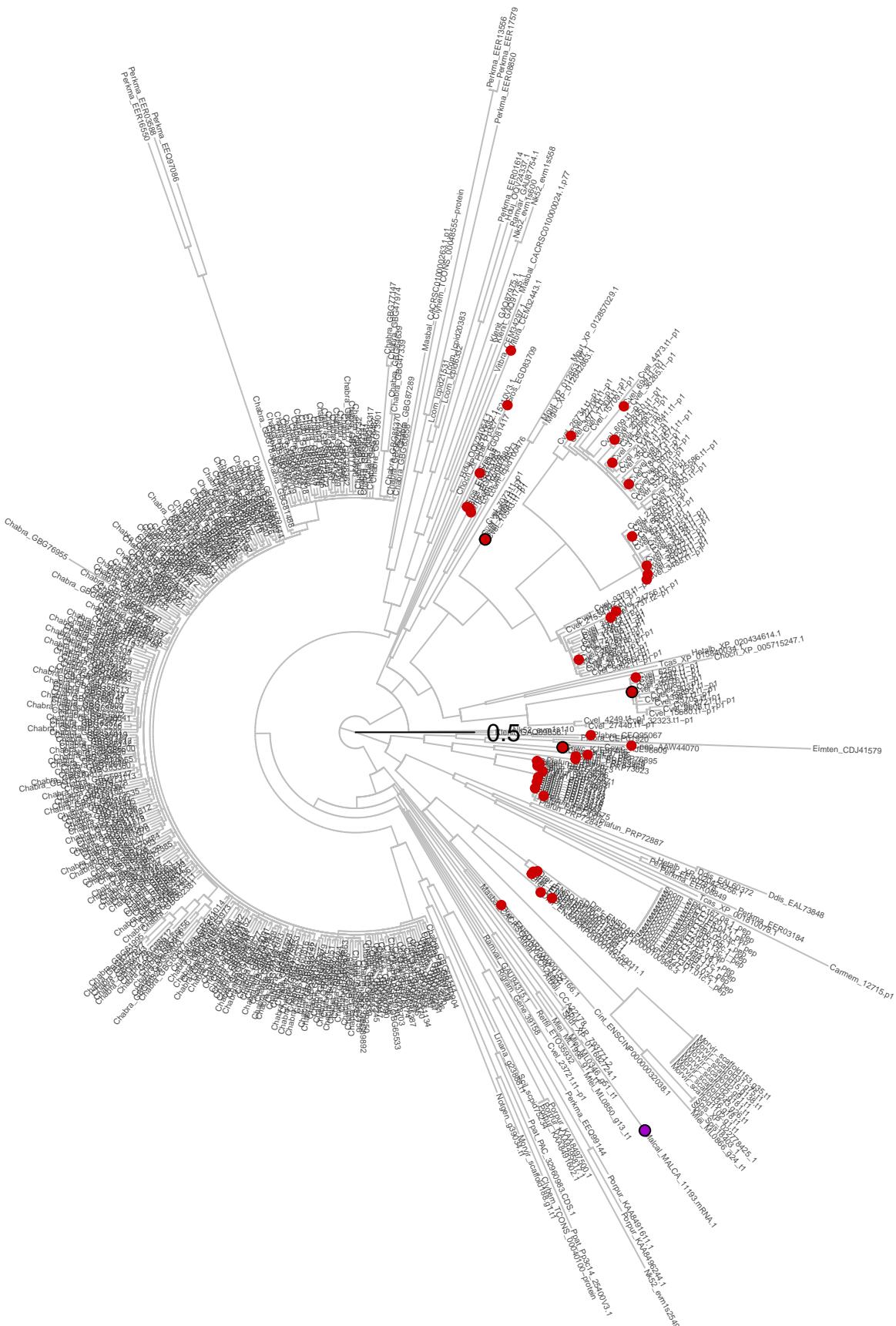
# Integrase\_H2C2.HG5



- Chromo.HG2.1:NA n = 6
- Chromo.HG3.3:NA n = 2

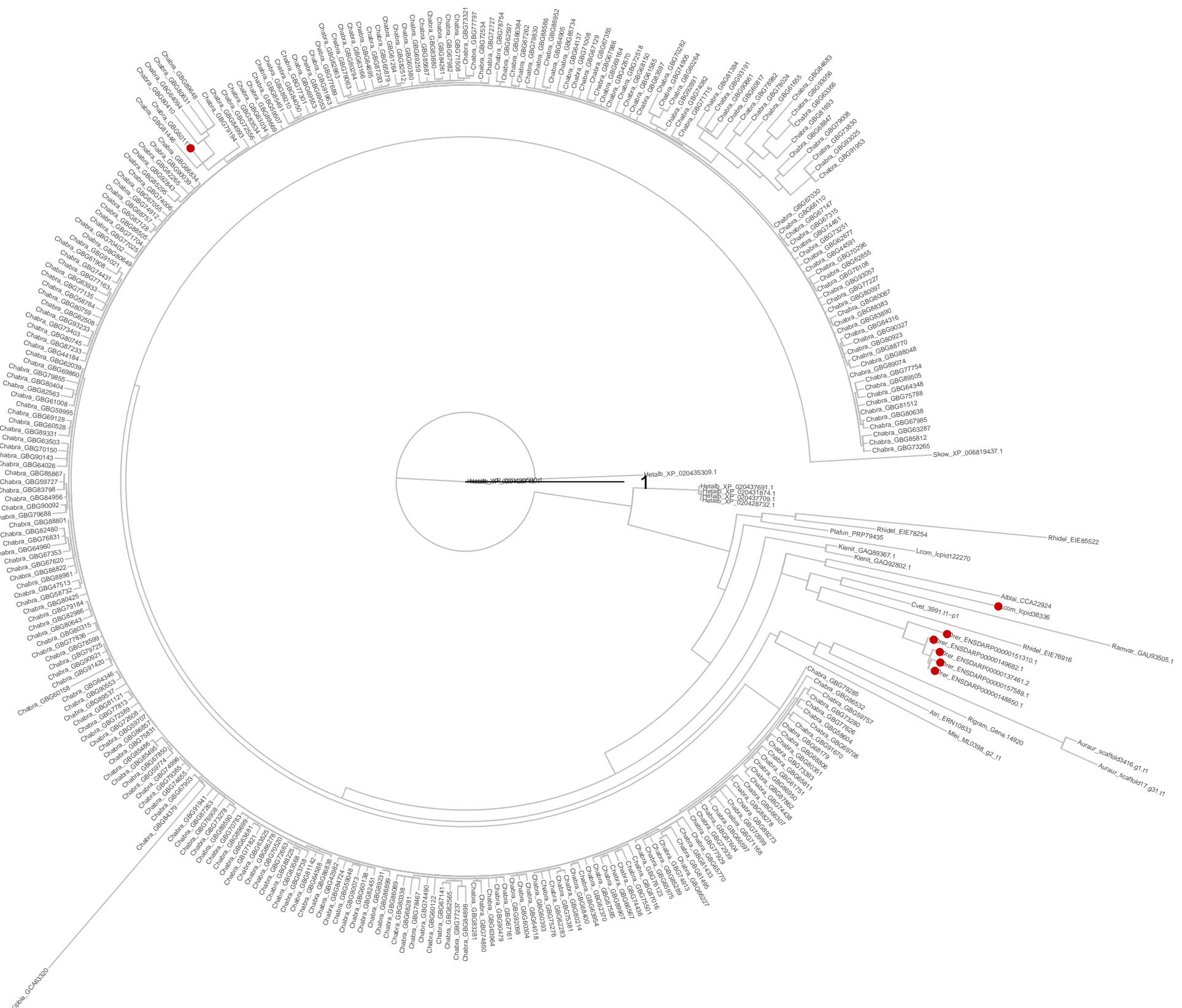
706 genes  
n=8 fusion(s) with Chromo domains  
(4 pass filters)

Integrase\_H2C2.HG7



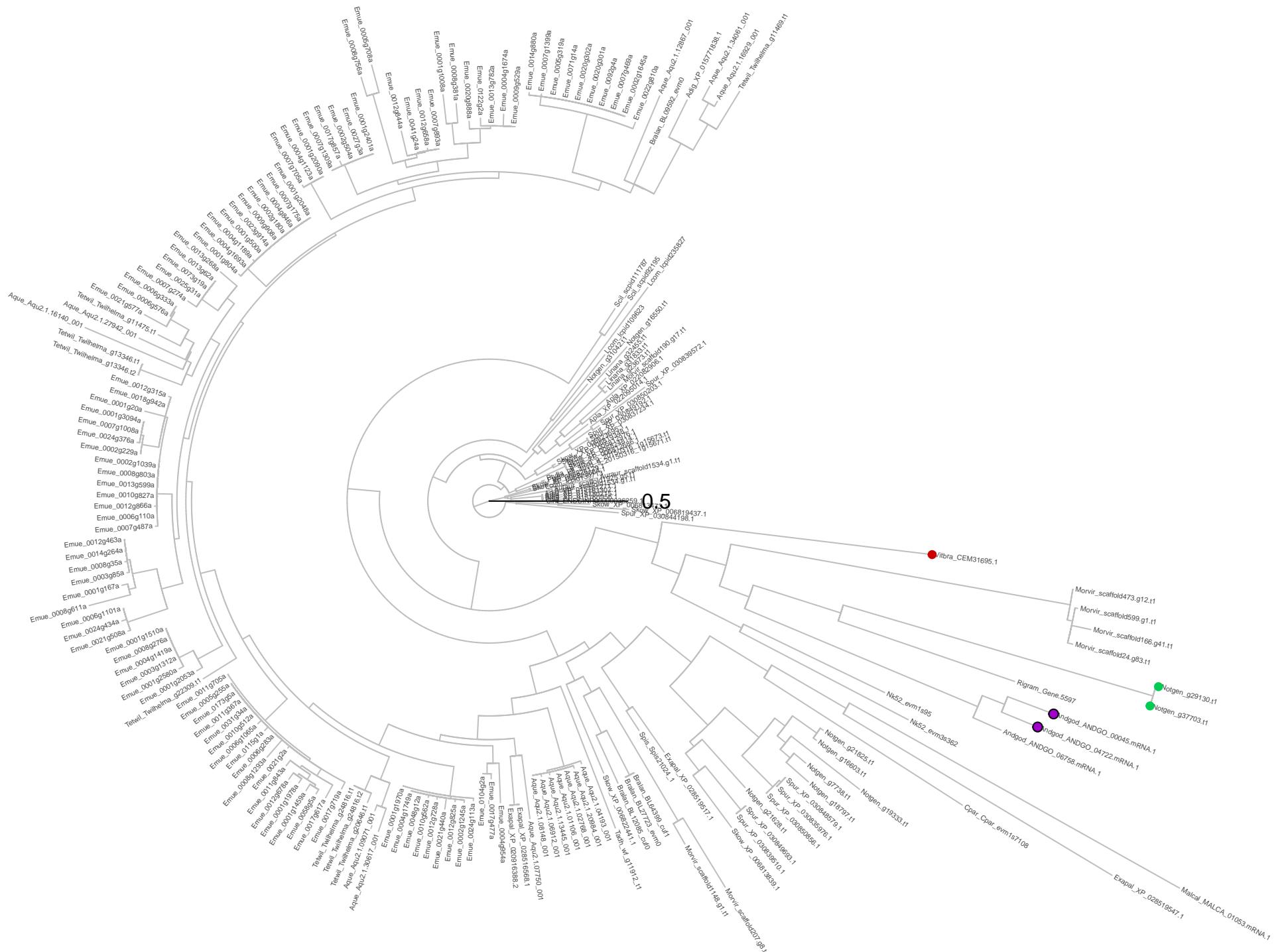
520 genes  
n=42 fusion(s) with Chromo domains  
(4 pass filters)

# Integrase\_H2C2.HG9



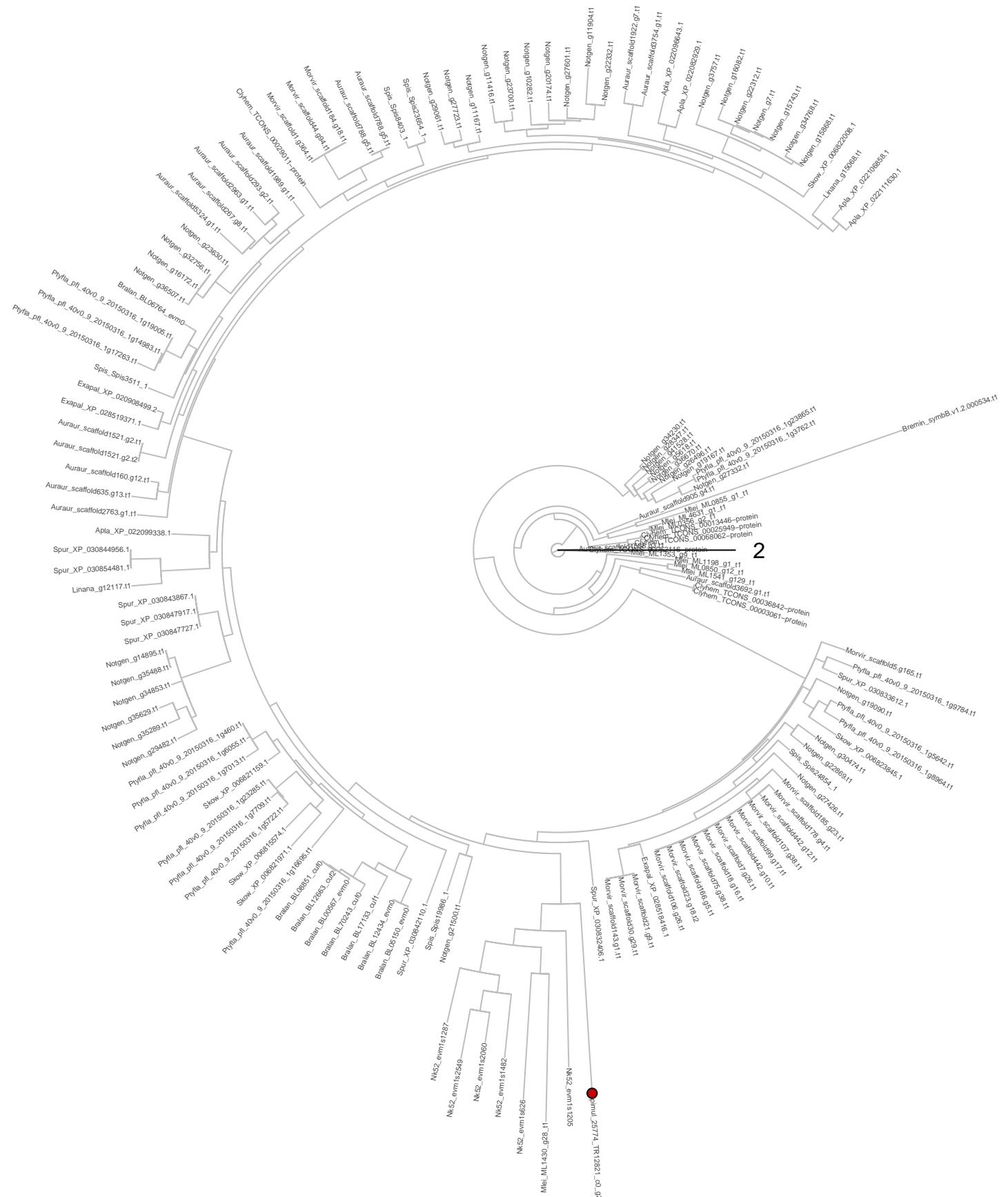
328 genes  
n=7 fusion(s) with Chromo domains  
(0 pass filters)

rve.HG10



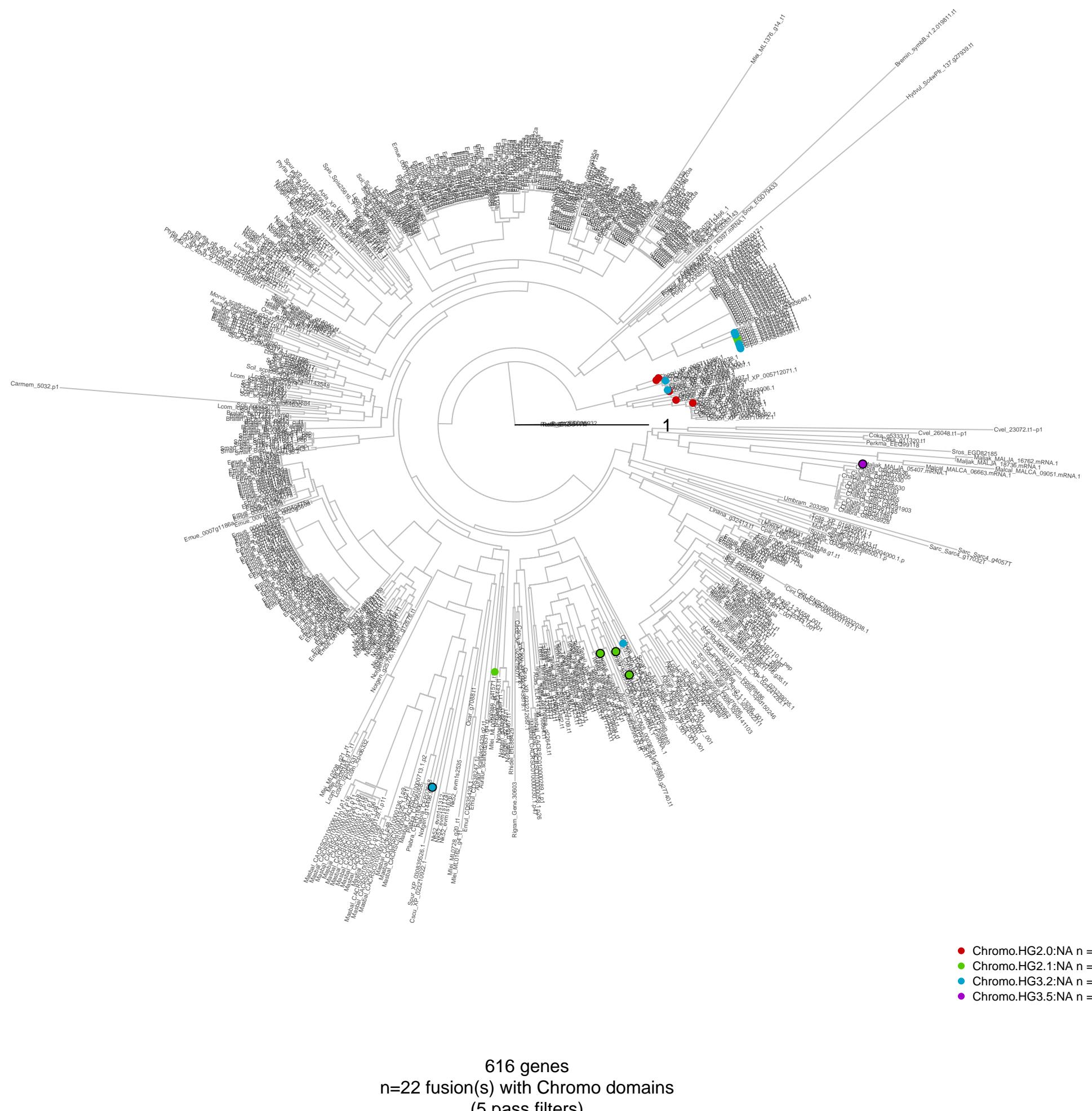
209 genes  
n=5 fusion(s) with Chromo domains  
(2 pass filters)

rve.HG14

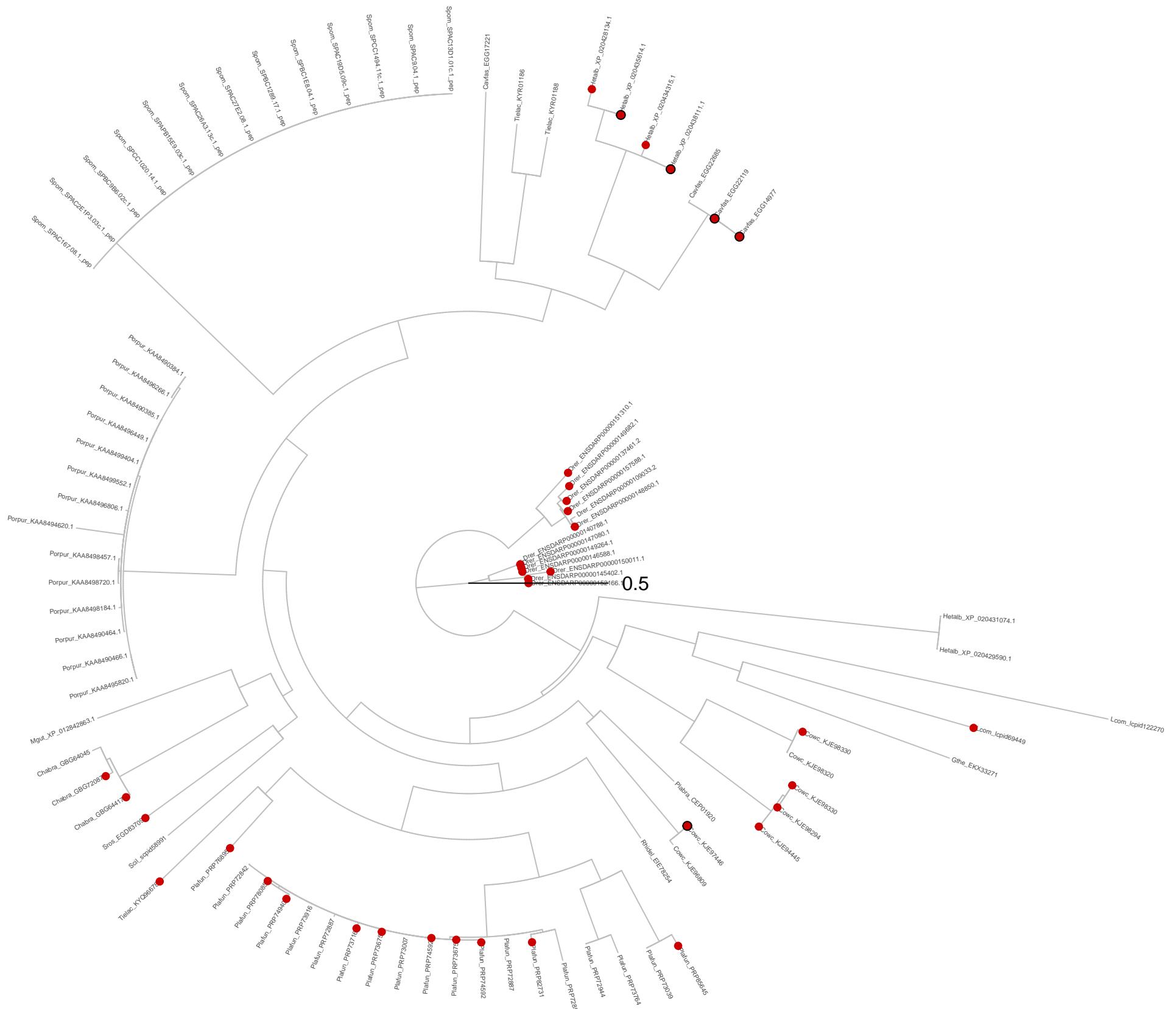


150 genes  
n=1 fusion(s) with Chromo domains  
(1 pass filters)

# rve.HG2

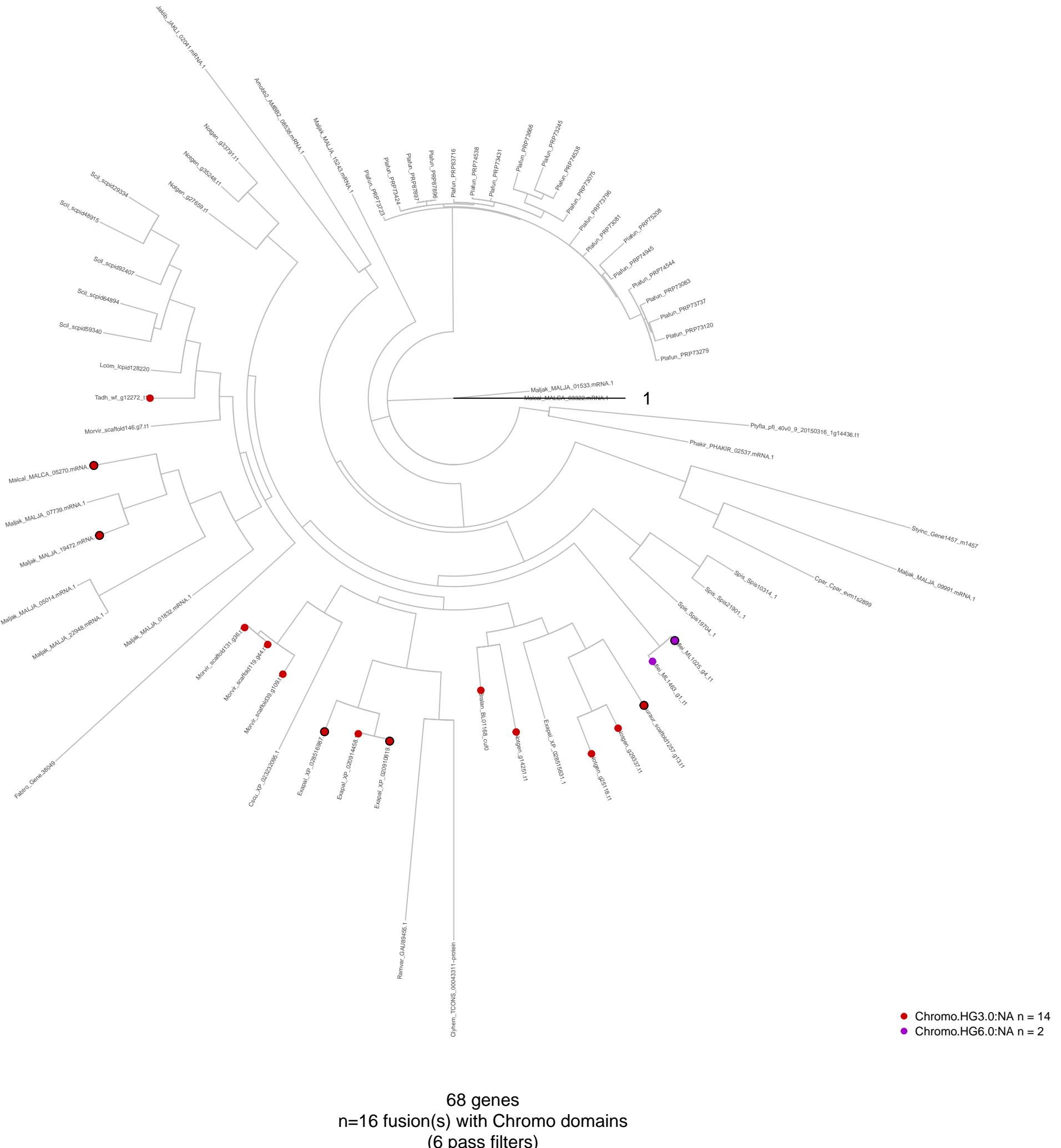


# rve.HG21

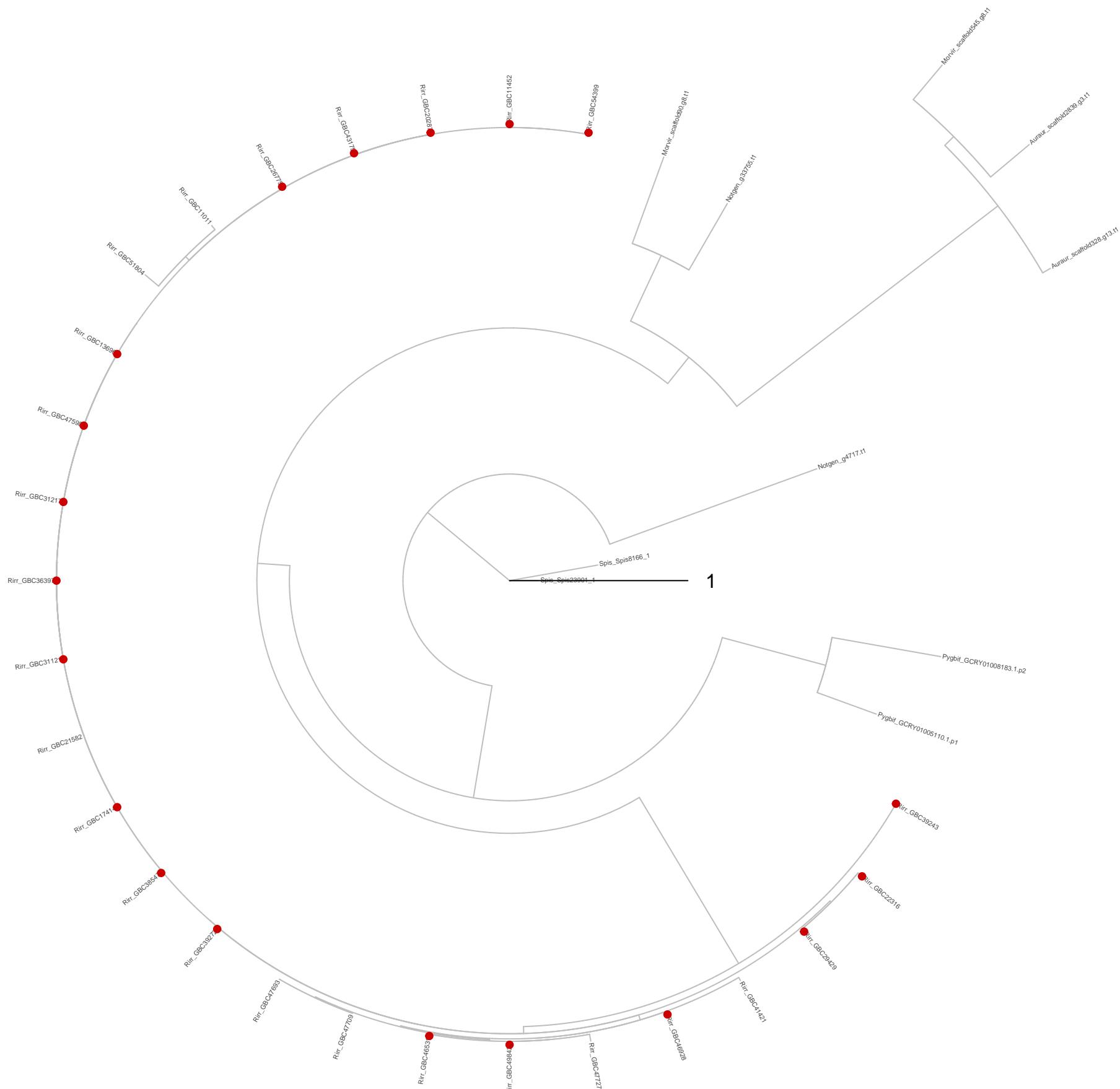


90 genes  
n=37 fusion(s) with Chromo domains  
(5 pass filters)

# rve.HG26



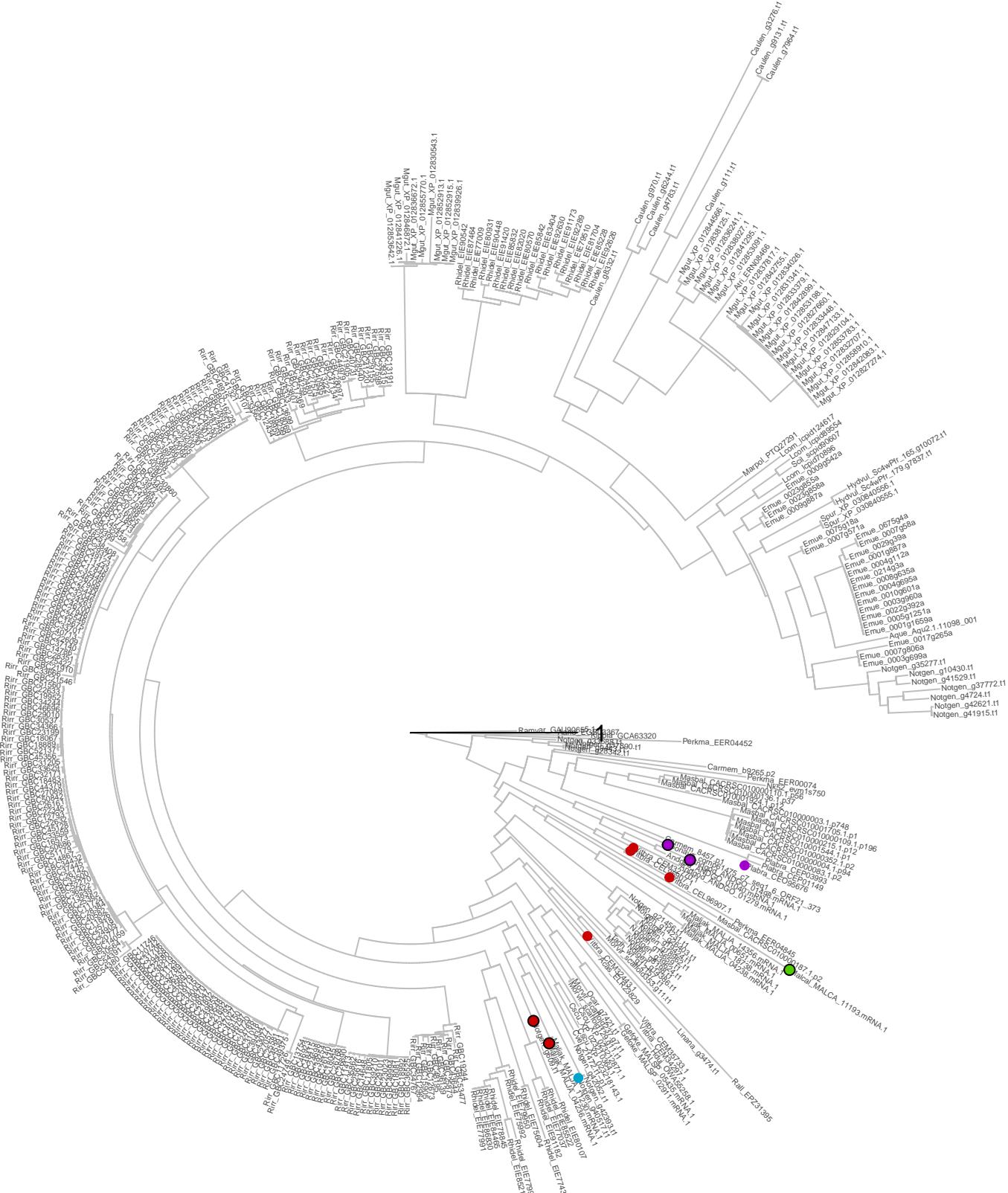
# rve.HG27



● Chromo.HG6.0:NA n = 19

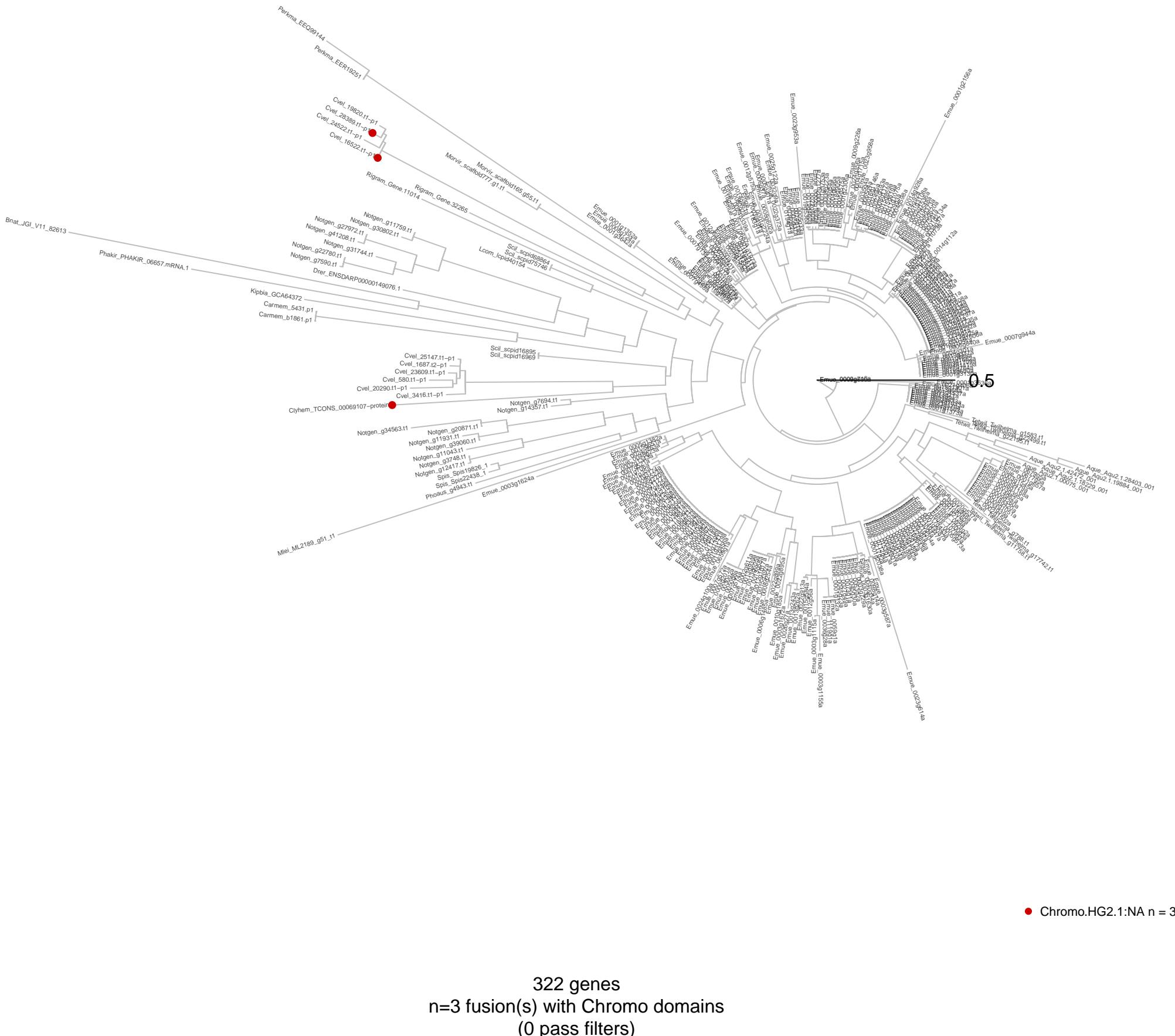
36 genes  
n=19 fusion(s) with Chromo domains  
(0 pass filters)

# rve.HG4

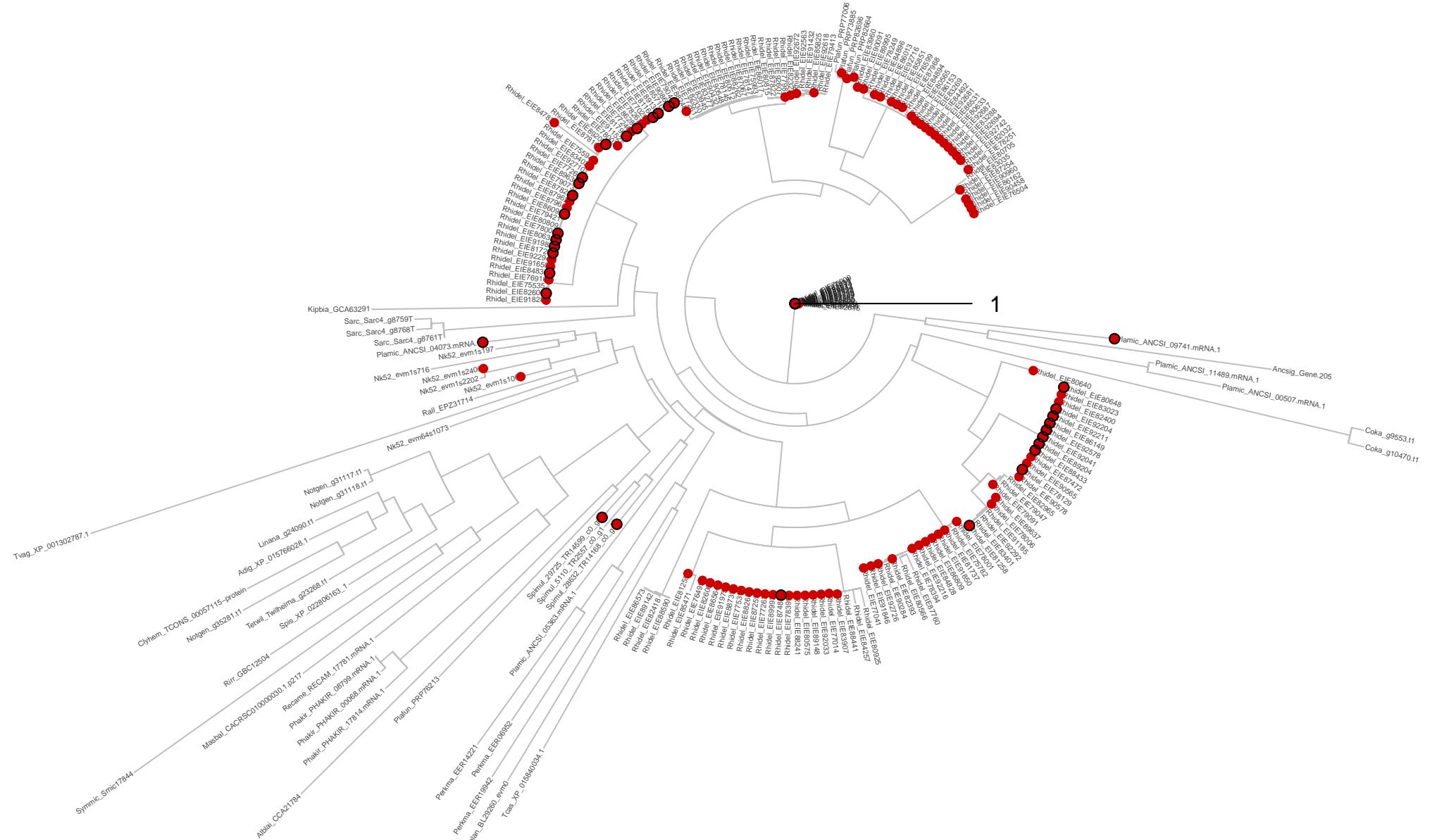


382 genes  
n=11 fusion(s) with Chromo domains  
(5 pass filters)

# rve.HG5

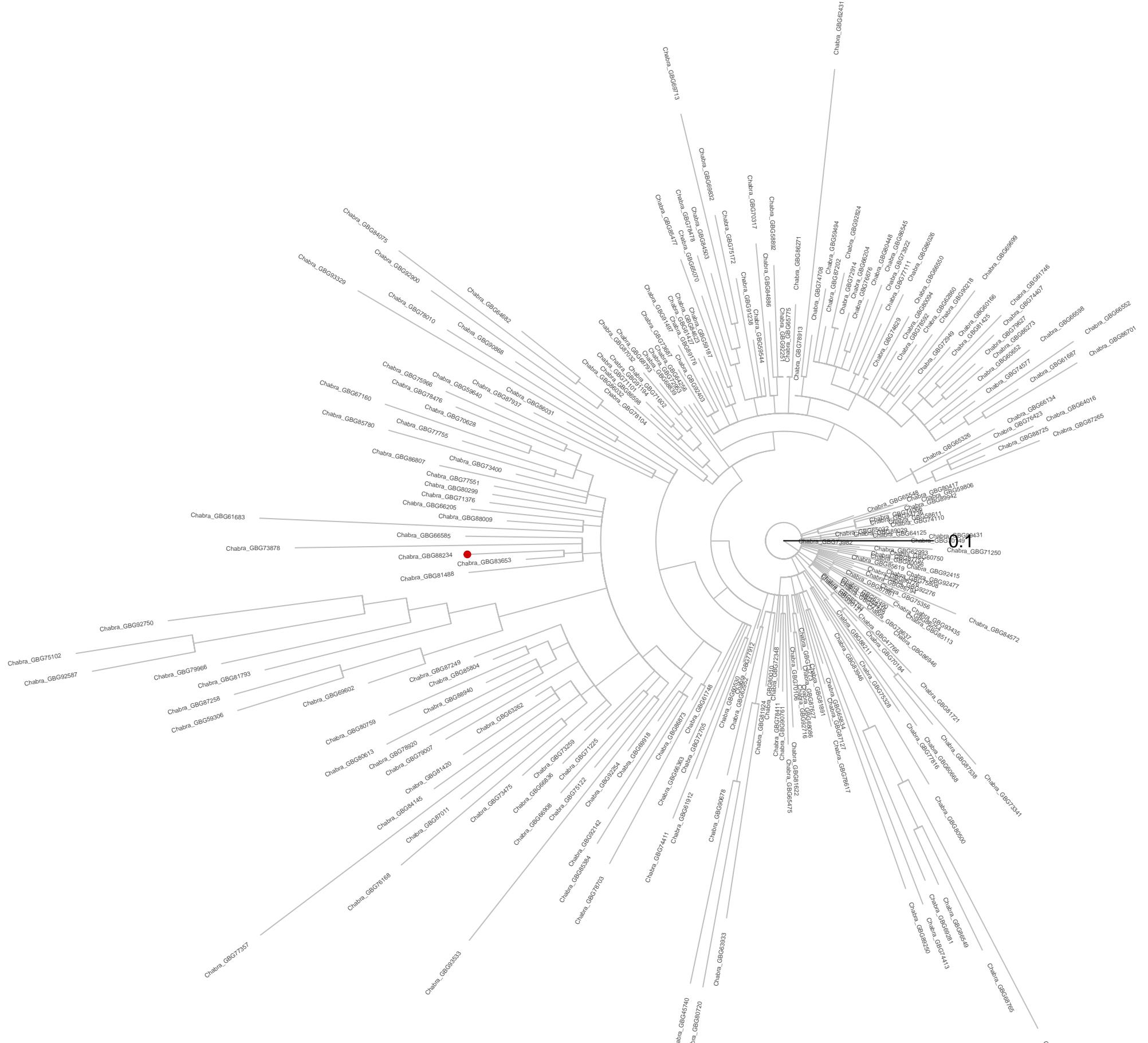


rve.HG7



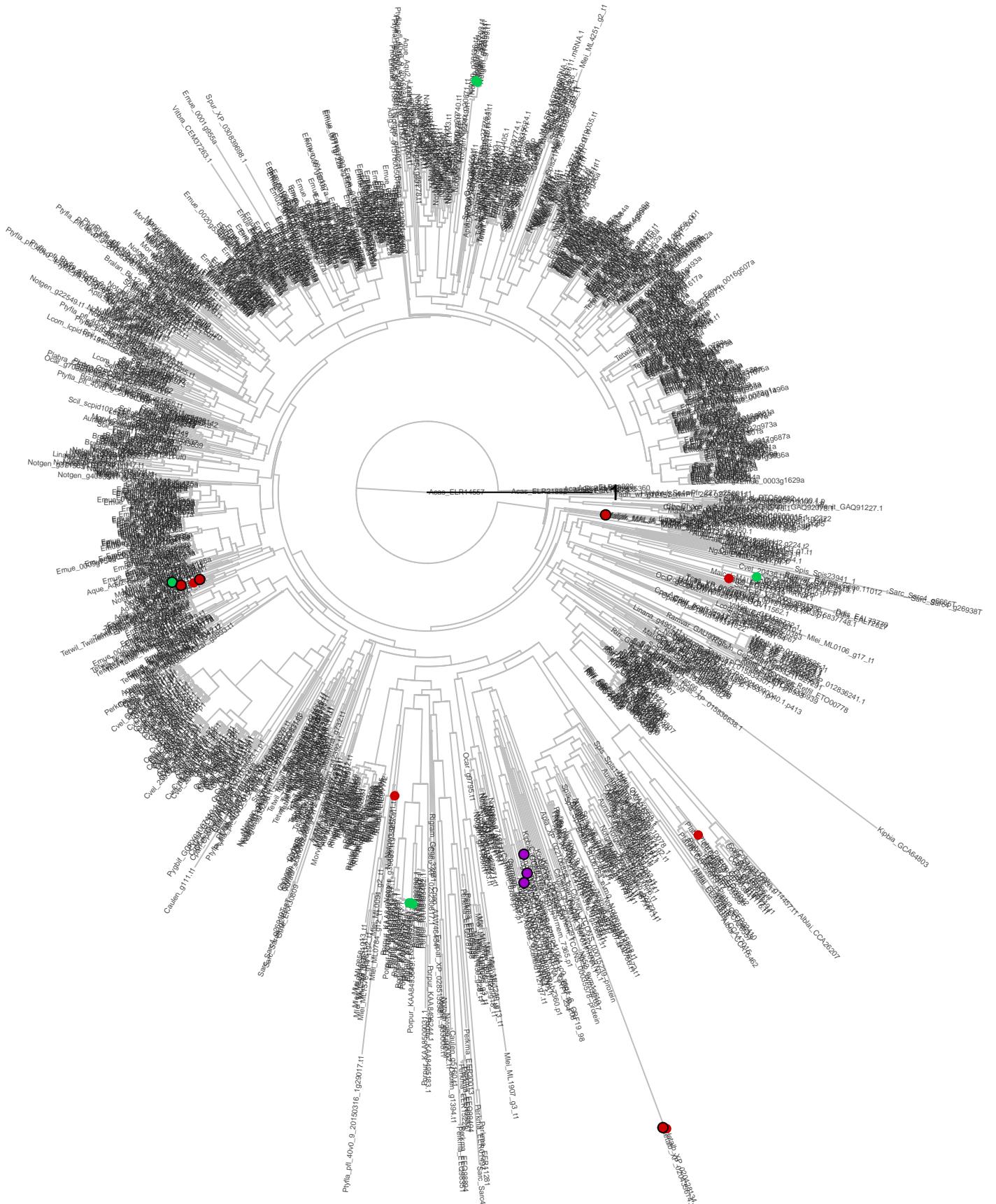
229 genes  
n=137 fusion(s) with Chromo domains  
(37 pass filters)

rve.HG8



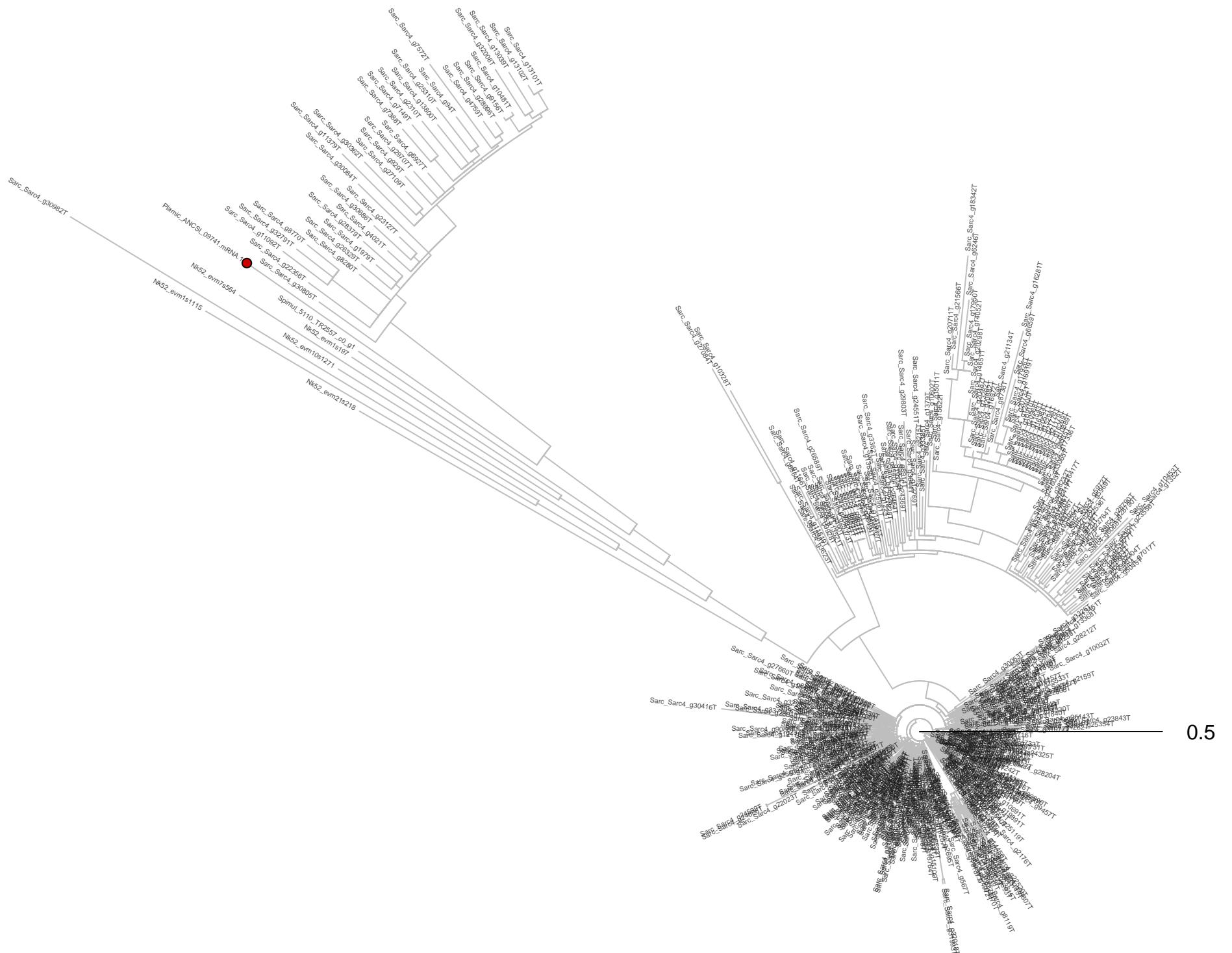
219 genes  
n=1 fusion(s) with Chromo domains  
(0 pass filters)

RVT.HG1



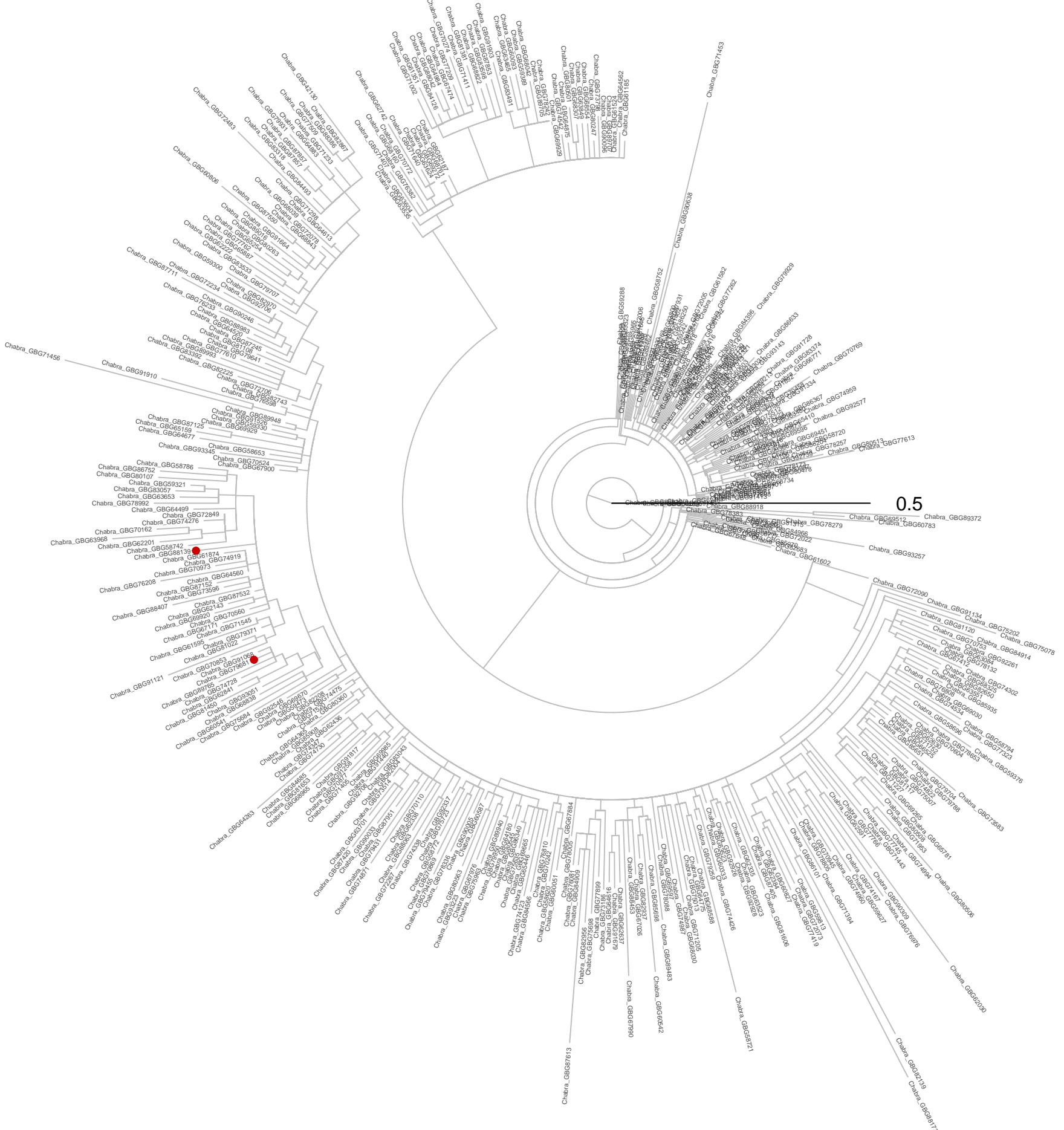
1401 genes  
n=19 fusion(s) with Chromo domains  
(8 pass filters)

RVT.HG10



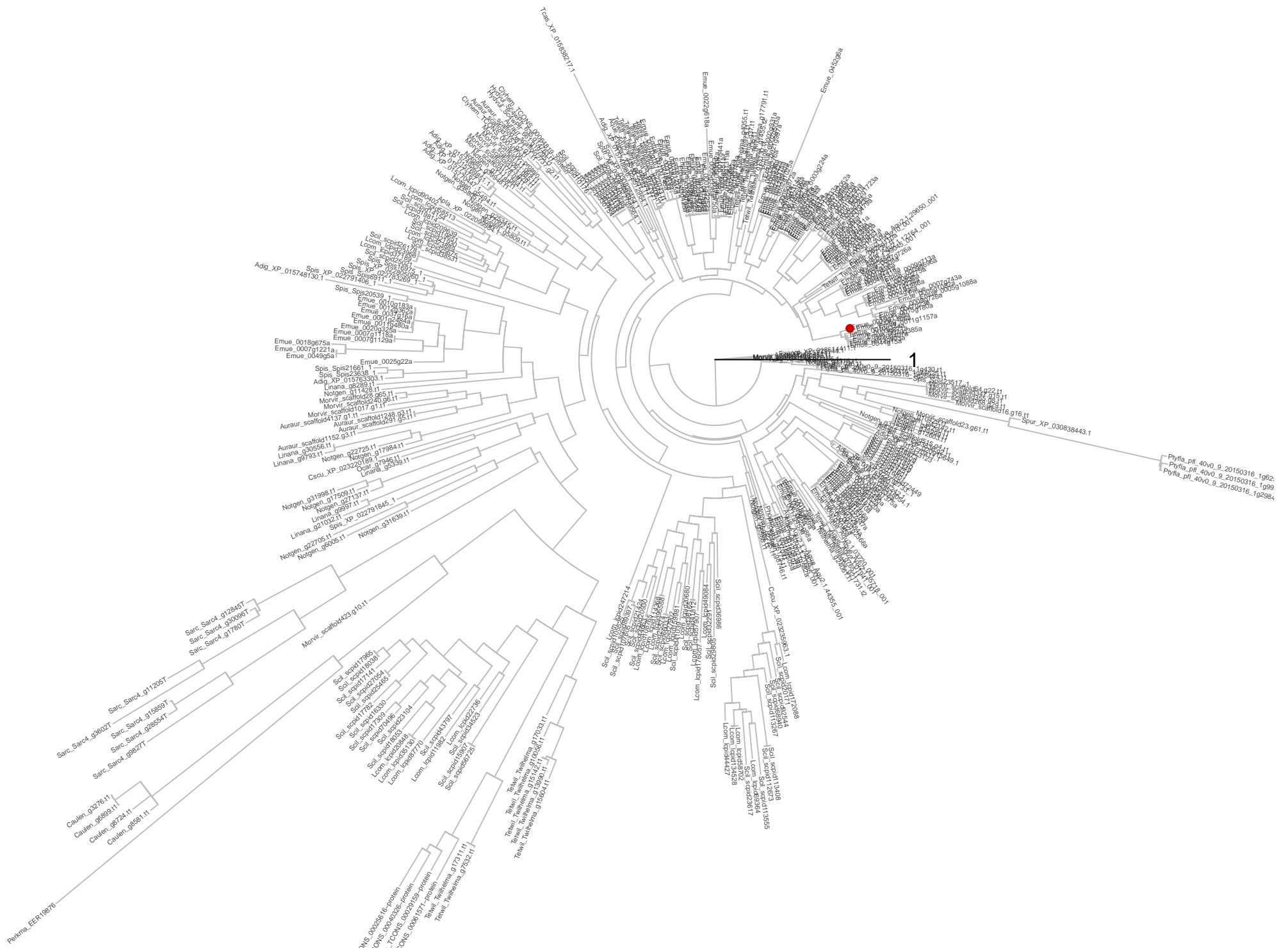
496 genes  
n=1 fusion(s) with Chromo domains  
(1 pass filters)

# RVT.HG12



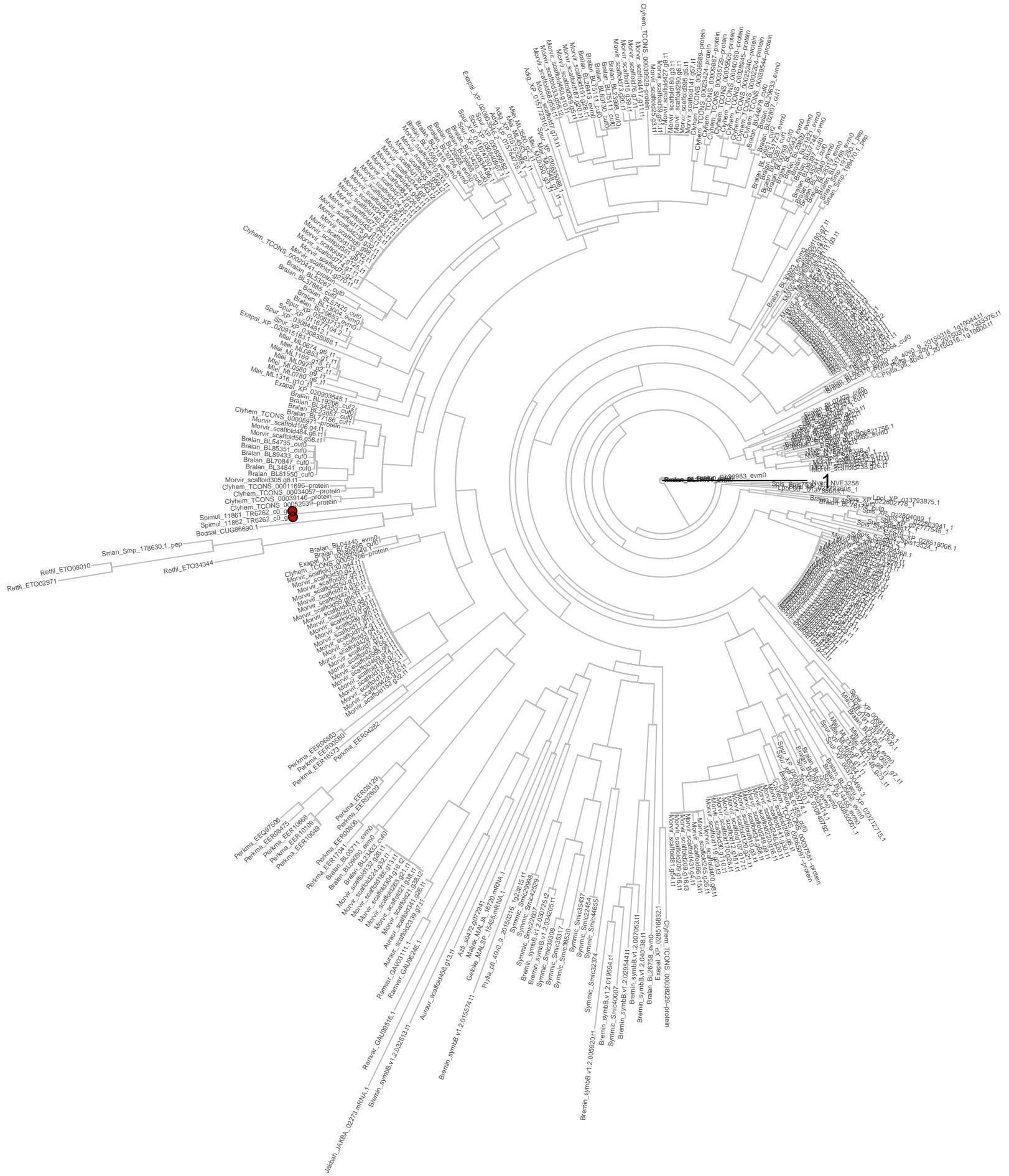
442 genes  
n=2 fusion(s) with Chromo domains  
(0 pass filters)

RVT.HG18



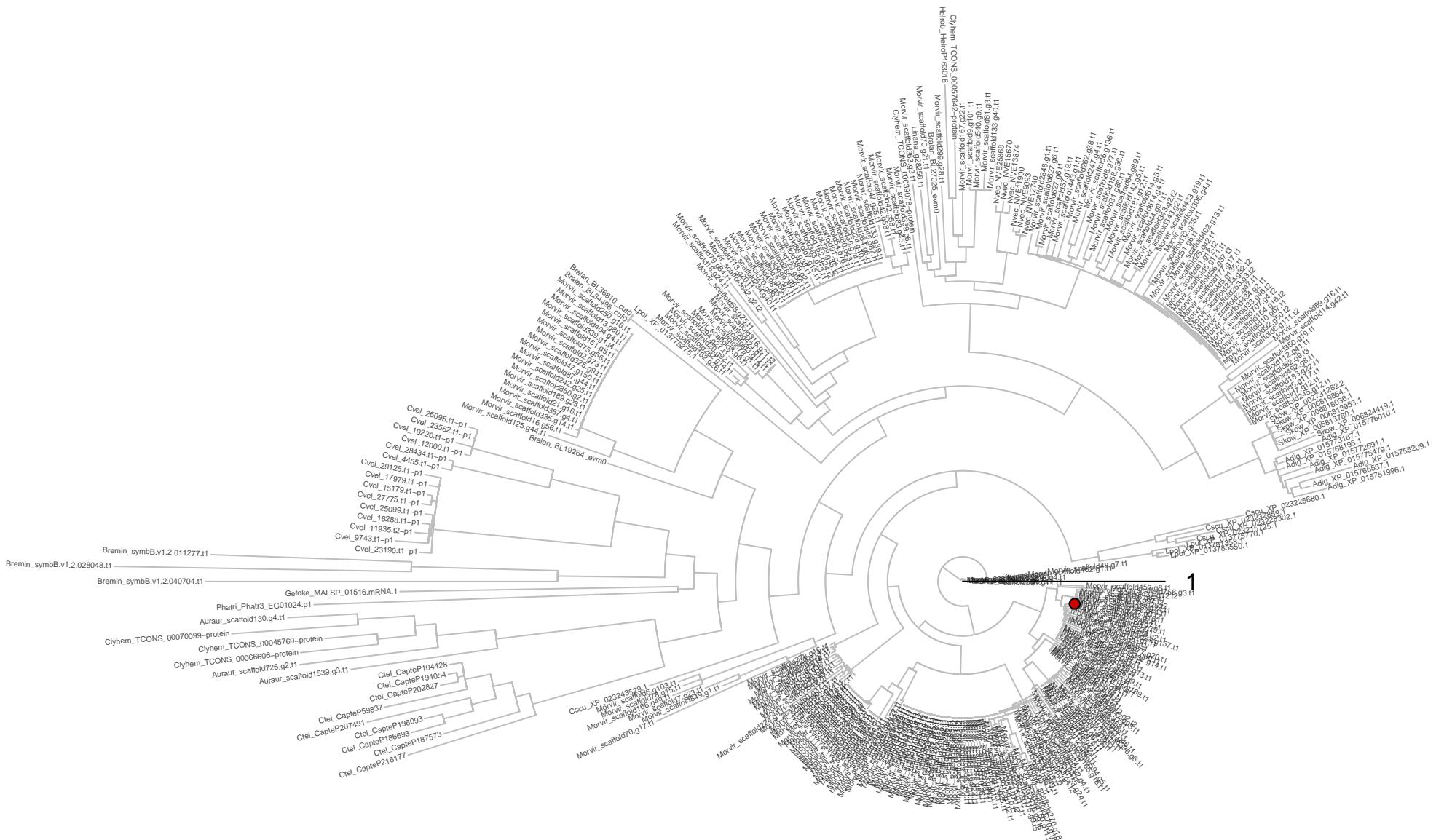
387 genes  
n=1 fusion(s) with Chromo domains  
(0 pass filters)

RVT.HG21



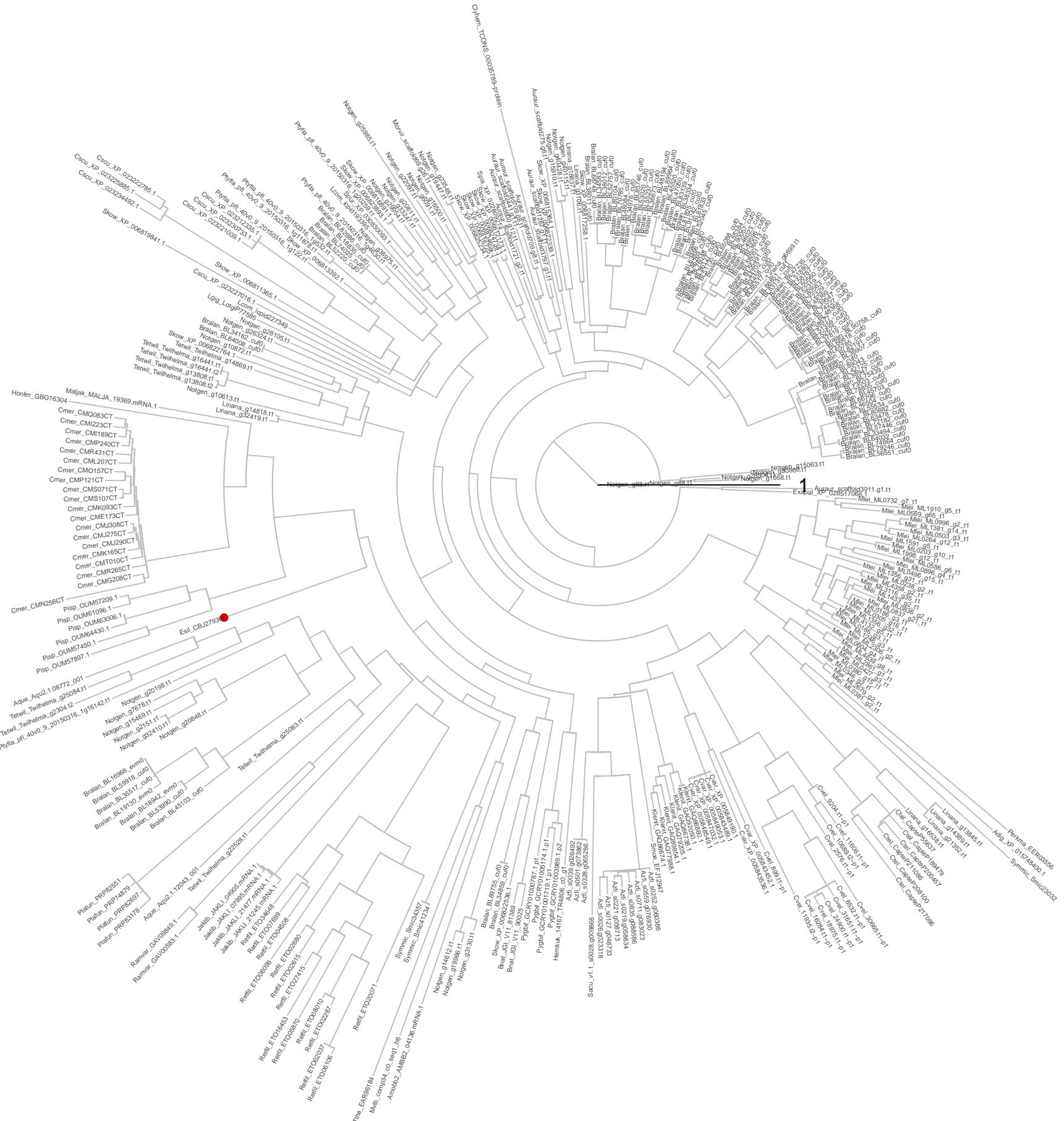
382 genes  
n=2 fusion(s) with Chromo domains  
(2 pass filters)

RVT.HG25



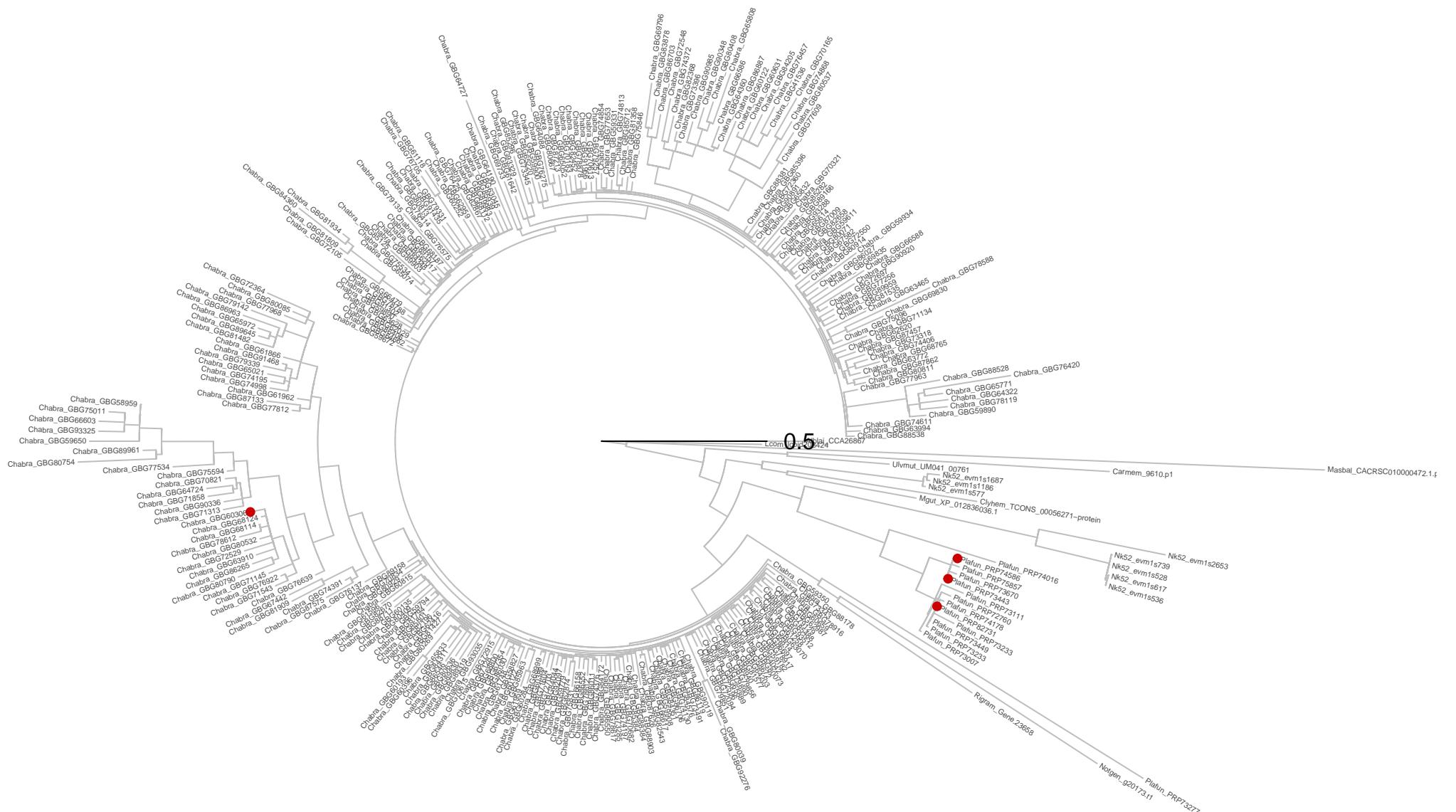
352 genes  
n=1 fusion(s) with Chromo domains  
(1 pass filters)

RVT.HG26



341 genes  
n=1 fusion(s) with Chromo domains  
(0 pass filters)

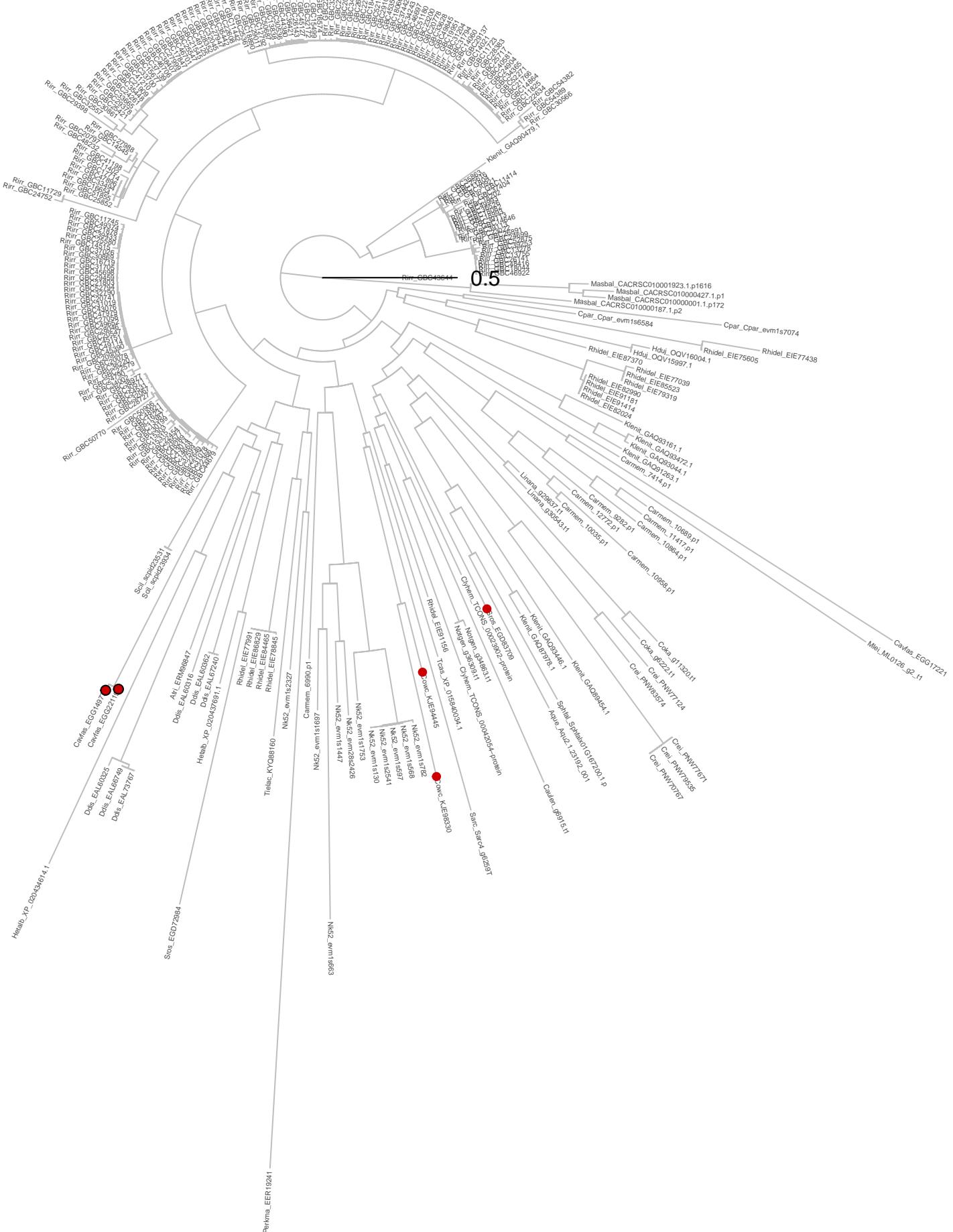
RVT.HG29



- Chromo HG2.1:NA n = 4

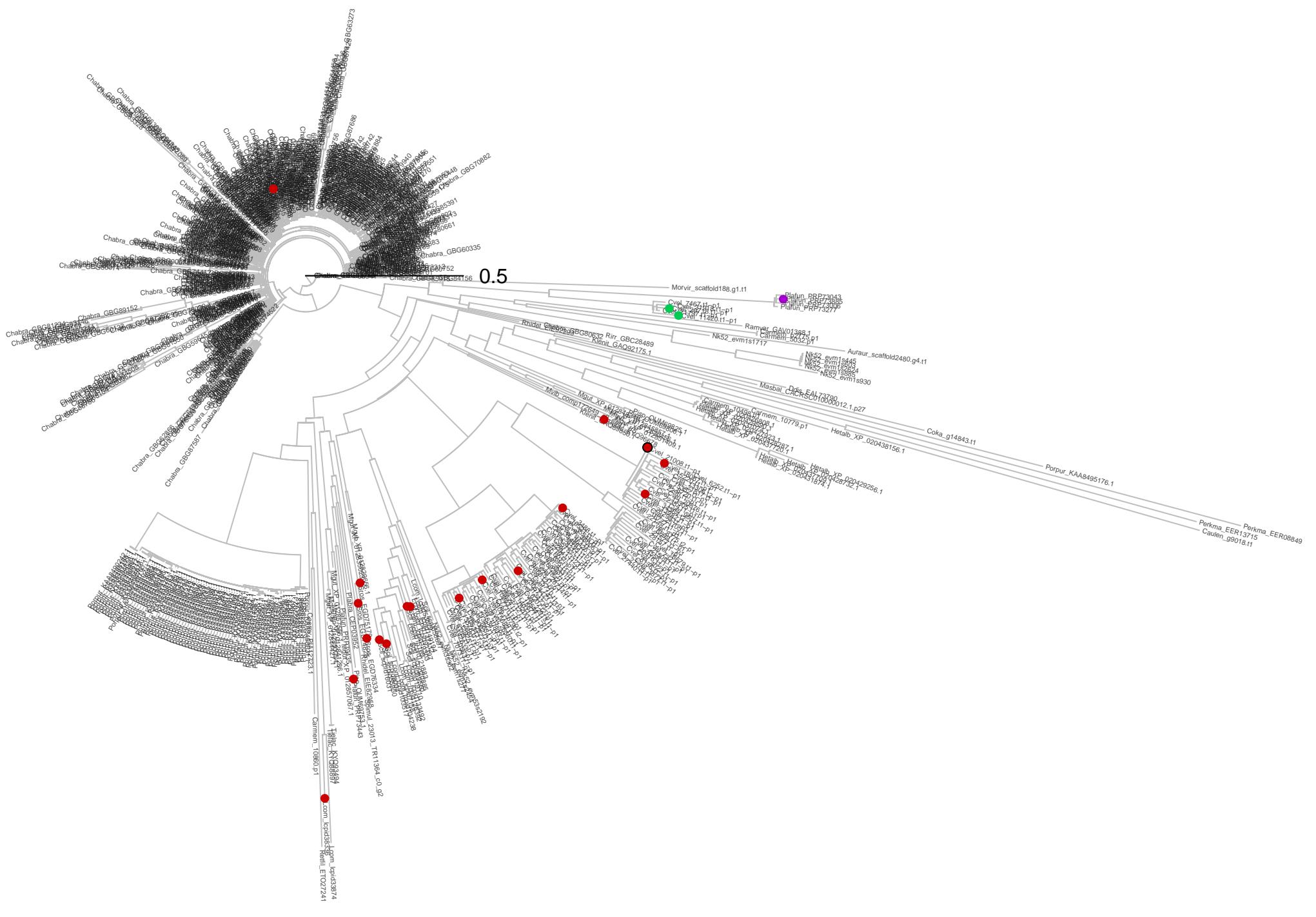
316 genes  
n=4 fusion(s) with Chromo domains  
(0 pass filters)

RVT.HG33



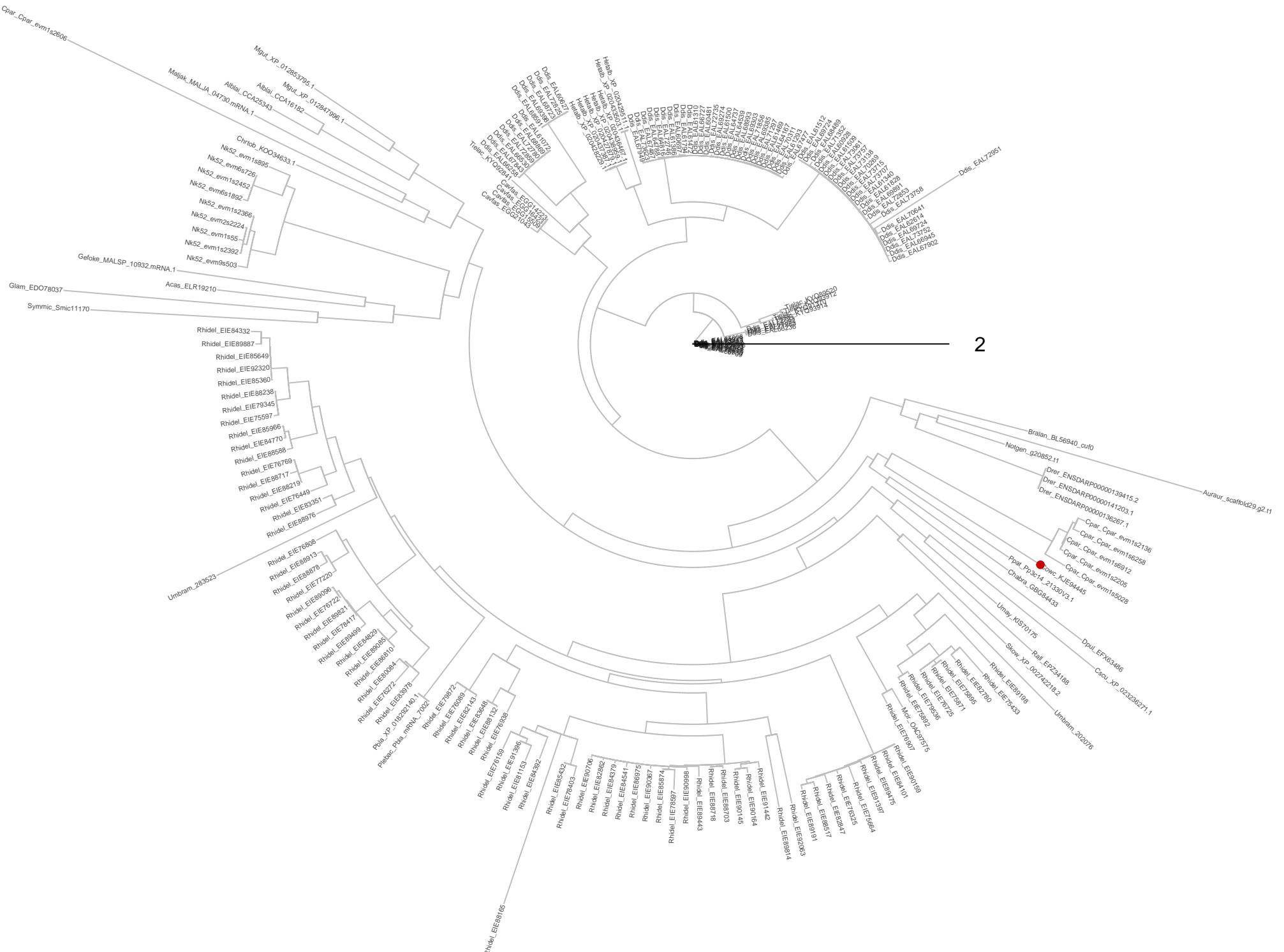
266 genes  
n=5 fusion(s) with Chromo domains  
(2 pass filters)

# RVT.HG4



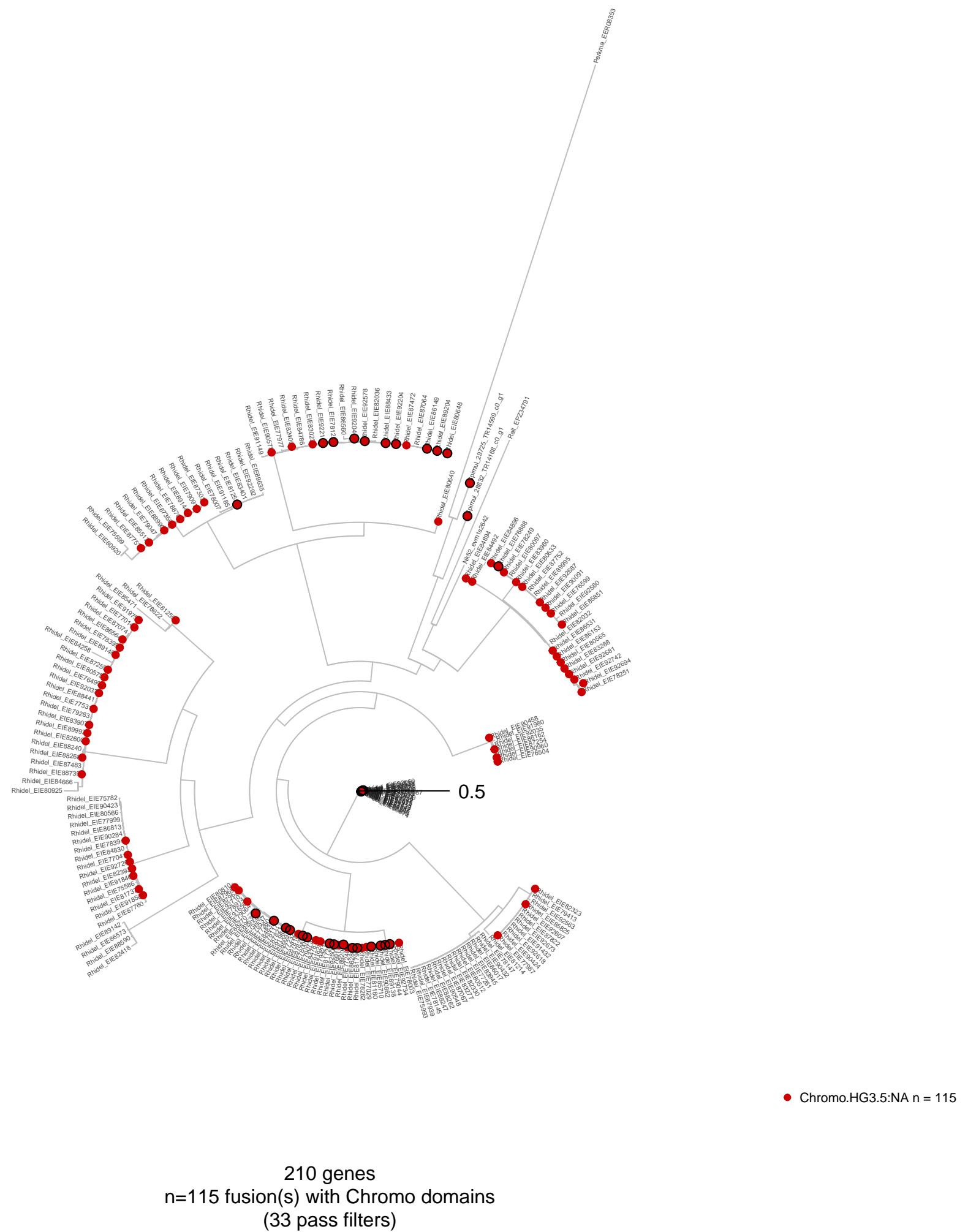
770 genes  
n=21 fusion(s) with Chromo domains  
(1 pass filters)

RVT.HG40

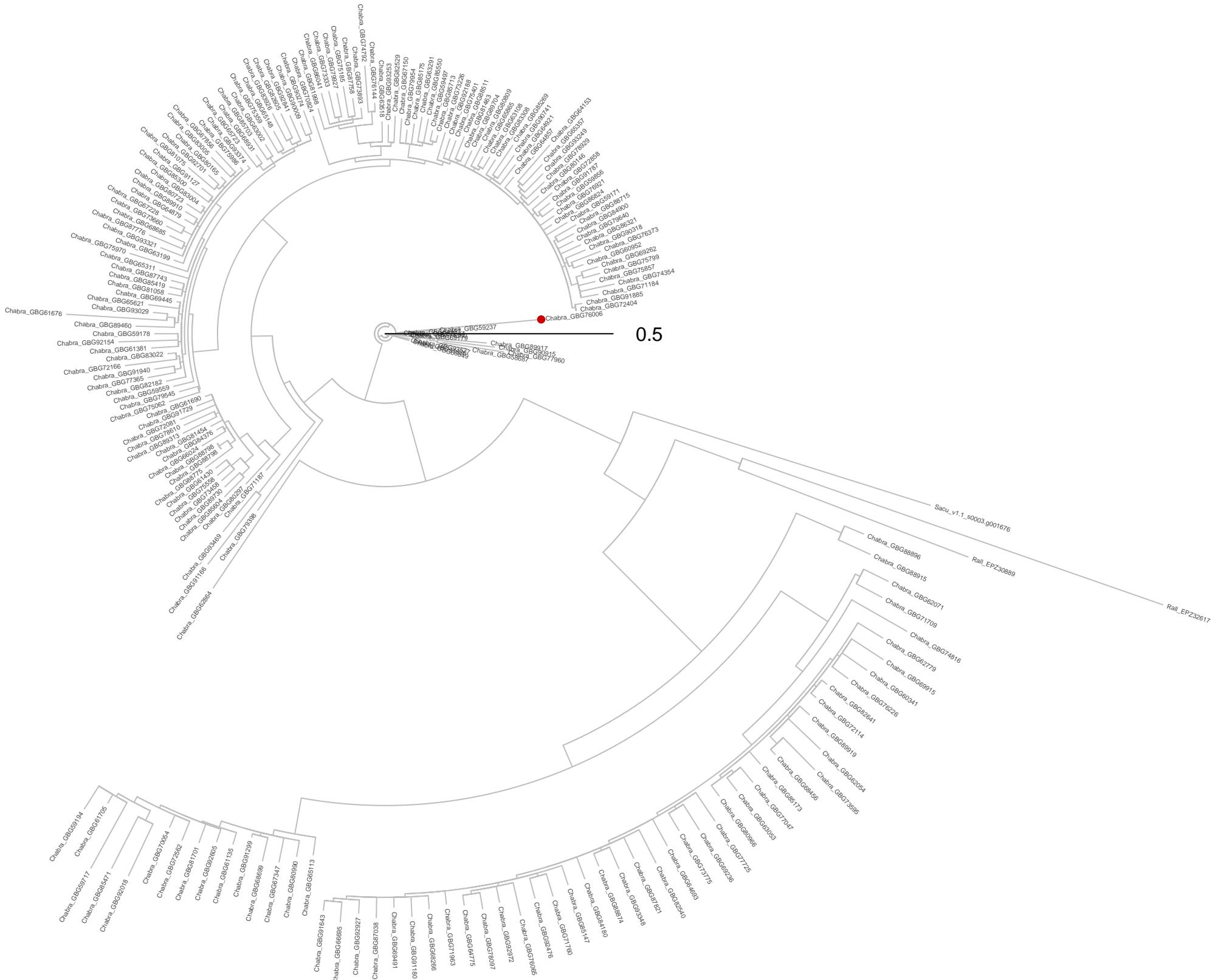


222 genes  
n=1 fusion(s) with Chromo domains  
(0 pass filters)

# RVT.HG41



RVT.HG43



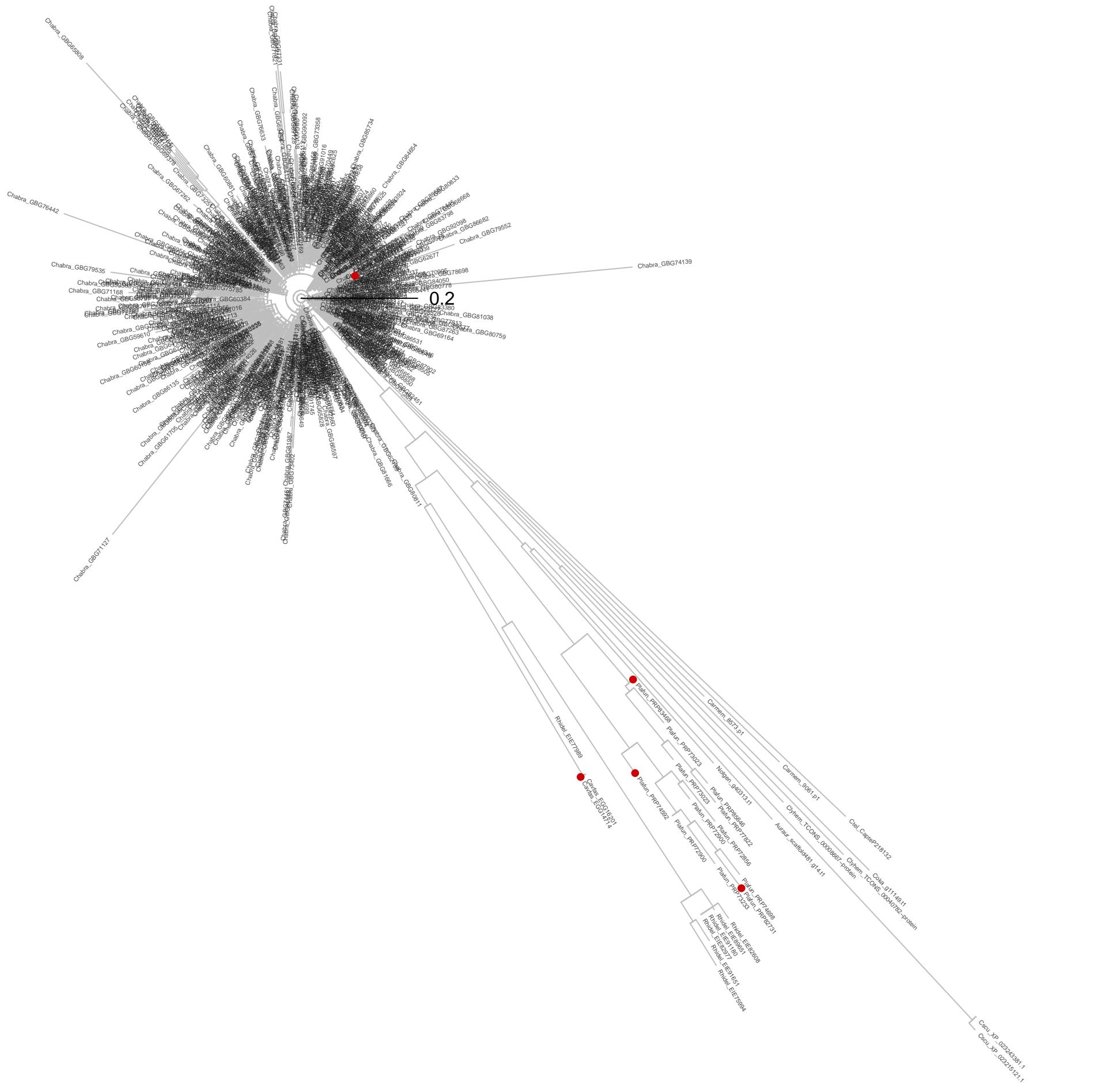
206 genes  
n=1 fusion(s) with Chromo domains  
(0 pass filters)

RVT.HG50

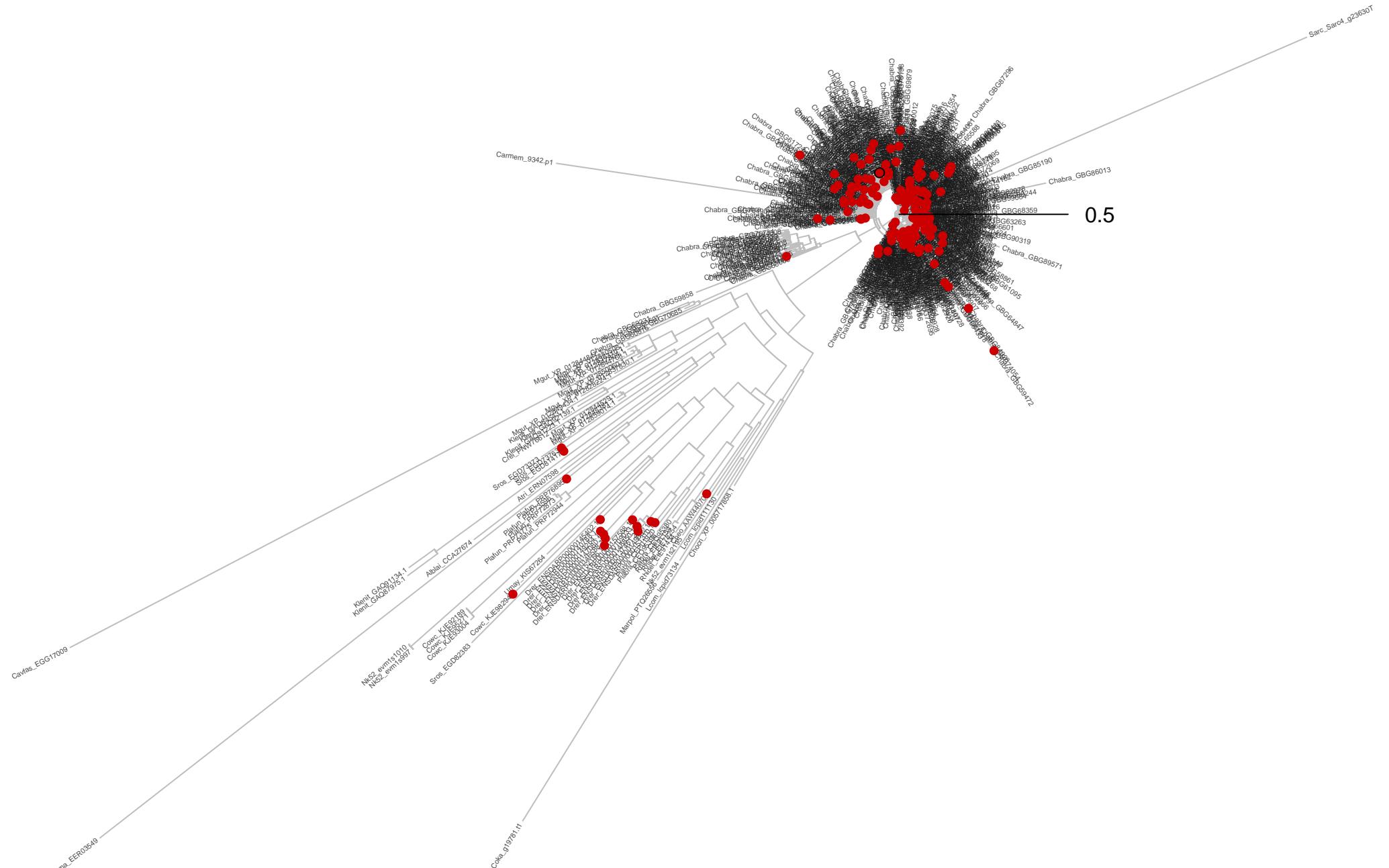


155 genes  
n=1 fusion(s) with Chromo domains  
(1 pass filters)

# RVT.HG6



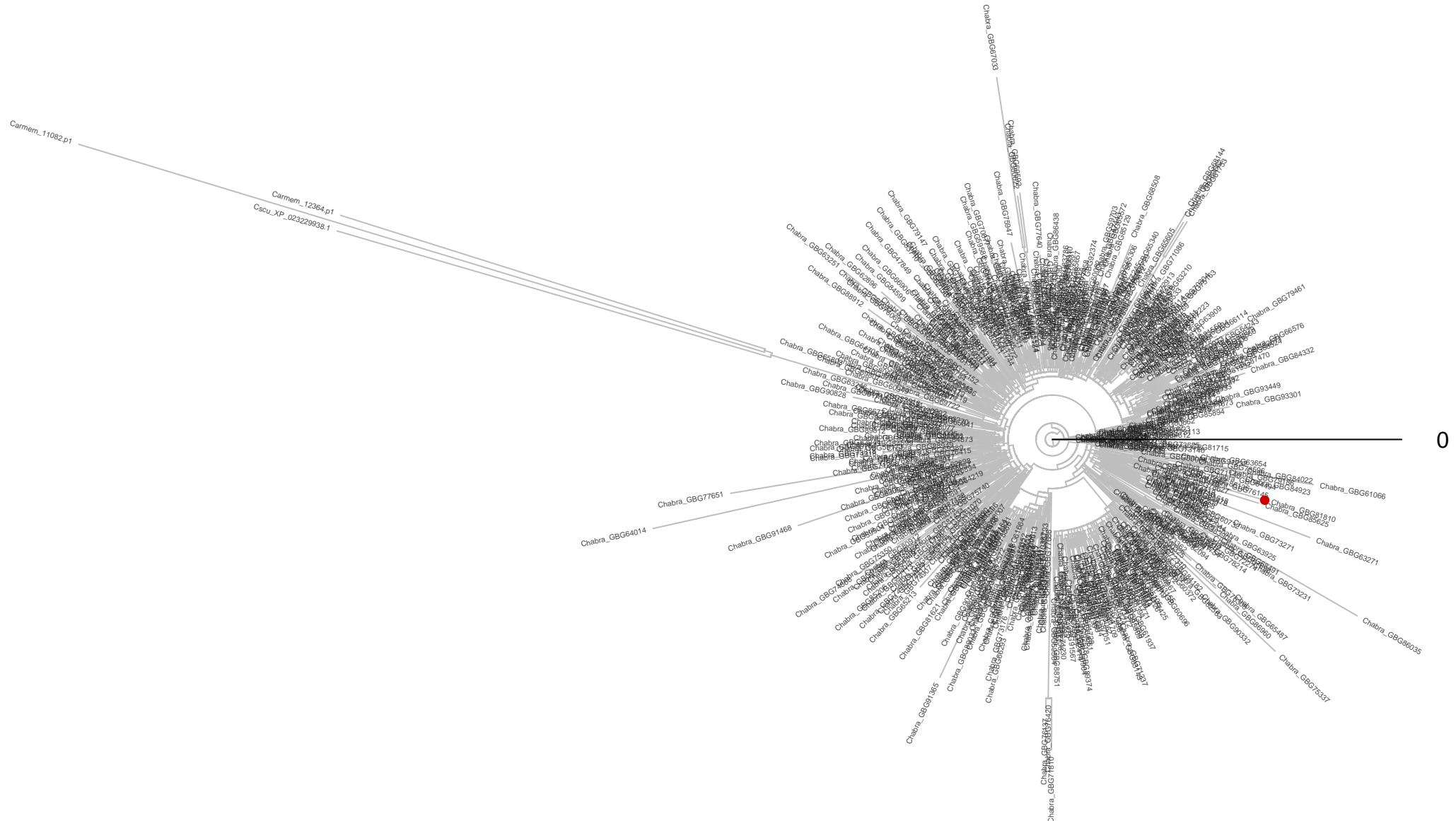
# RVT.HG7



659 genes  
n=152 fusion(s) with Chromo domains  
(1 pass filters)

● Chromo.HG2.1:NA n = 152

RVT.HG8



587 genes  
n=1 fusion(s) with Chromo domains  
(0 pass filters)