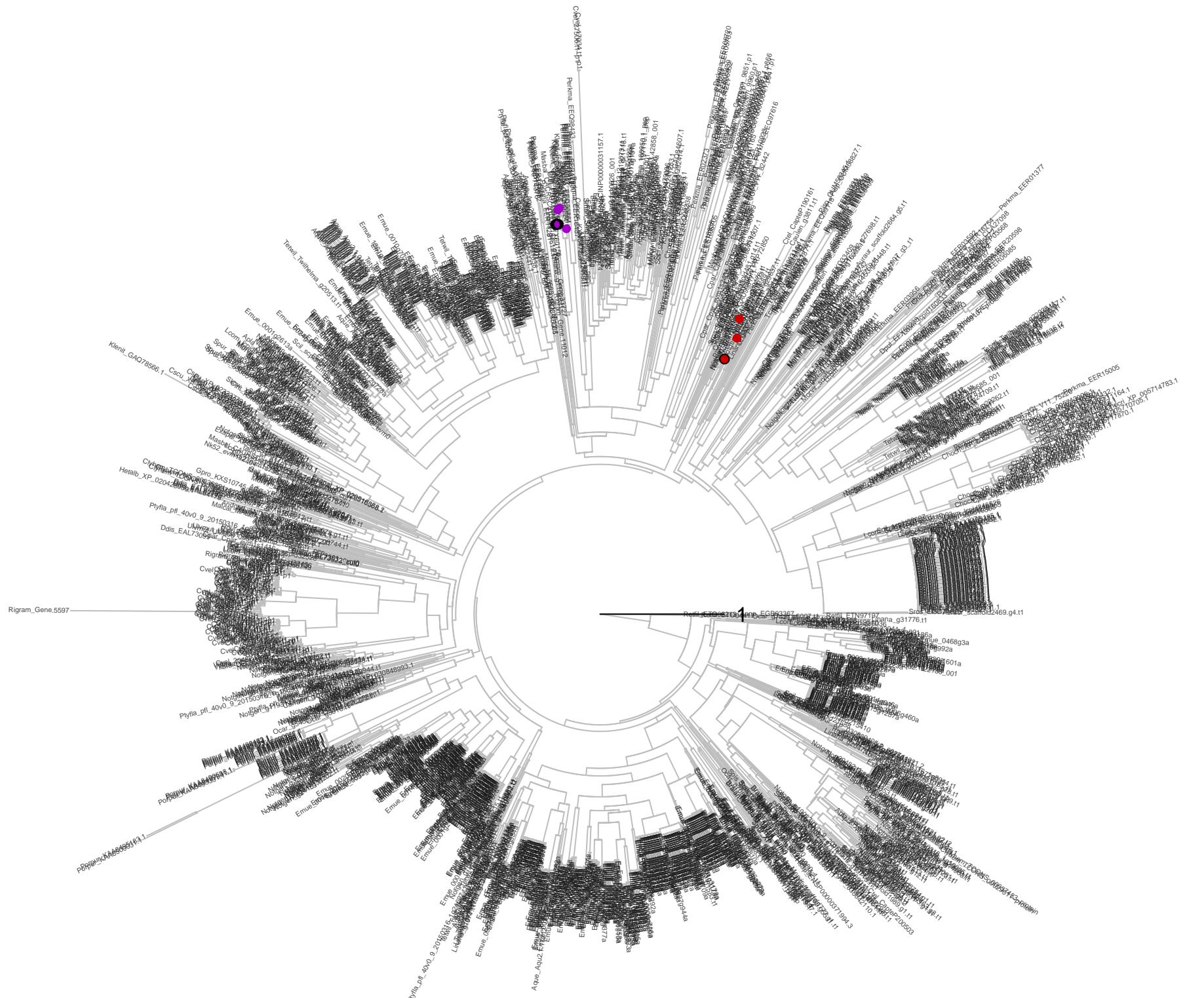
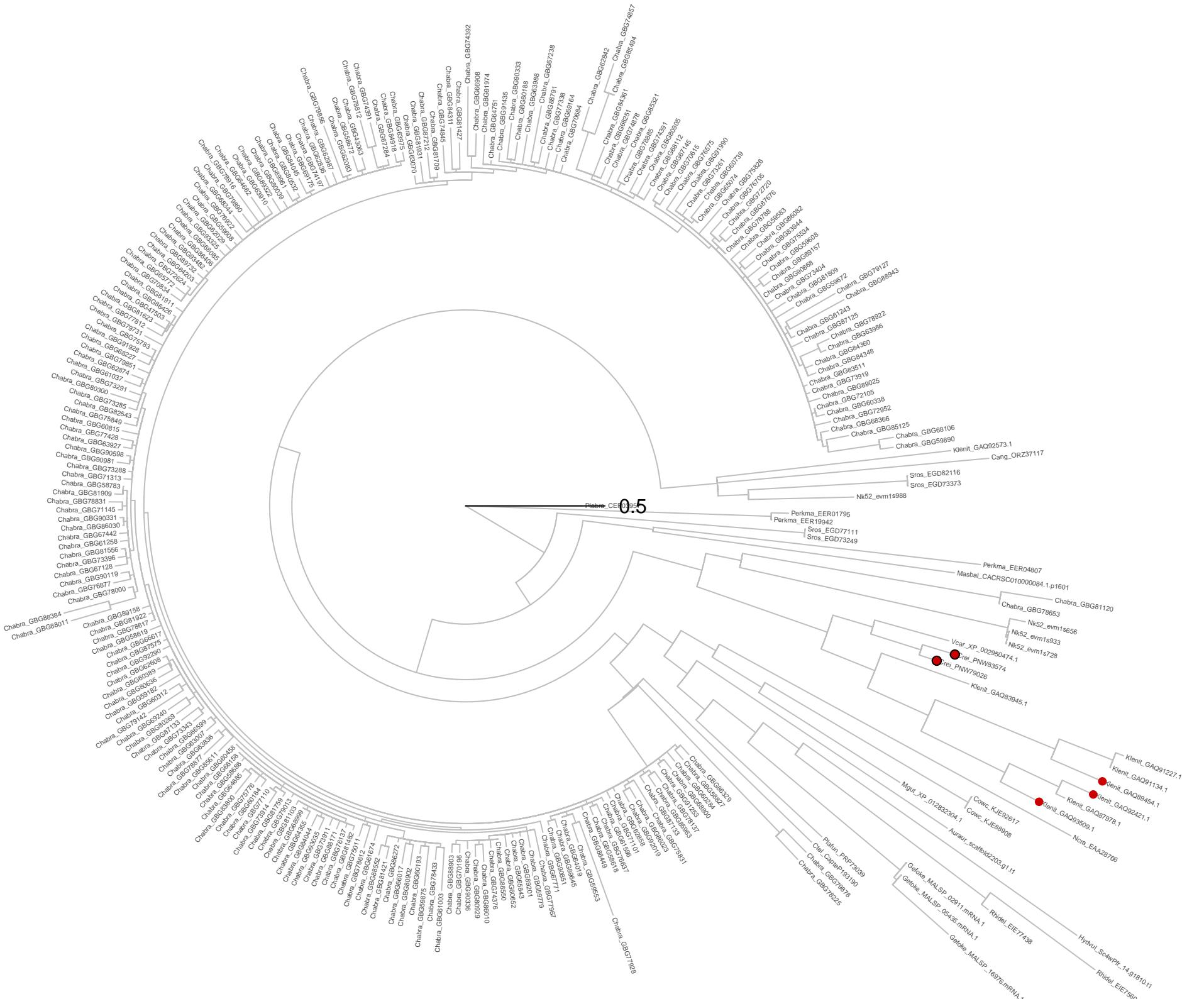


## Integrase\_H2C2.HG1



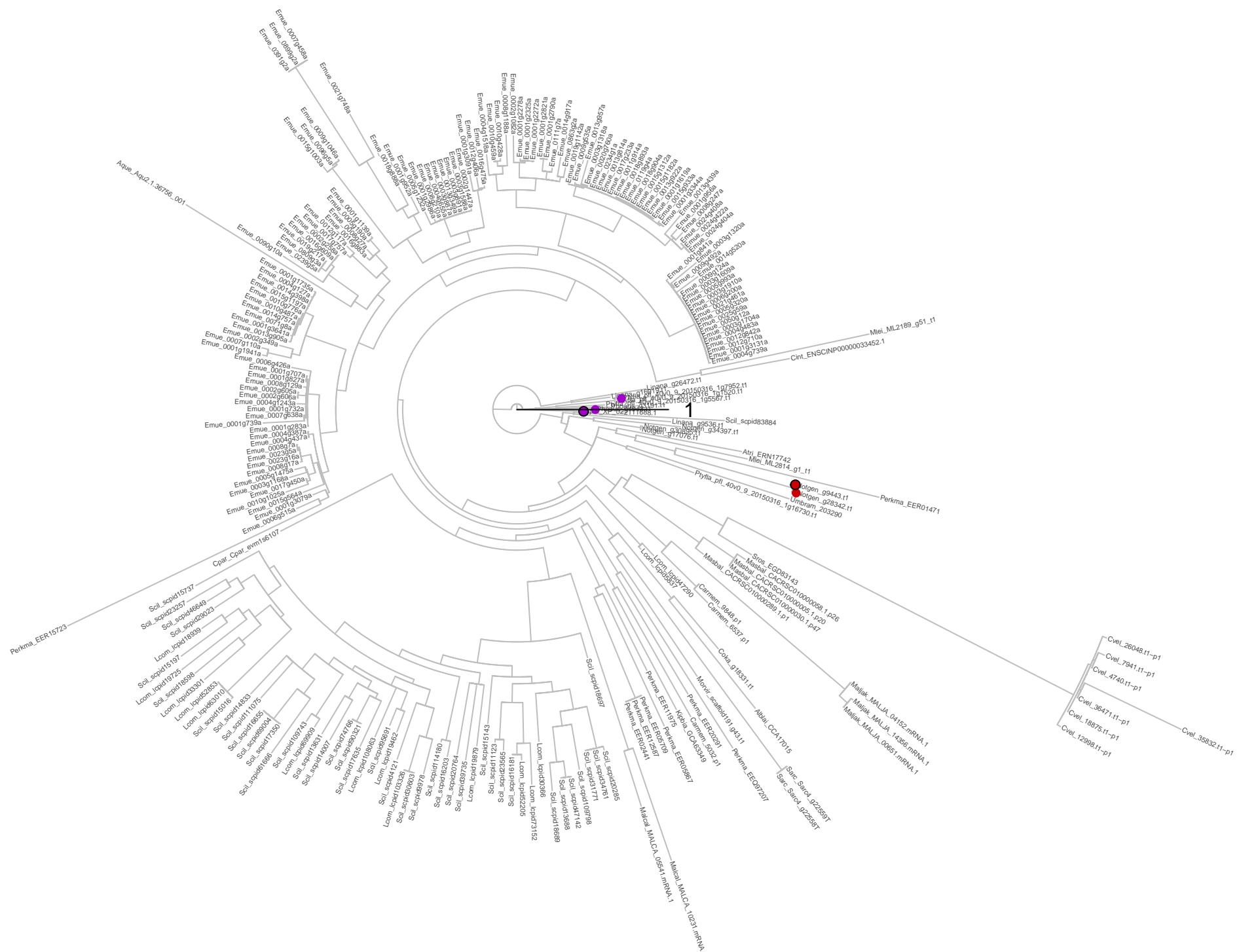
1906 genes  
n=9 fusion(s) with PHD domains  
(4 pass filters)

# Integrase\_H2C2.HG11



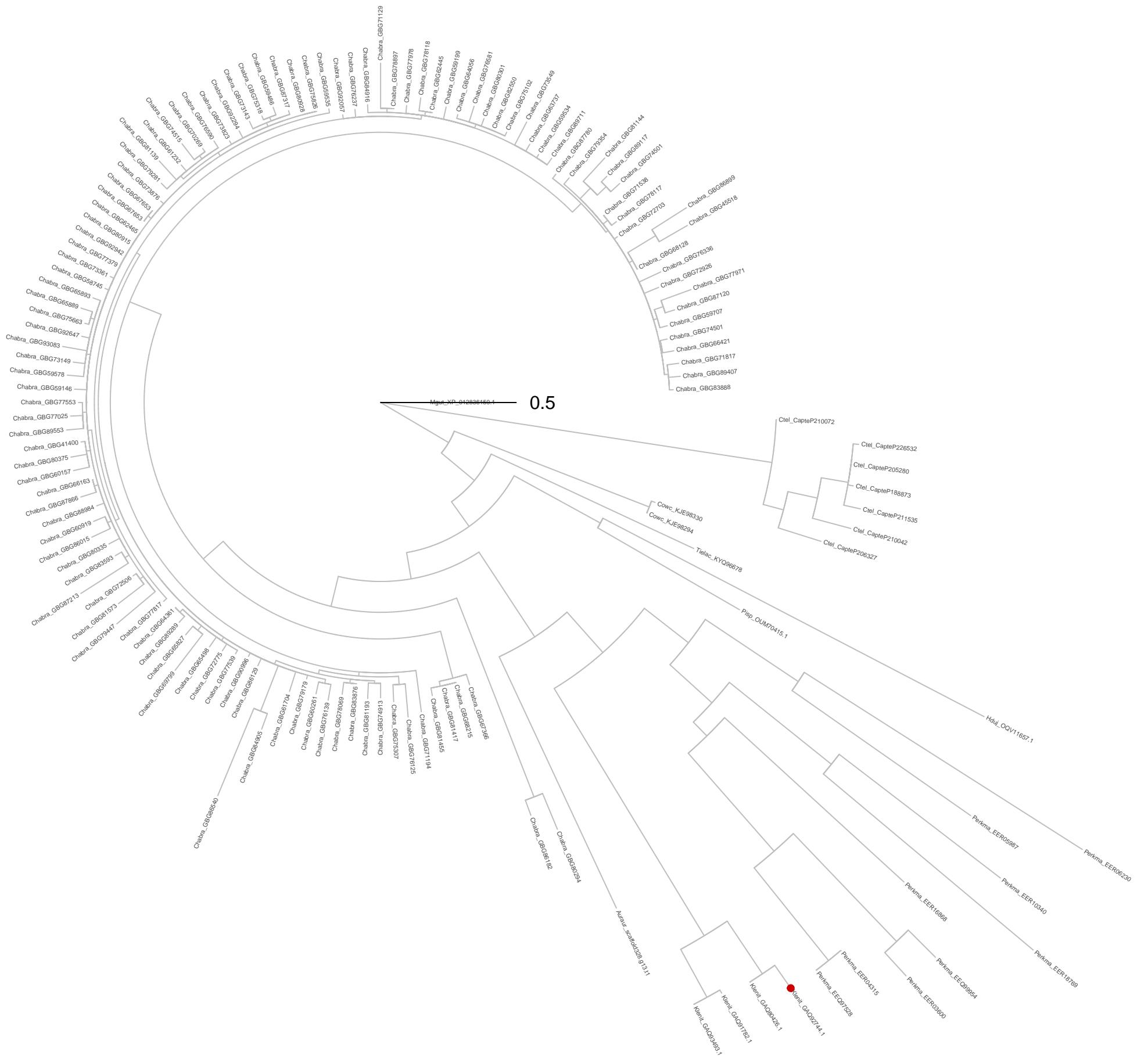
277 genes  
n=5 fusion(s) with PHD domains  
(2 pass filters)

## Integrase\_H2C2.HG13



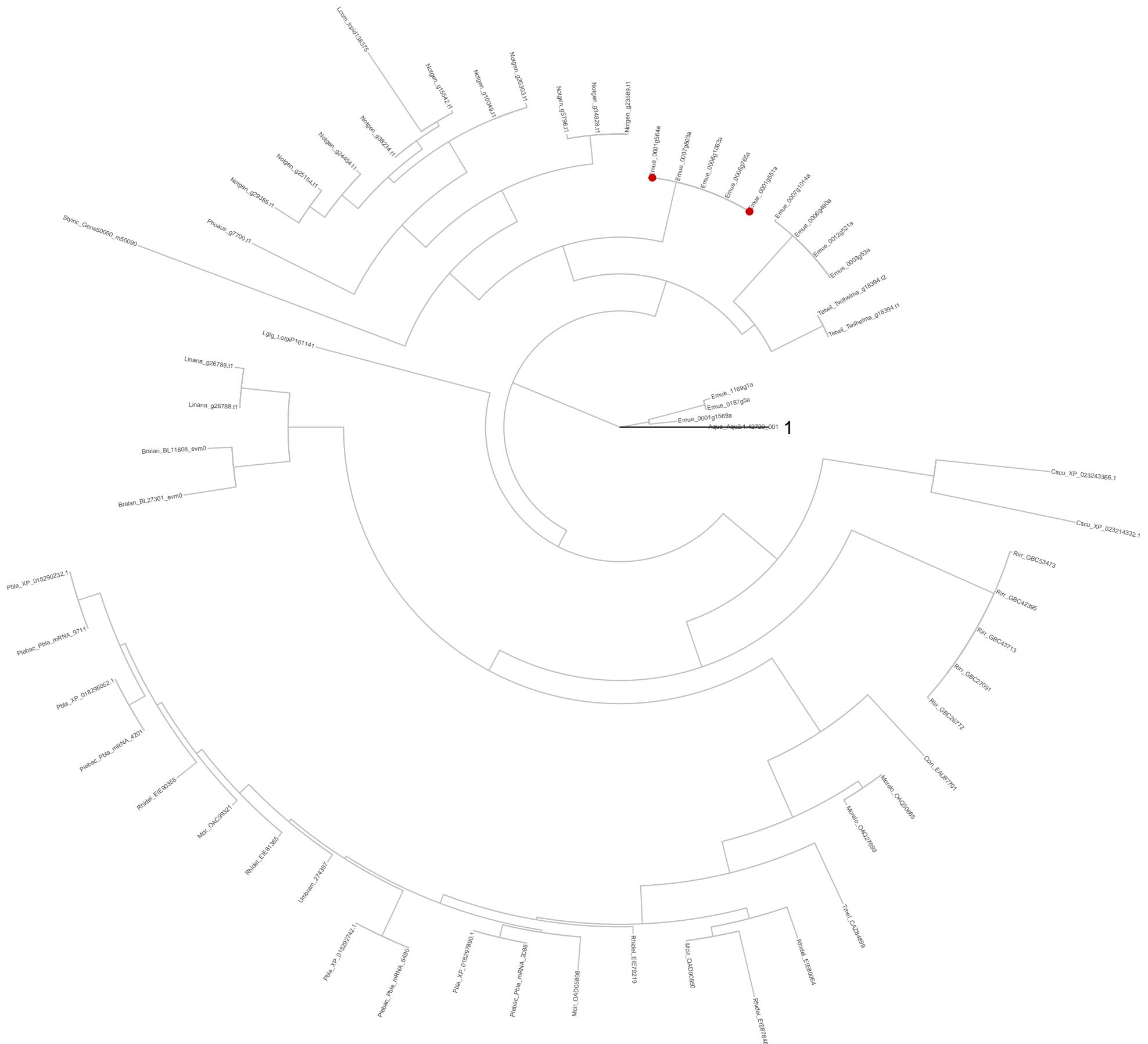
238 genes  
n=5 fusion(s) with PHD domains  
(2 pass filters)

## **Integrase\_H2C2.HG16**



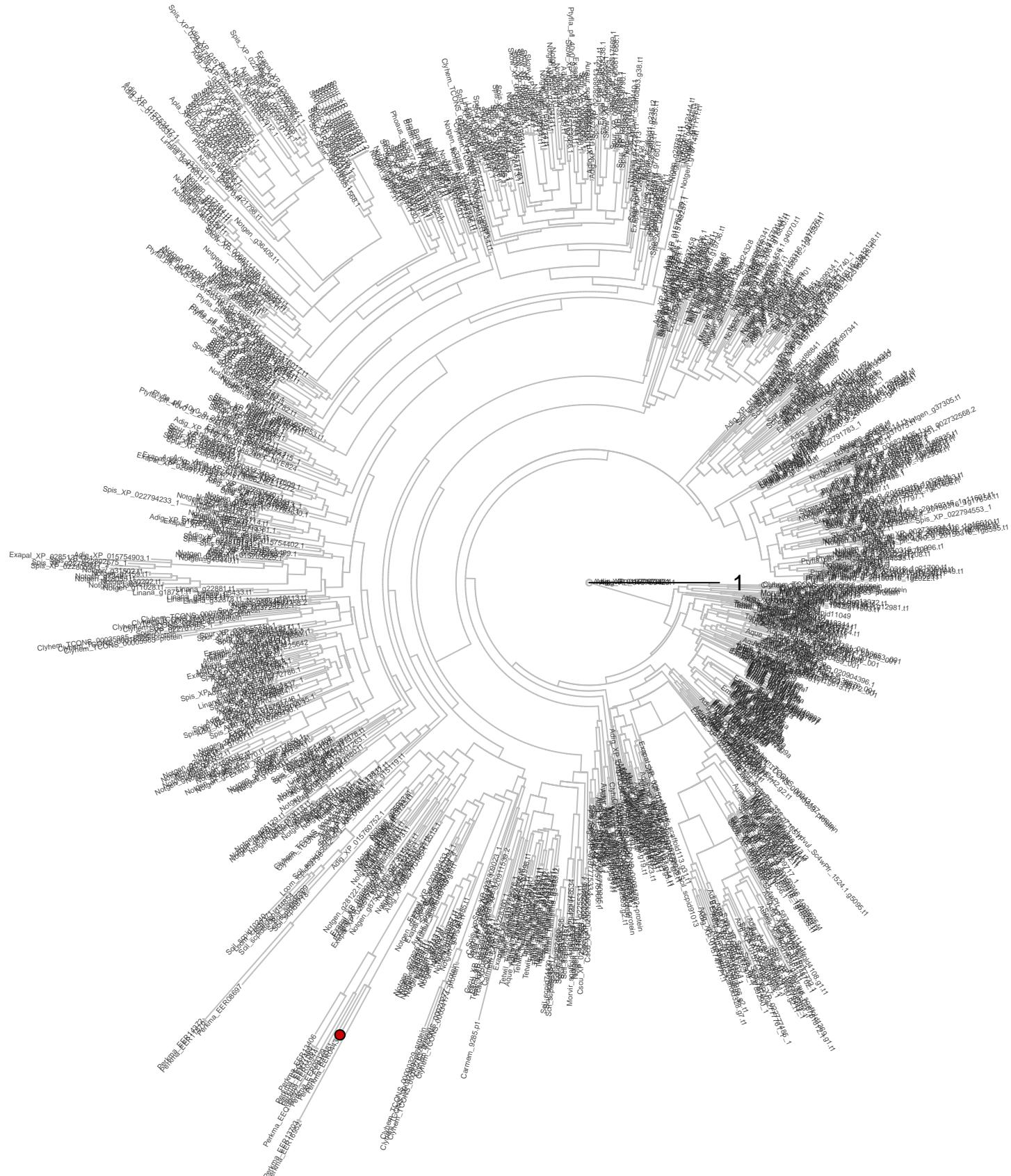
144 genes  
n=1 fusion(s) with PHD domains  
(0 pass filters)

## **Integrase\_H2C2.HG19**



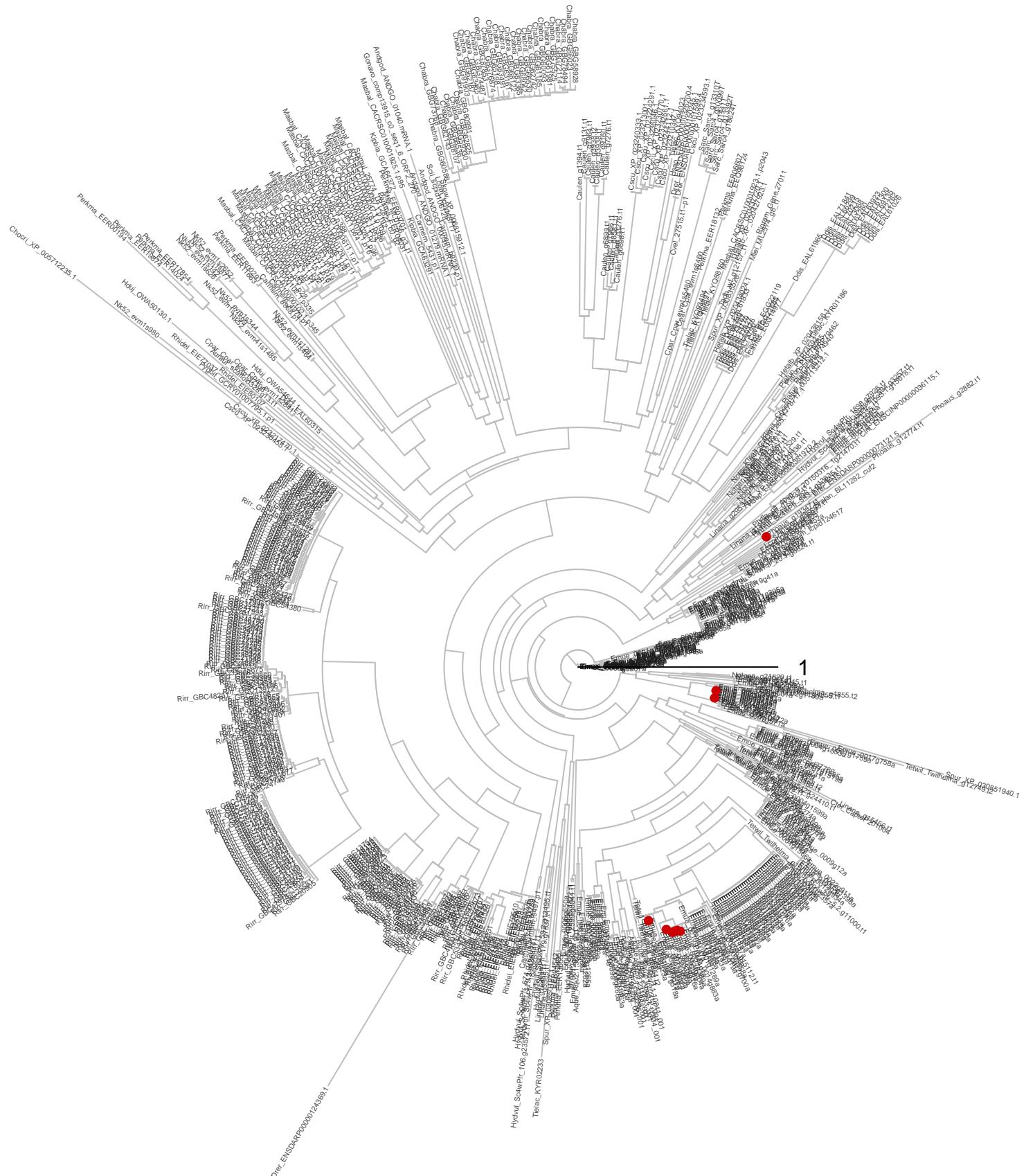
61 genes  
n=2 fusion(s) with PHD domains  
(0 pass filters)

## **Integrase\_H2C2.HG3**



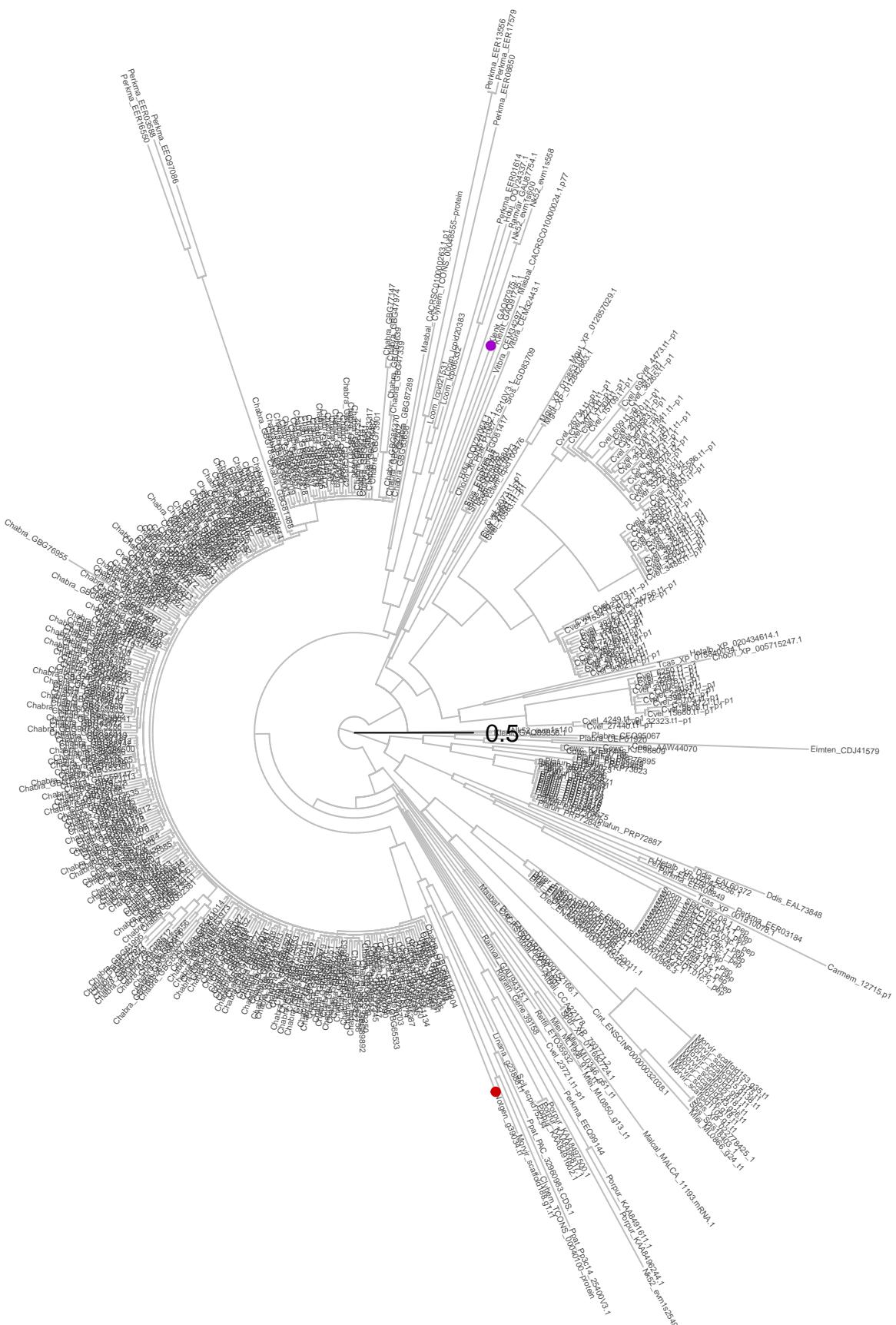
1054 genes  
n=1 fusion(s) with PHD domains  
(1 pass filters)

## **Integrase\_H2C2.HG5**



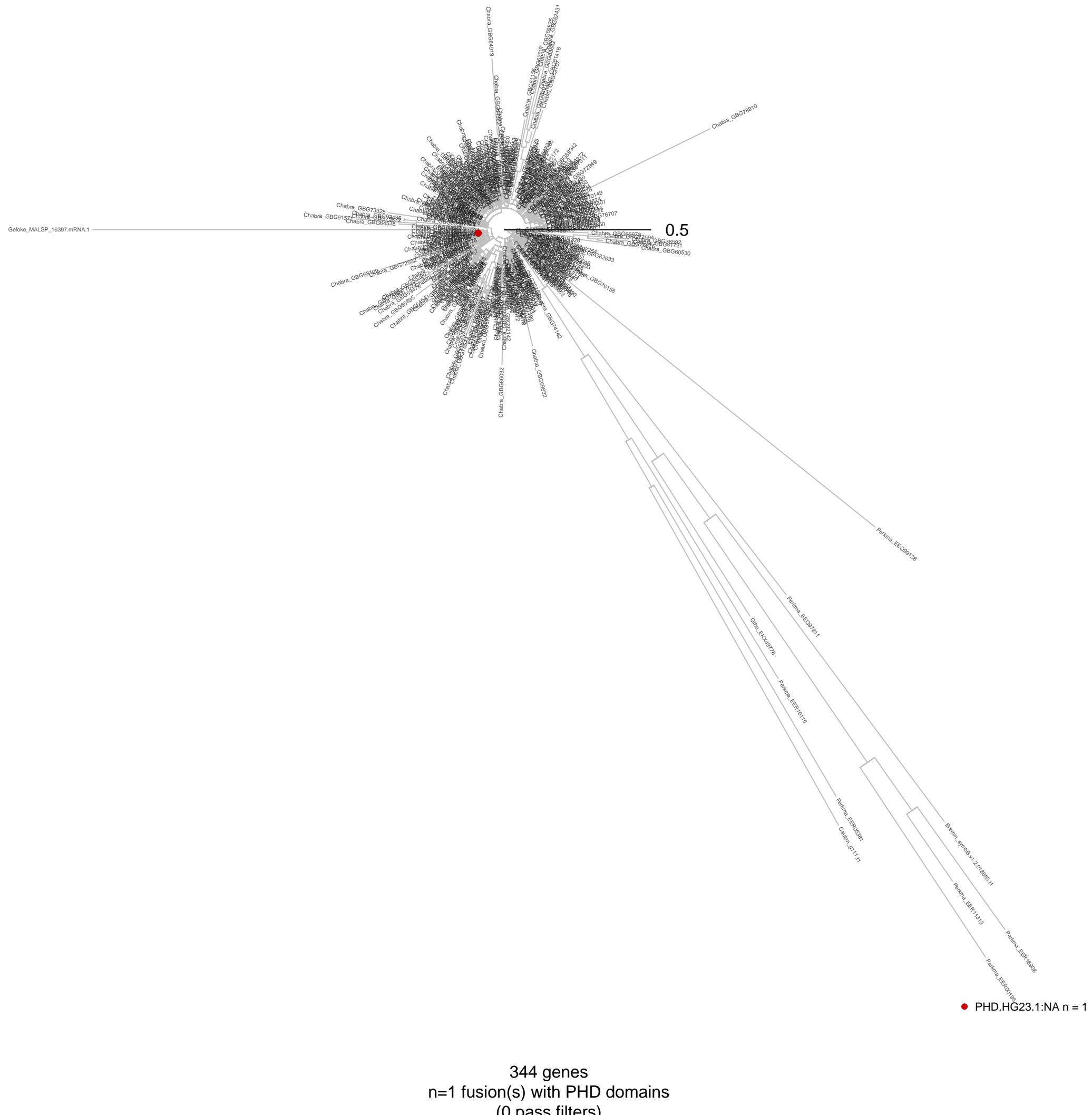
706 genes  
n=9 fusion(s) with PHD domains  
(0 pass filters)

## Integrase\_H2C2.HG7

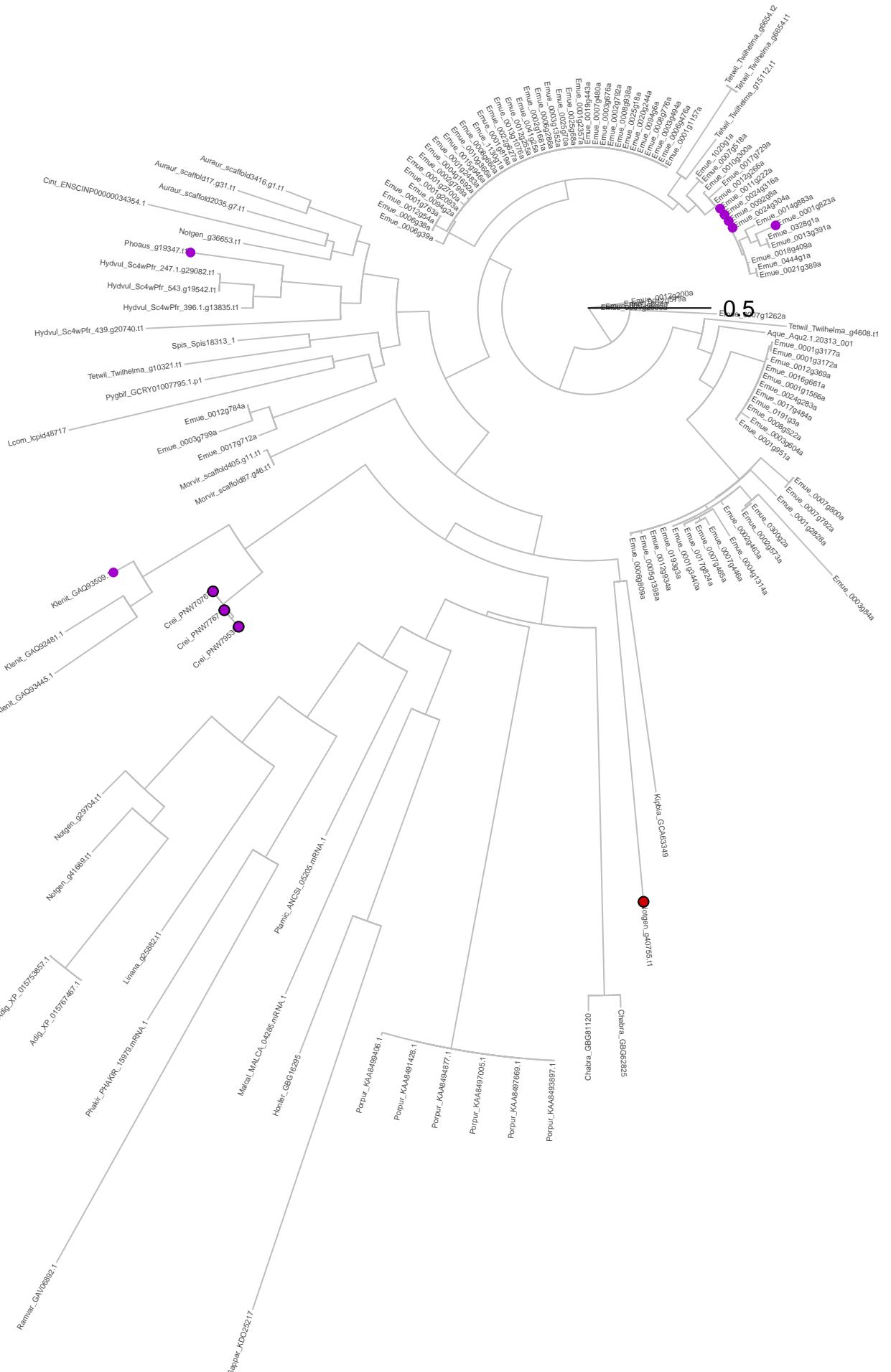


520 genes  
n=2 fusion(s) with PHD domains  
(0 pass filters)

# Integrase\_H2C2.HG8

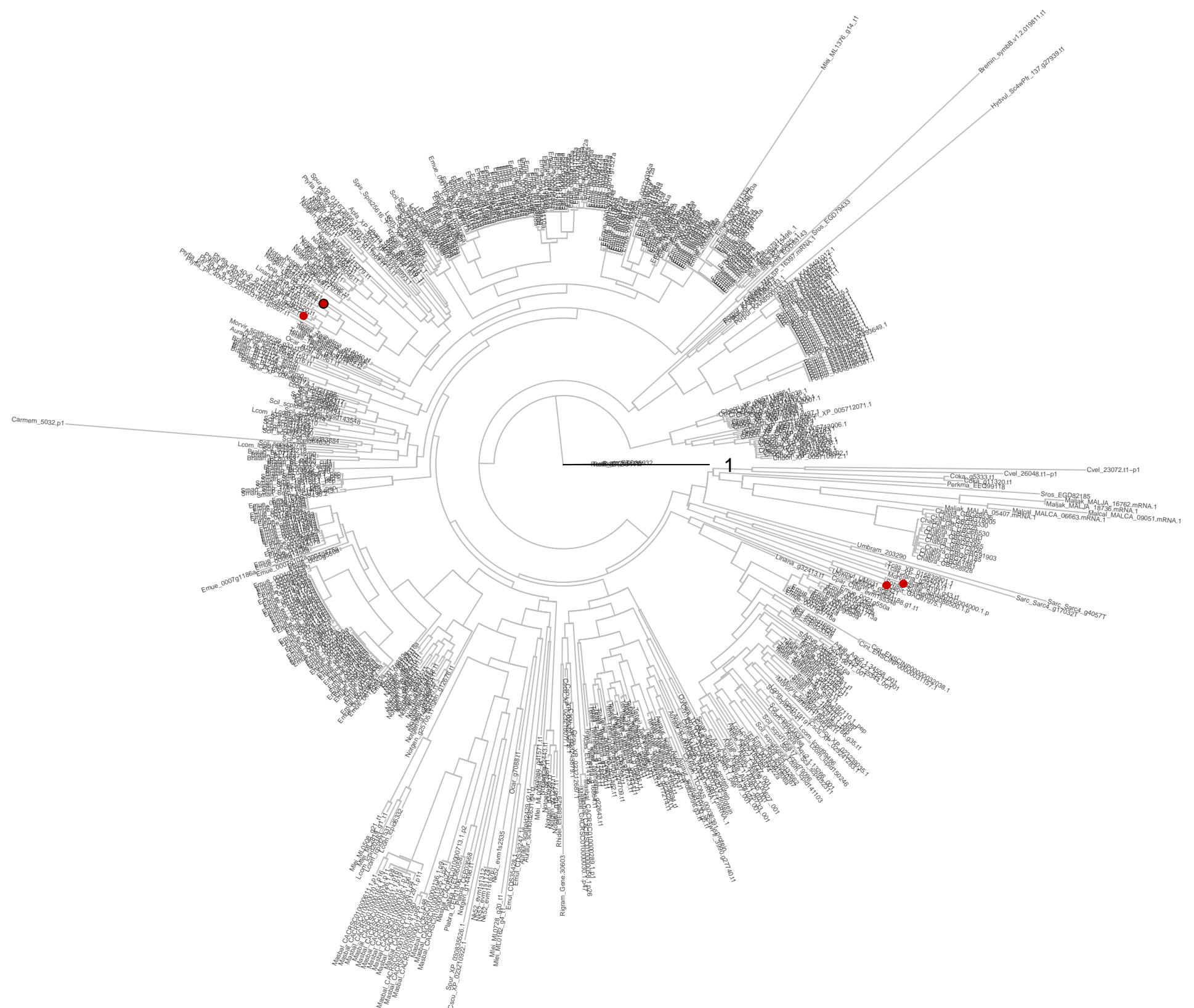


rve.HG16



136 genes  
n=11 fusion(s) with PHD domains  
(4 pass filters)

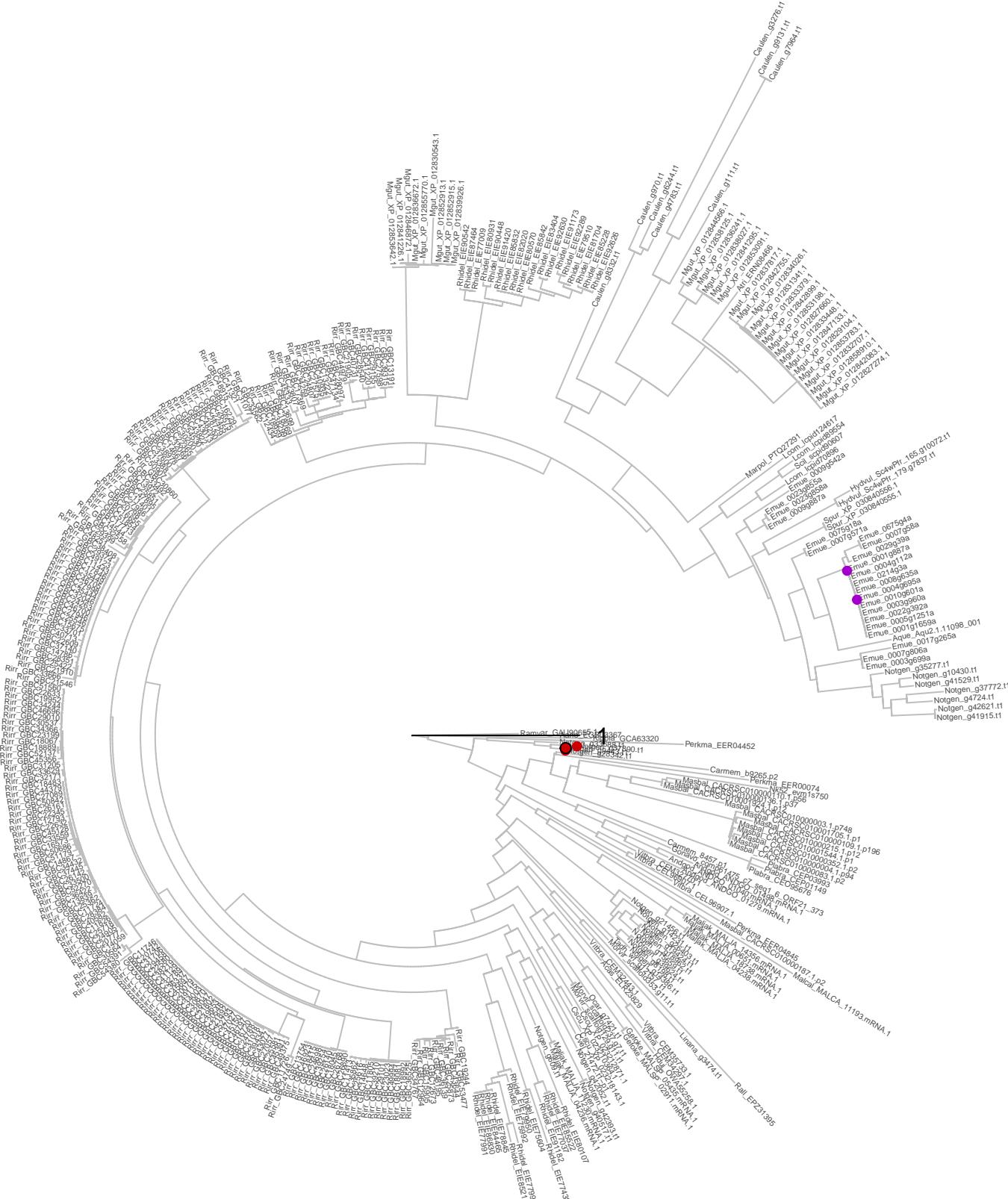
# rve.HG2



616 genes  
n=4 fusion(s) with PHD domains  
(1 pass filters)

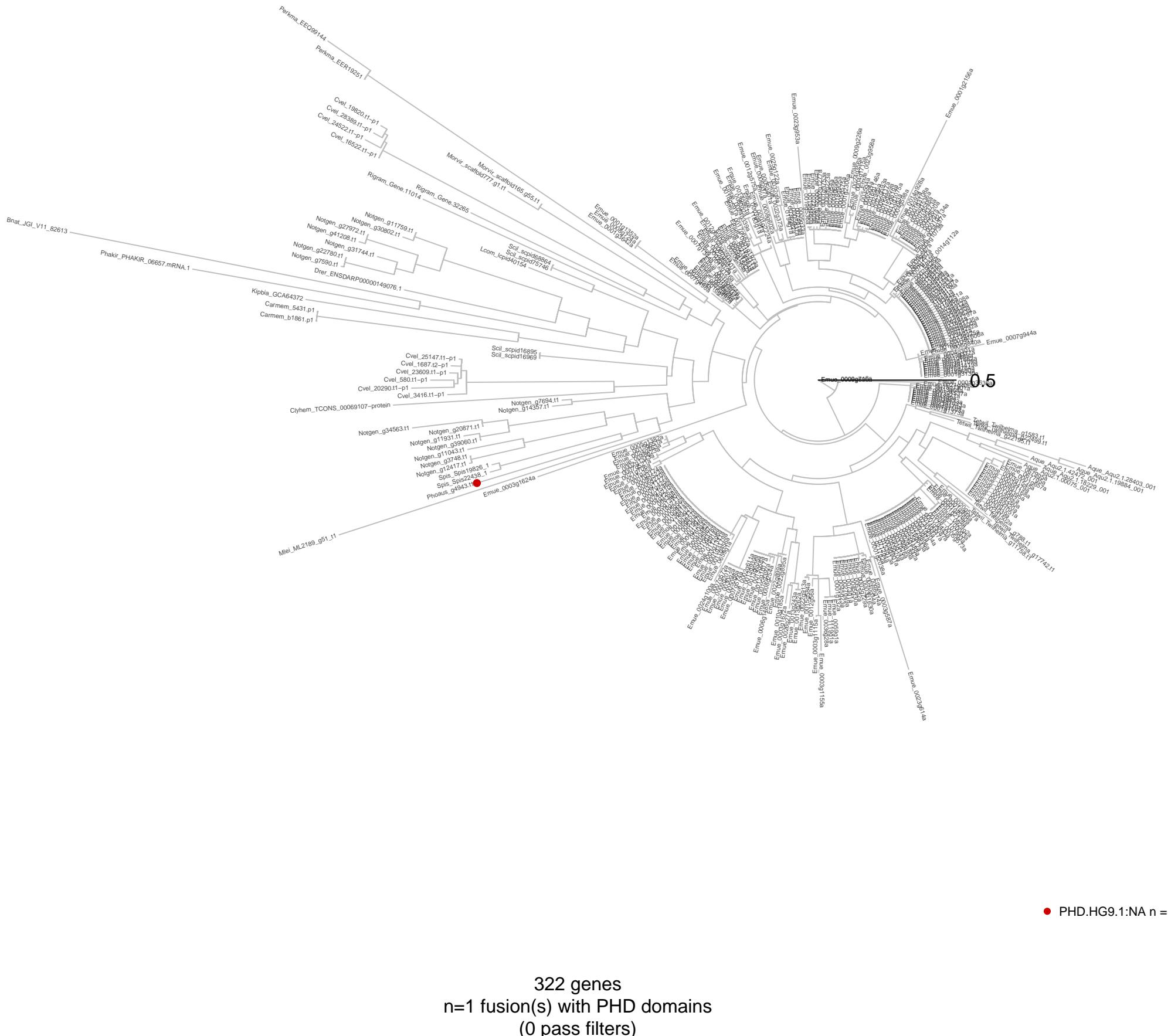
● PHD.HG9.1:NA n = 4

rve.HG4

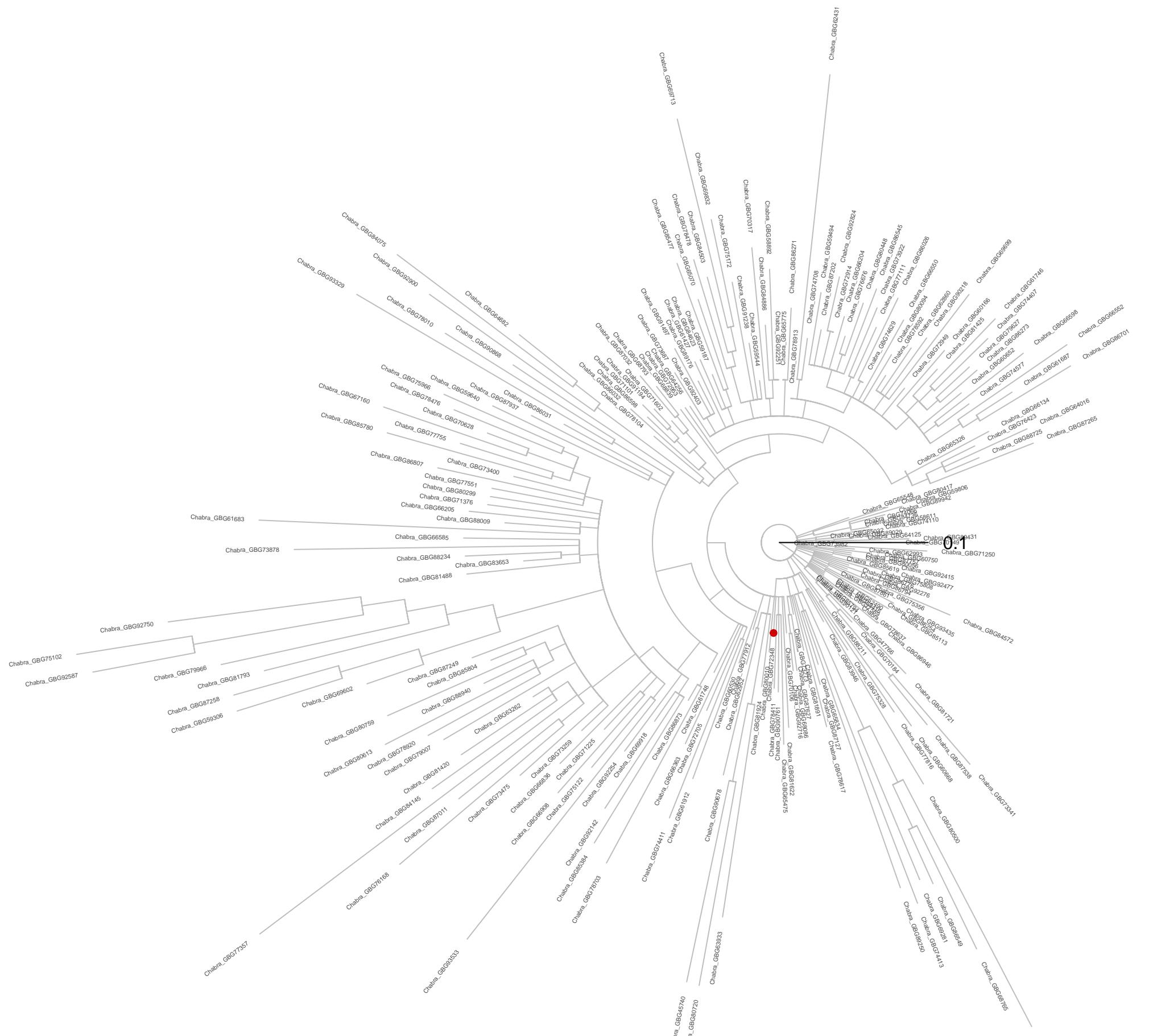


382 genes  
n=5 fusion(s) with PHD domains  
(1 pass filters)

# rve.HG5



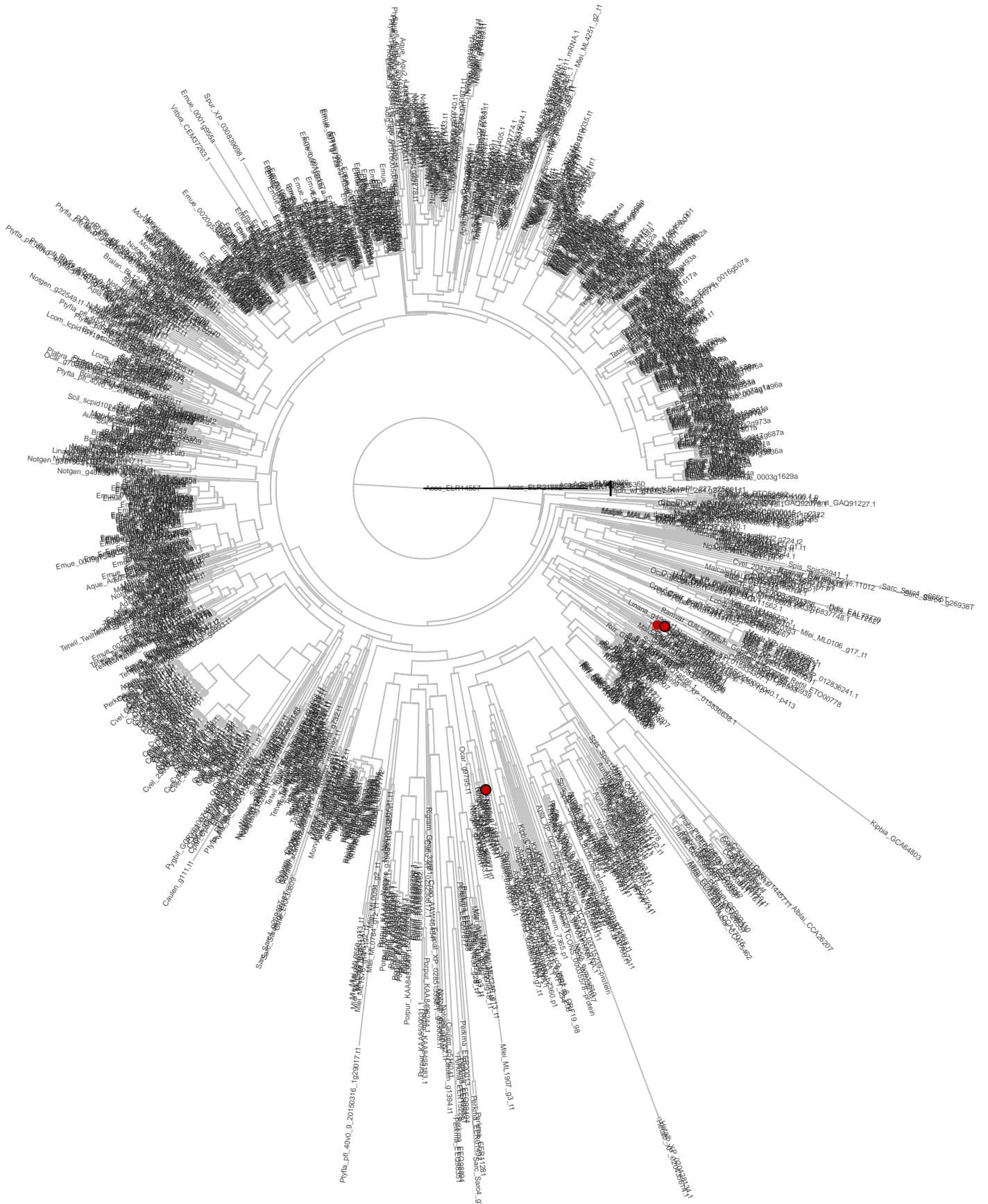
# rve.HG8



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

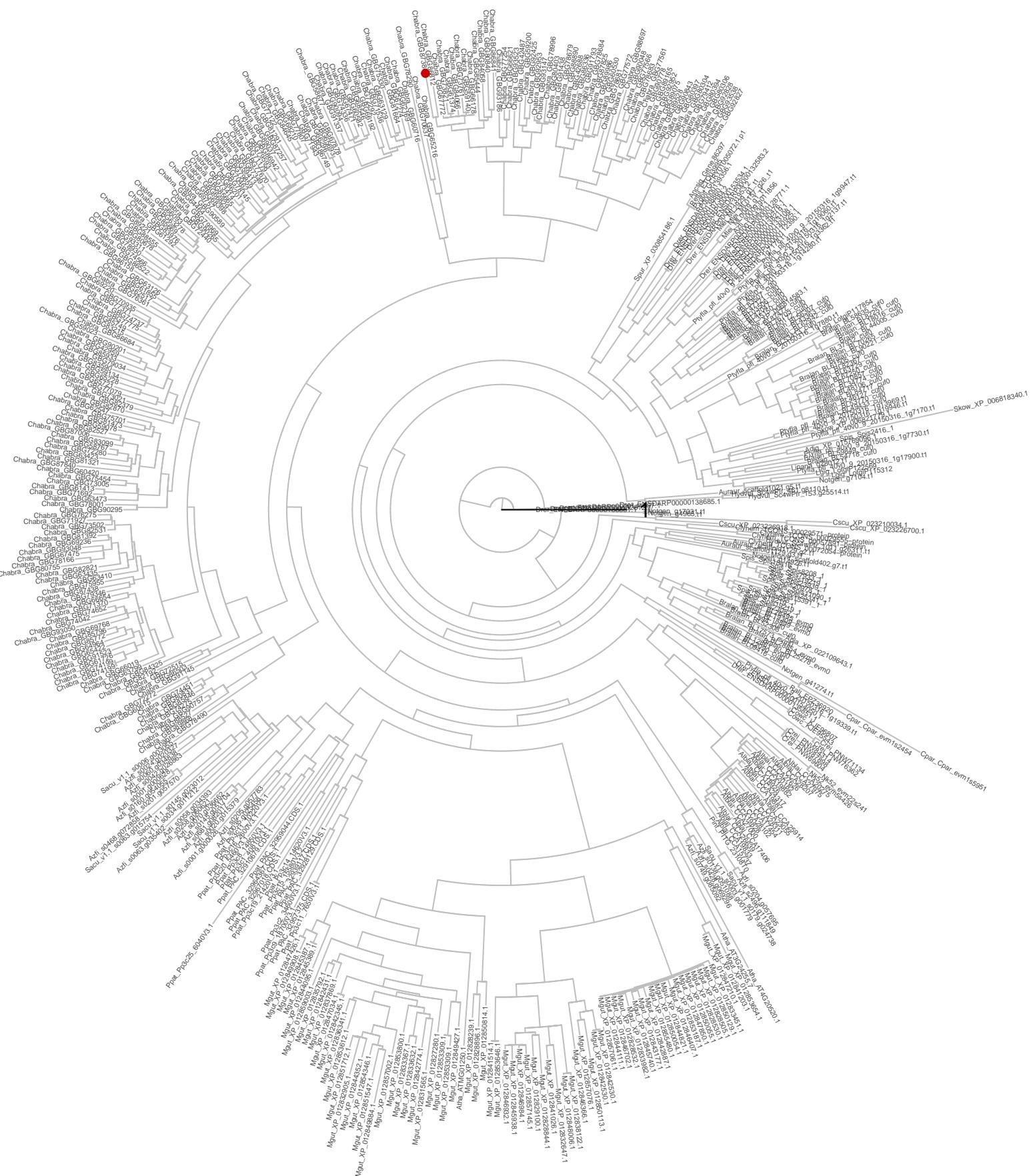
● PHD.HG23.1:NA n = 1

RVT.HG1



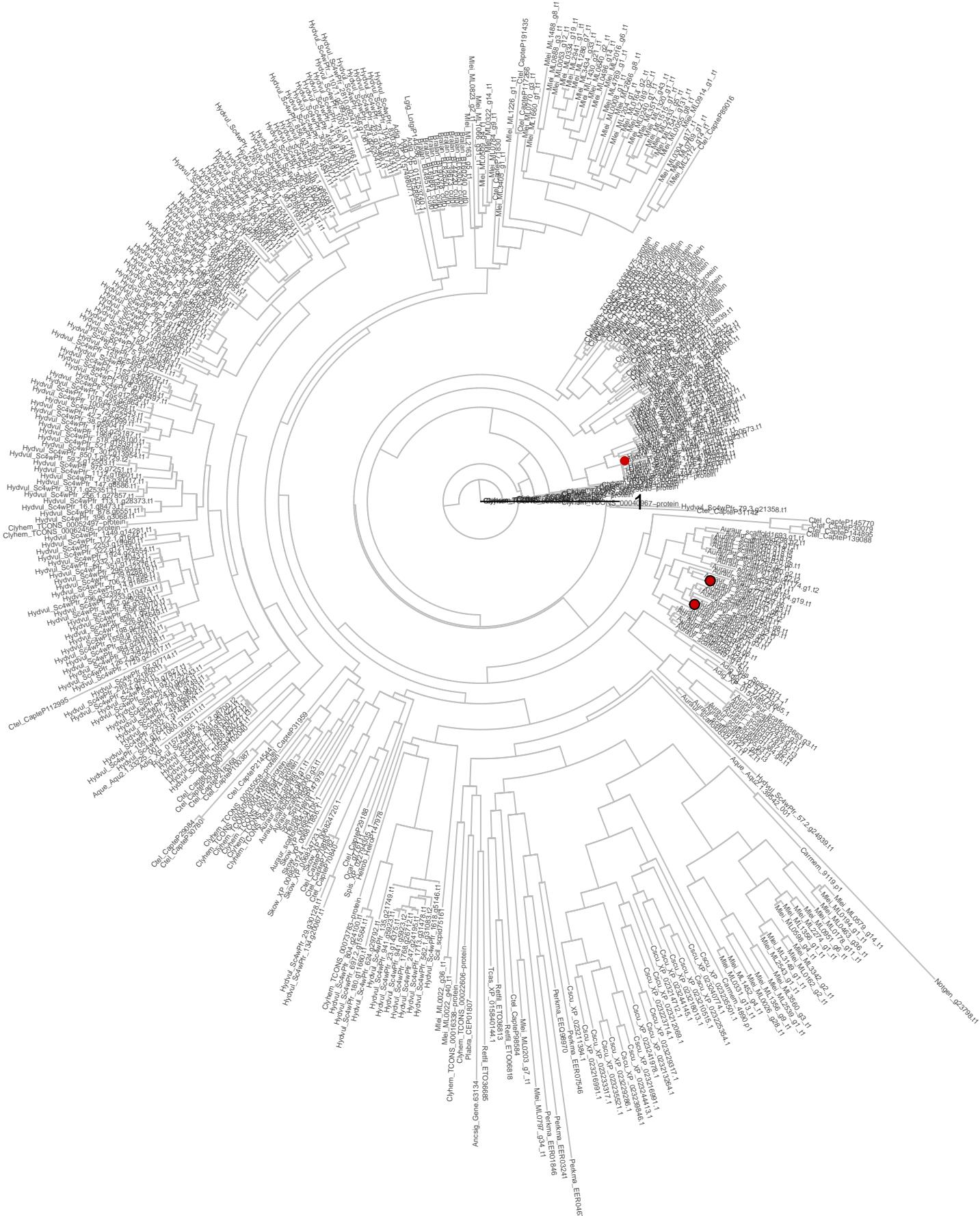
1401 genes  
n=3 fusion(s) with PHD domains  
(2 pass filters)

RVT.HG11



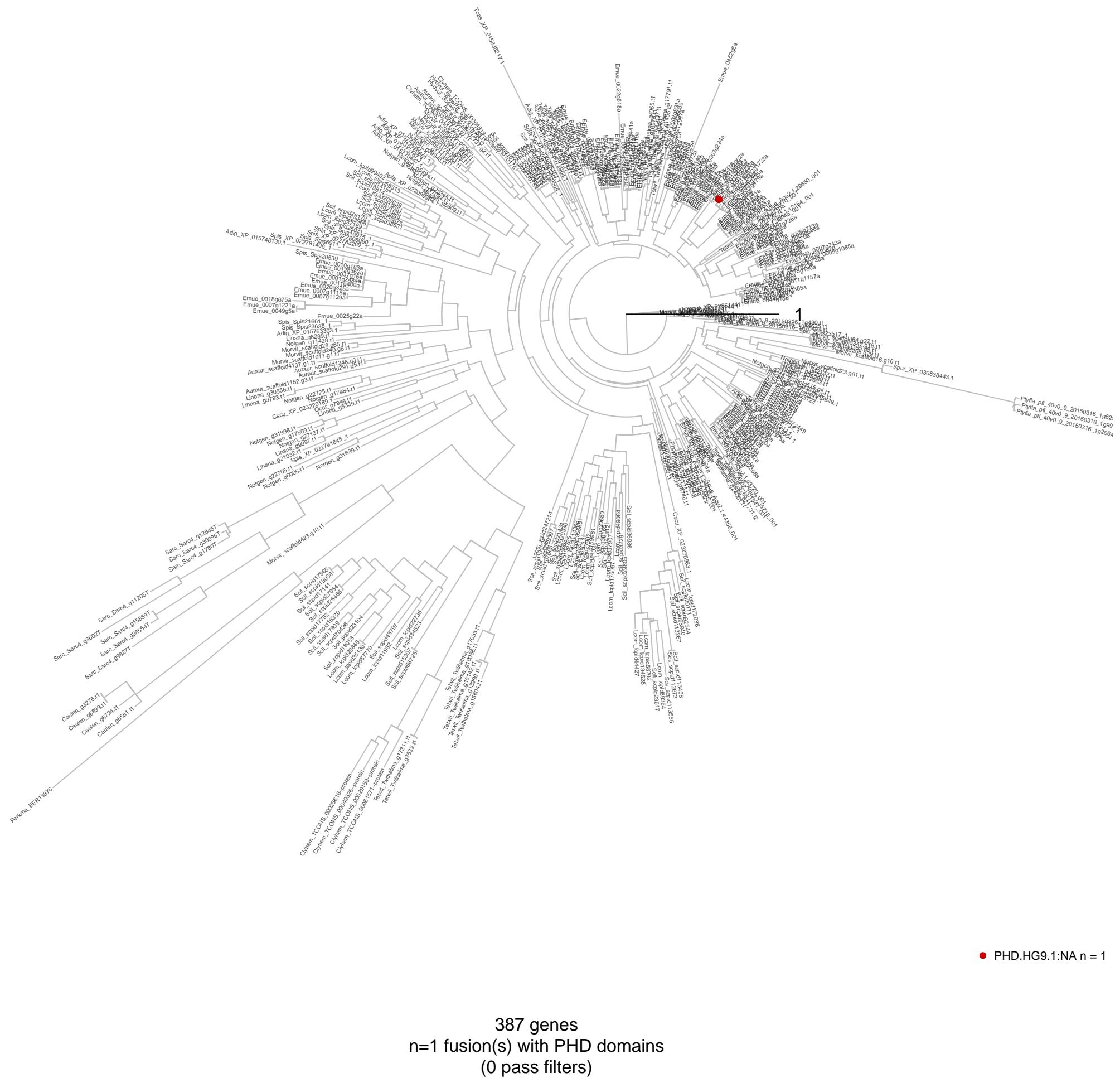
490 genes  
n=1 fusion(s) with PHD domains  
(0 pass filters)

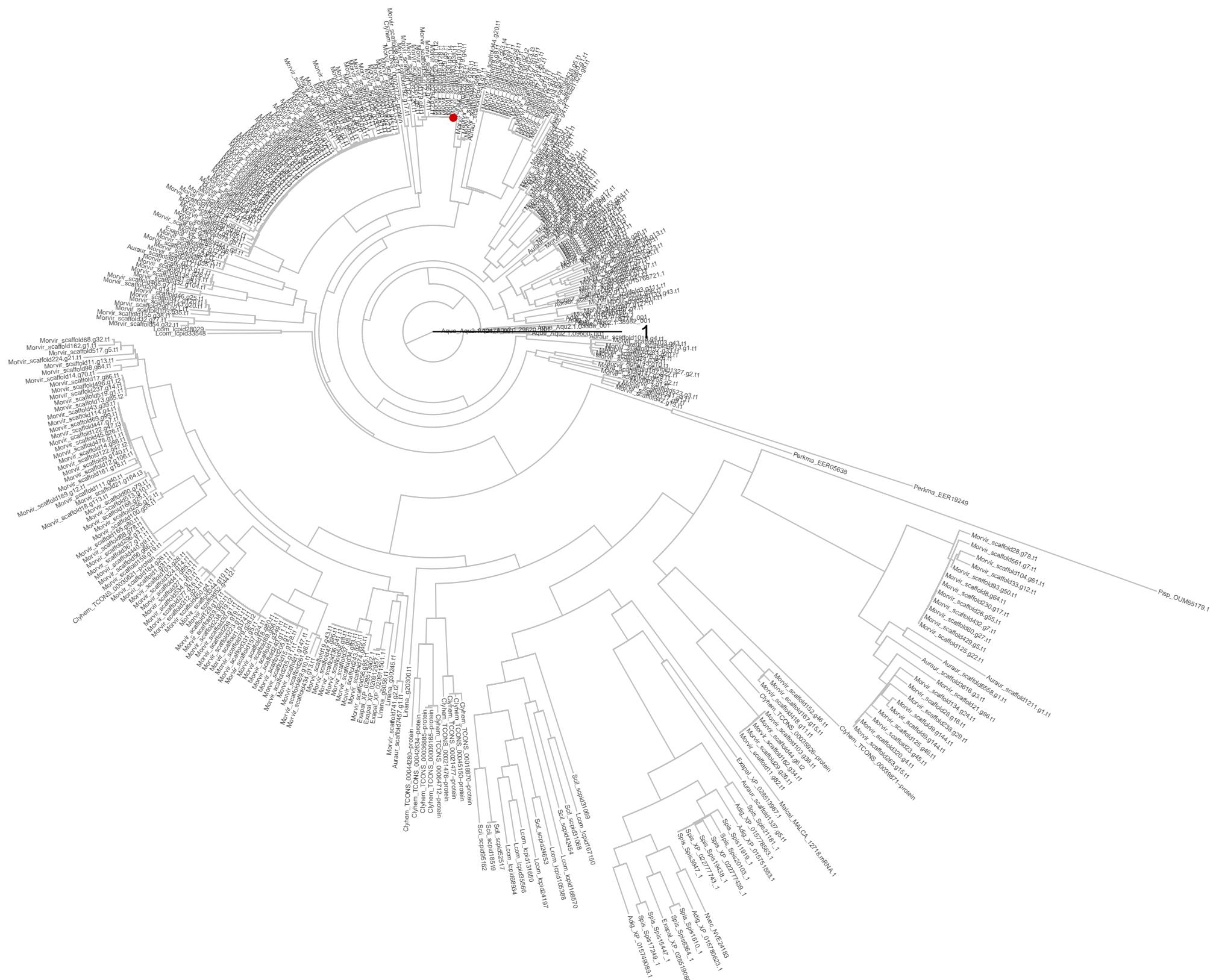
RVT.HG14



434 genes  
n=3 fusion(s) with PHD domains  
(2 pass filters)

# RVT.HG18

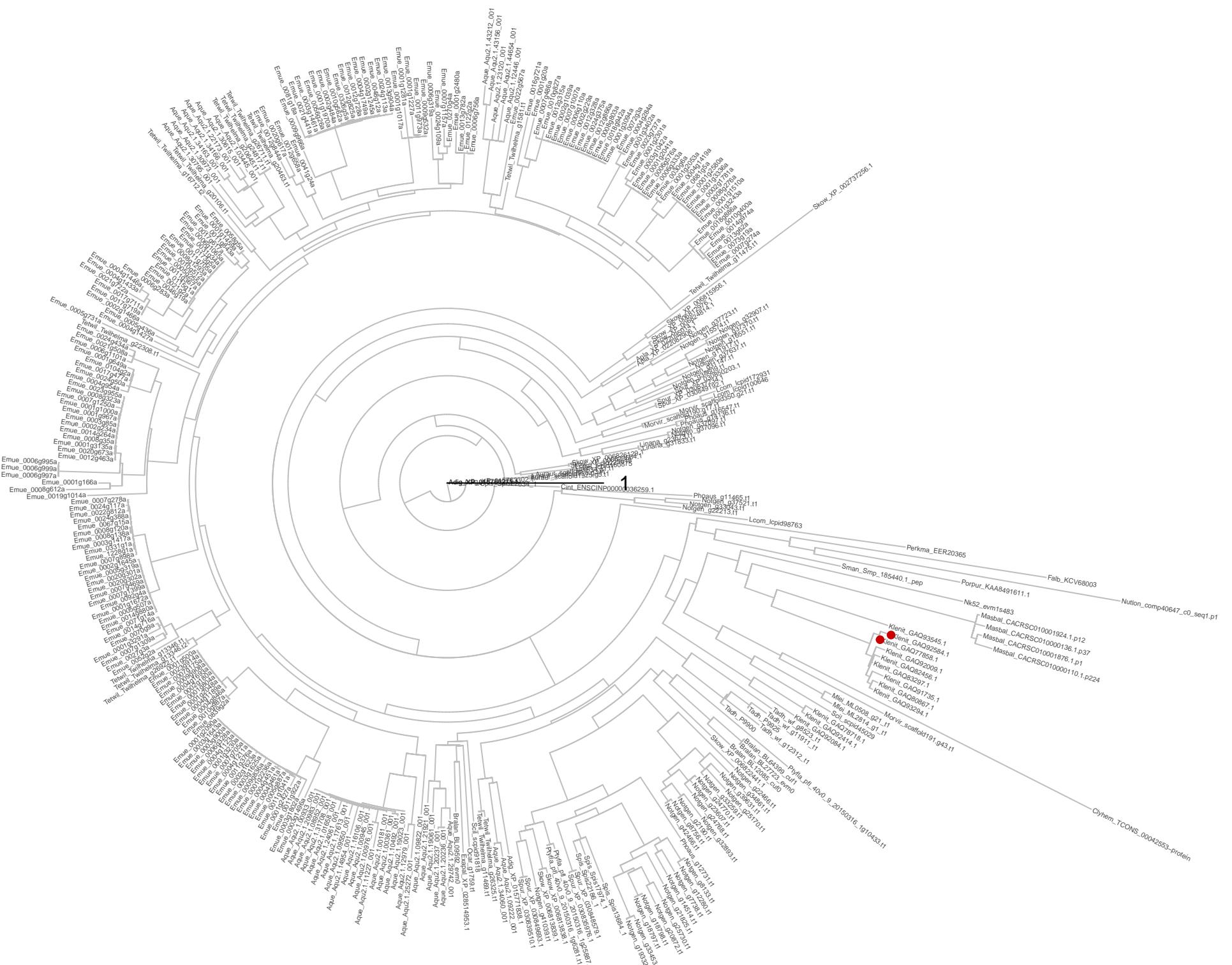




● PHD.HG11.1:like:PYGO1/PYGO2:likeclu:2 n = 1

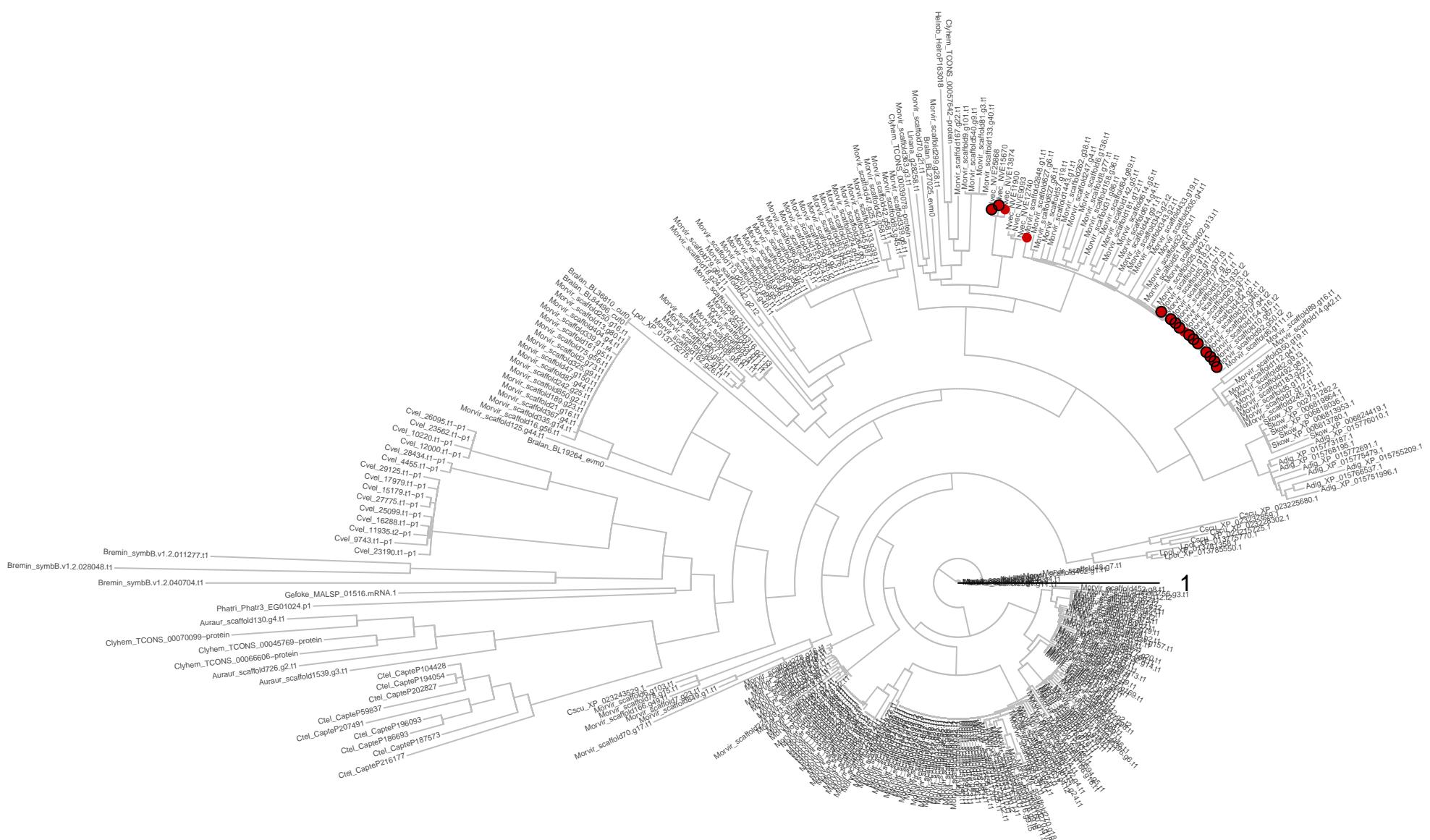
383 genes  
n=1 fusion(s) with PHD domains  
(0 pass filters)

RVT.HG24

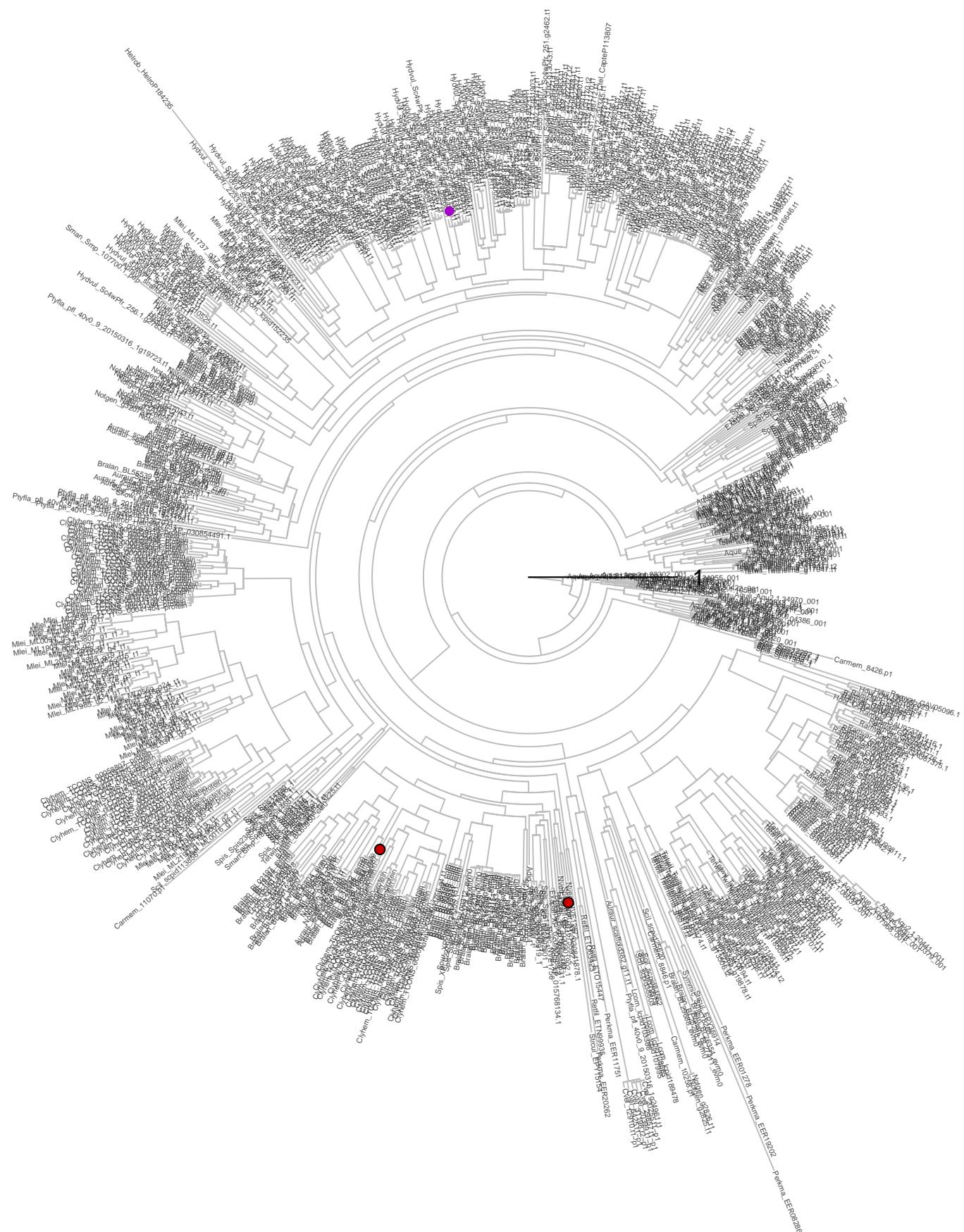


362 genes  
n=2 fusion(s) with PHD domains  
(0 pass filters)

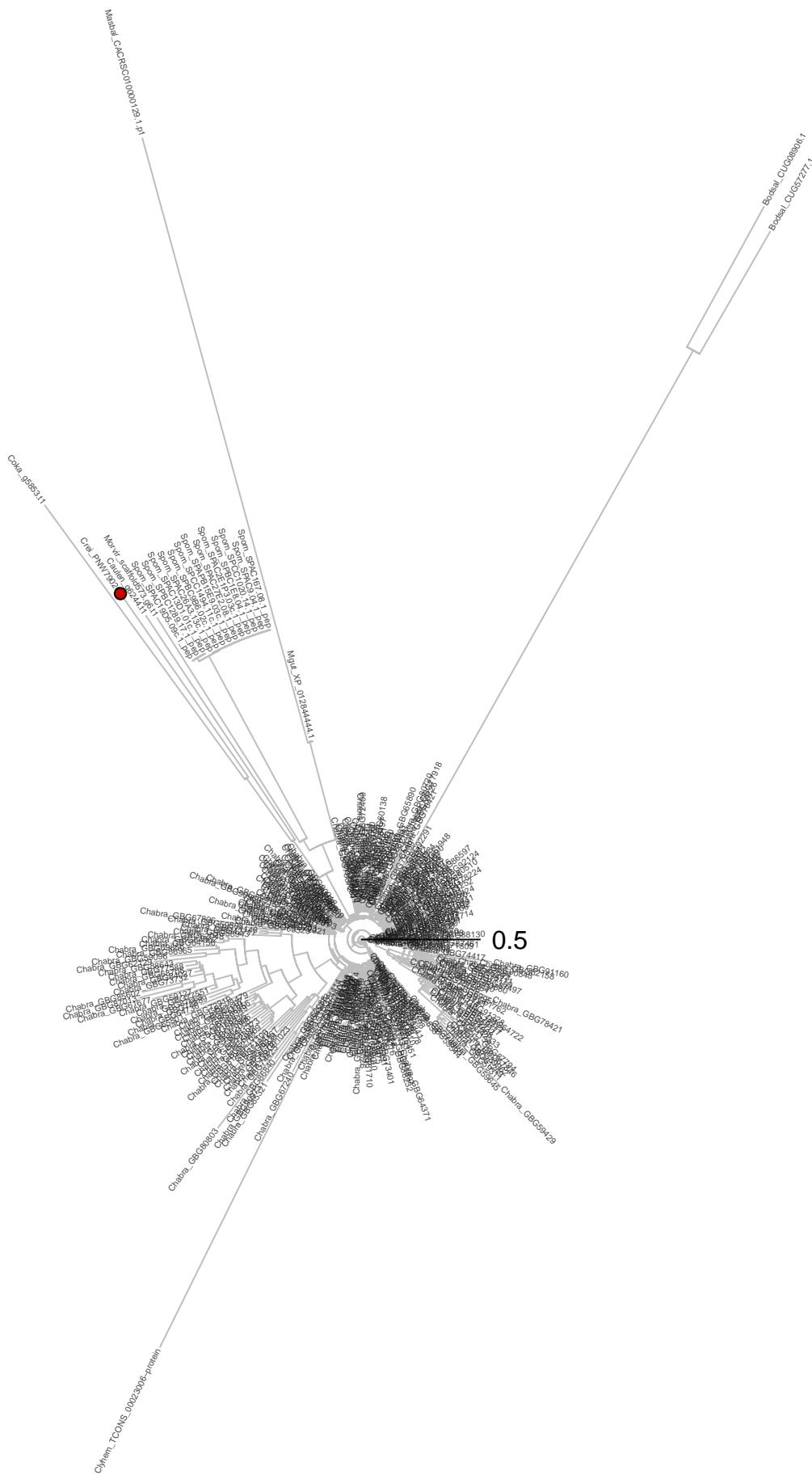
# RVT.HG25



# RVT.HG3

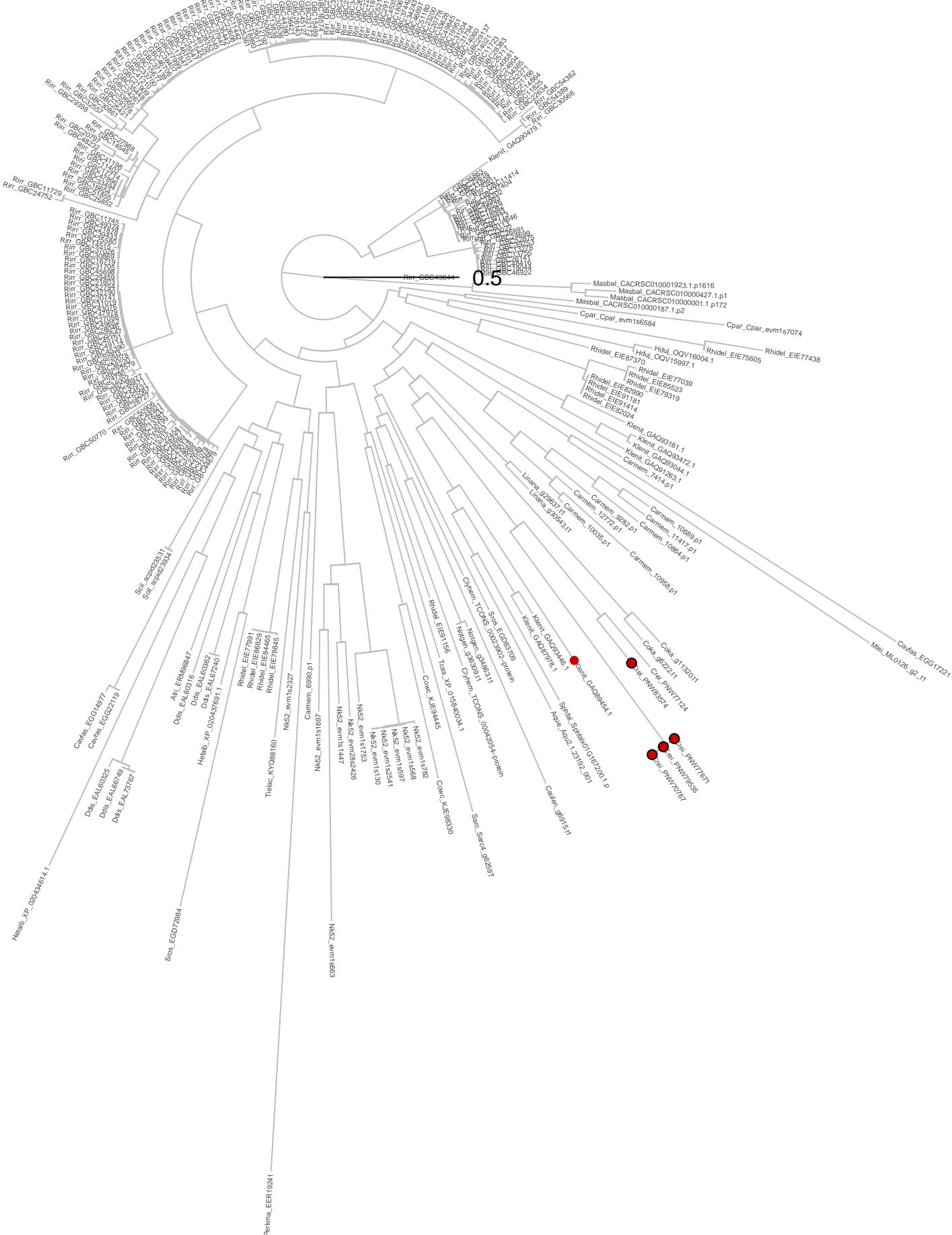


RVT.HG30



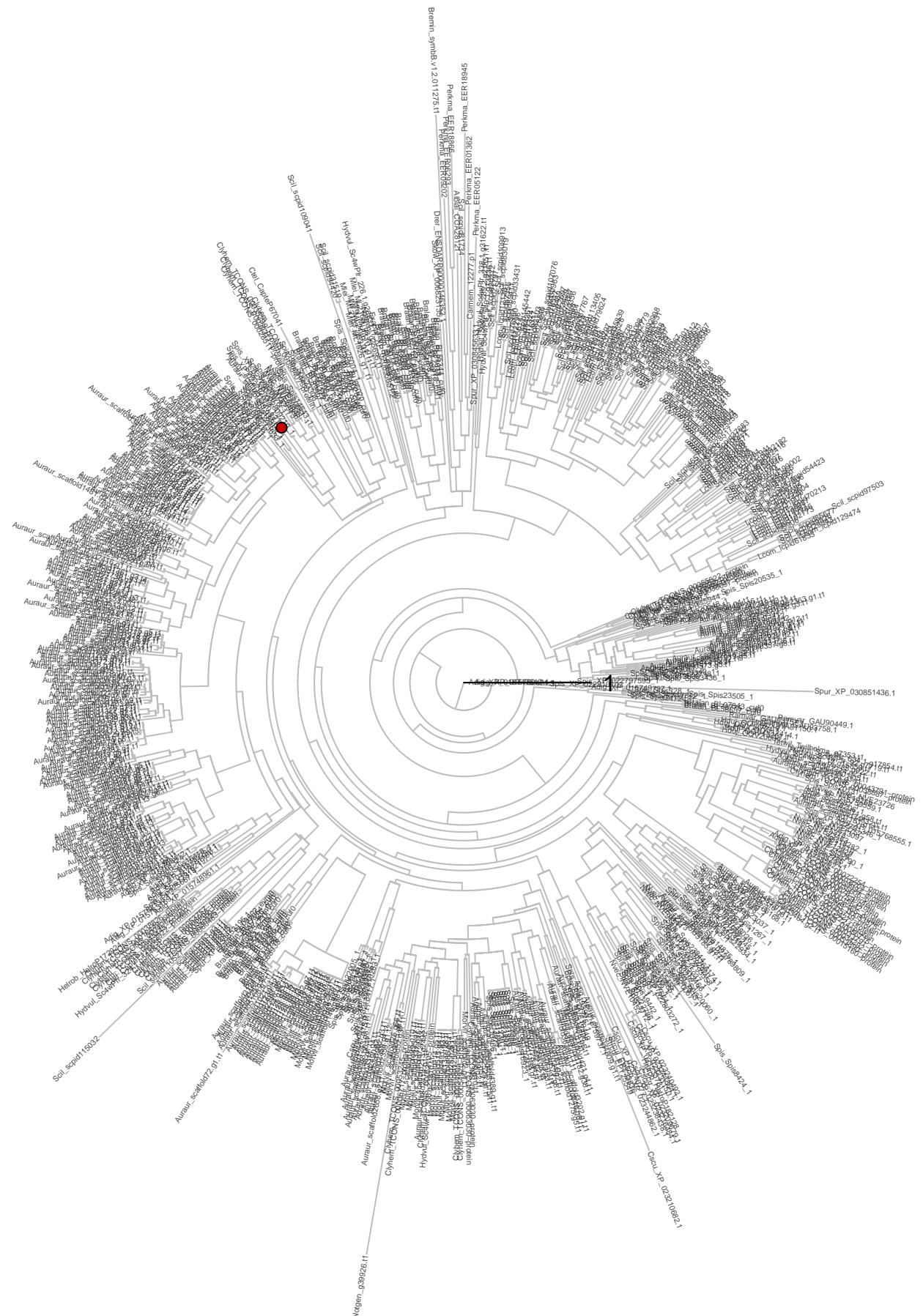
294 genes  
n=1 fusion(s) with PHD domains  
(1 pass filters)

RVT.HG33



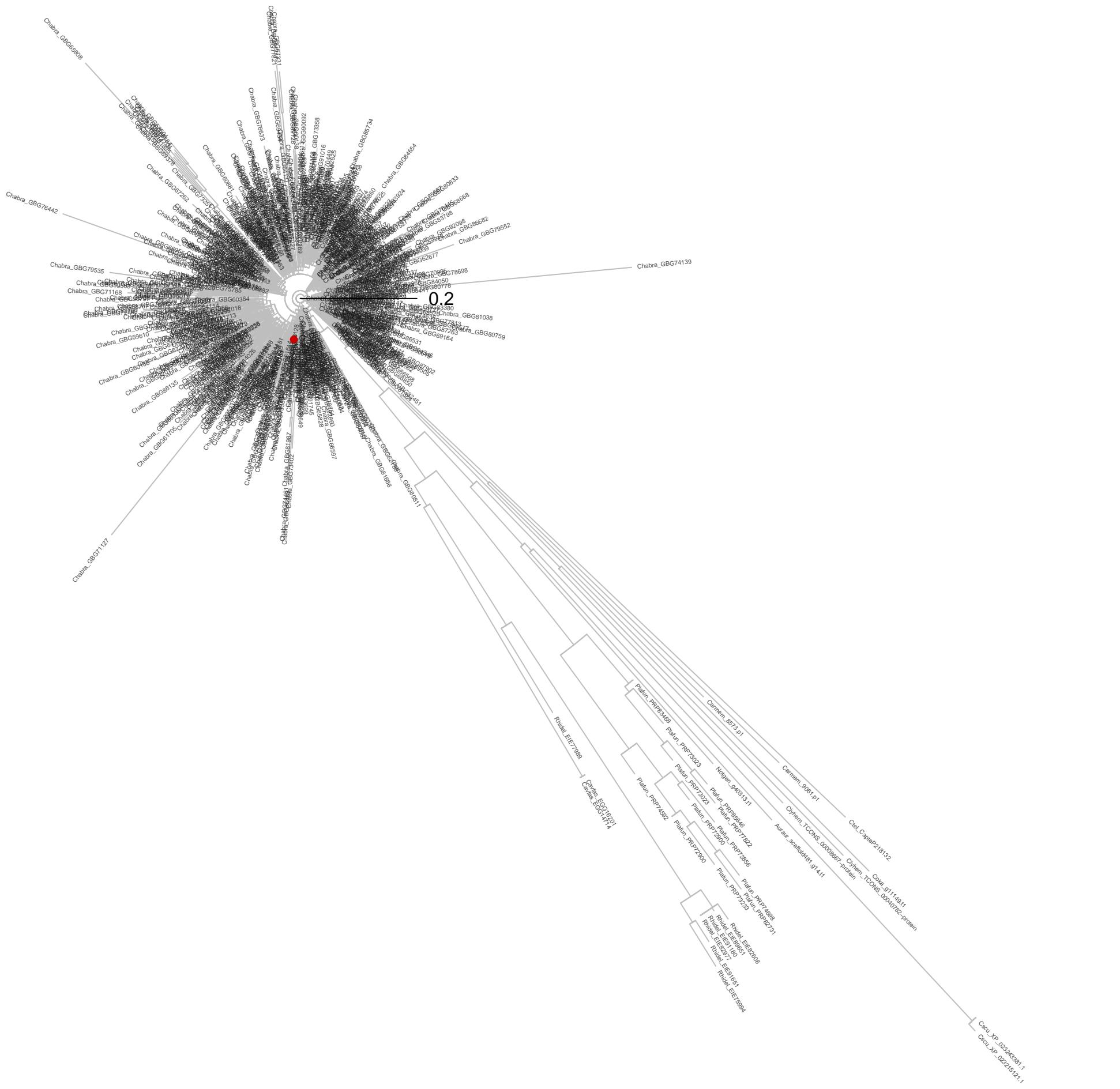
266 genes  
n=5 fusion(s) with PHD domains  
(4 pass filters)

RVT.HG5



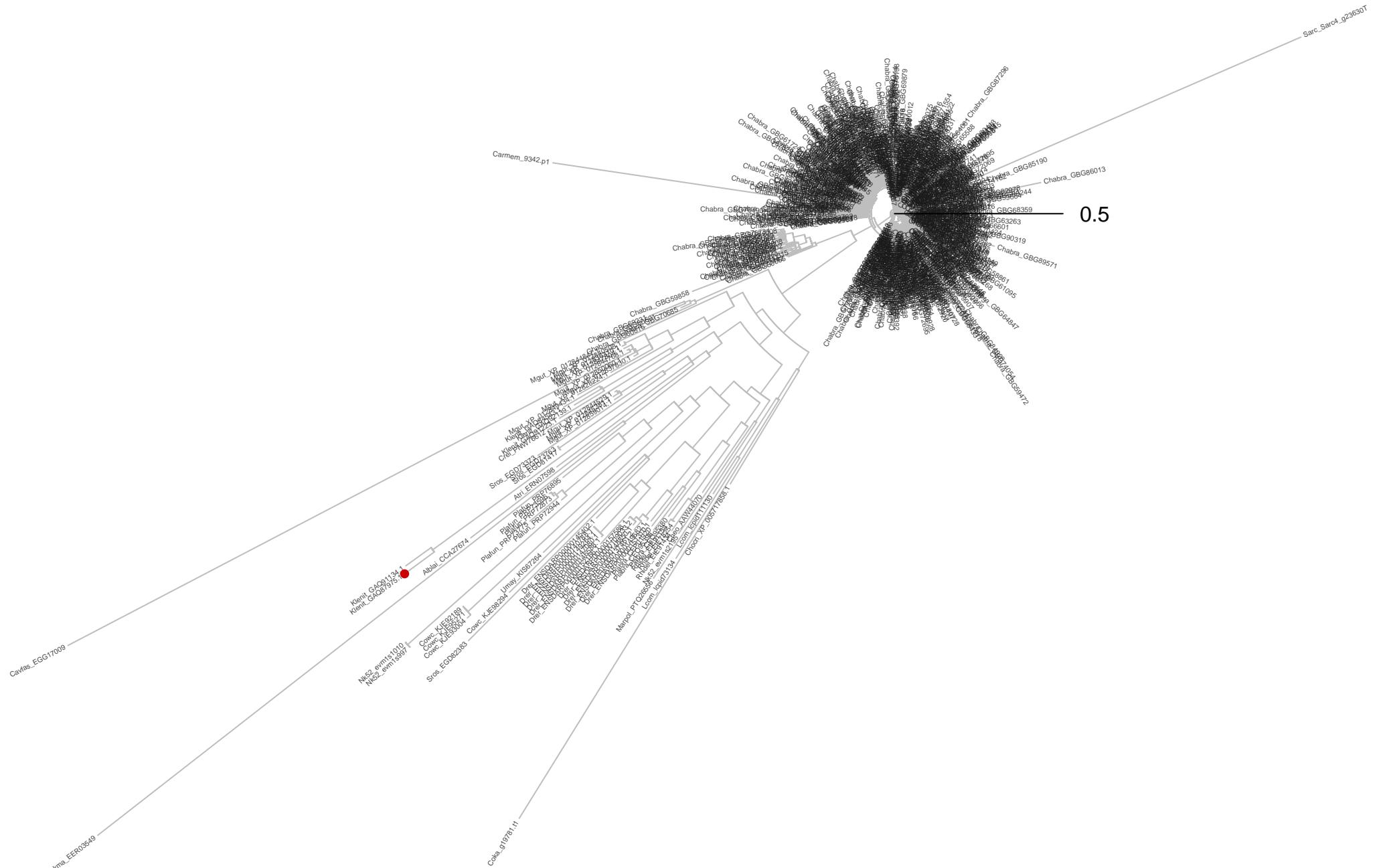
737 genes  
n=1 fusion(s) with PHD domains  
(1 pass filters)

RVT.HG6



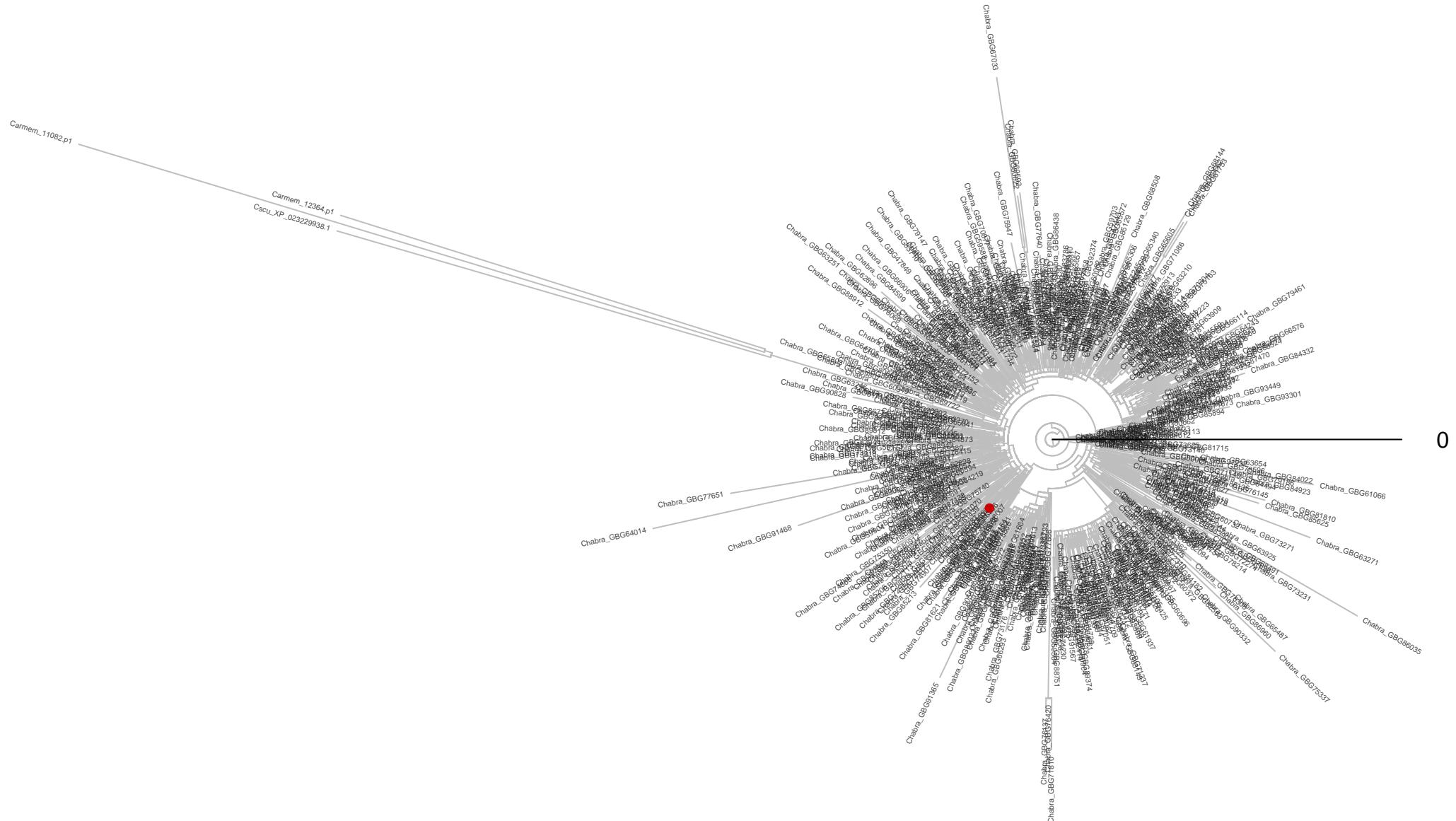
669 genes  
n=1 fusion(s) with PHD domains  
(0 pass filters)

RVT.HG7



659 genes  
n=1 fusion(s) with PHD domains  
(0 pass filters)

RVT.HG8



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