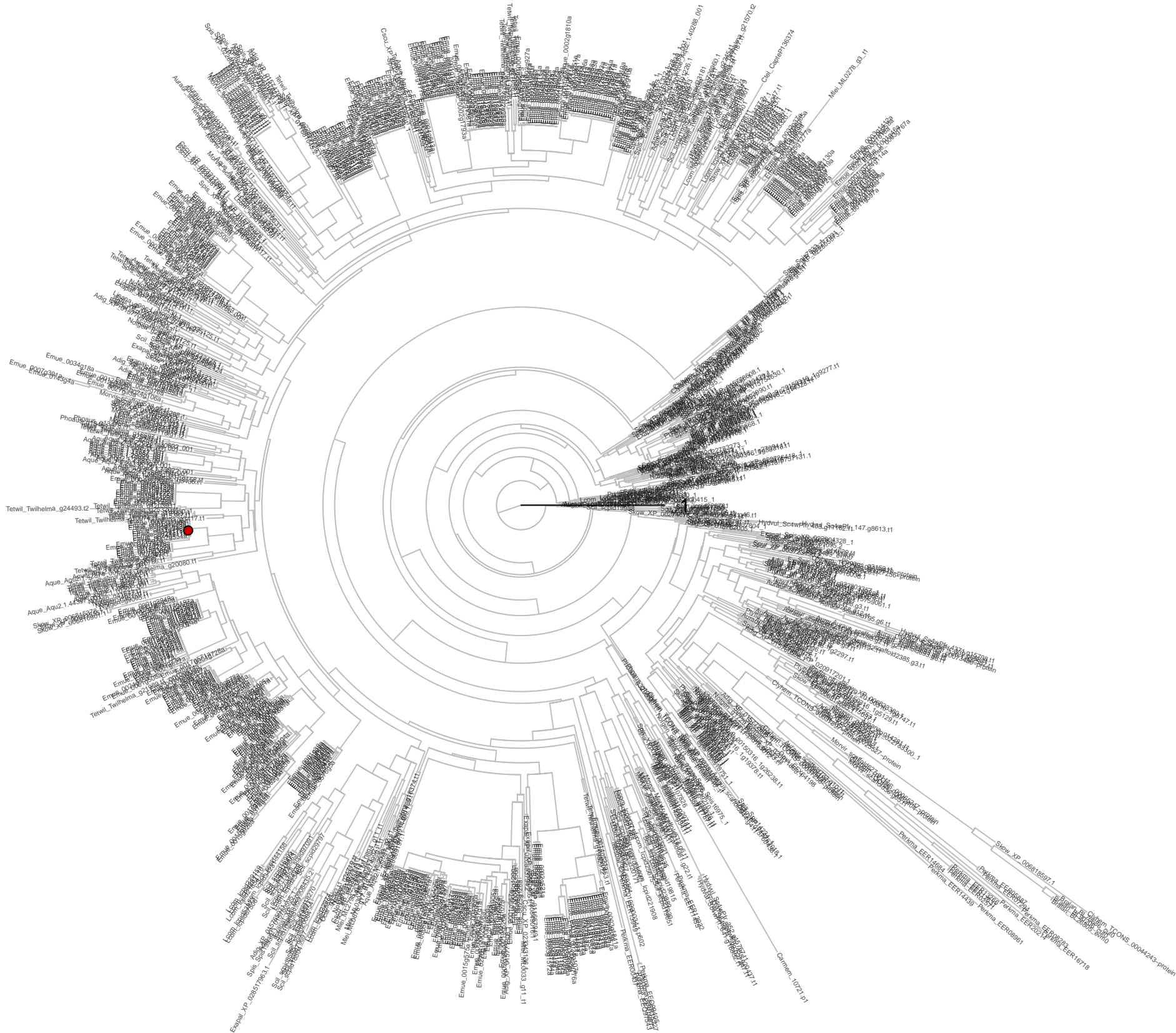


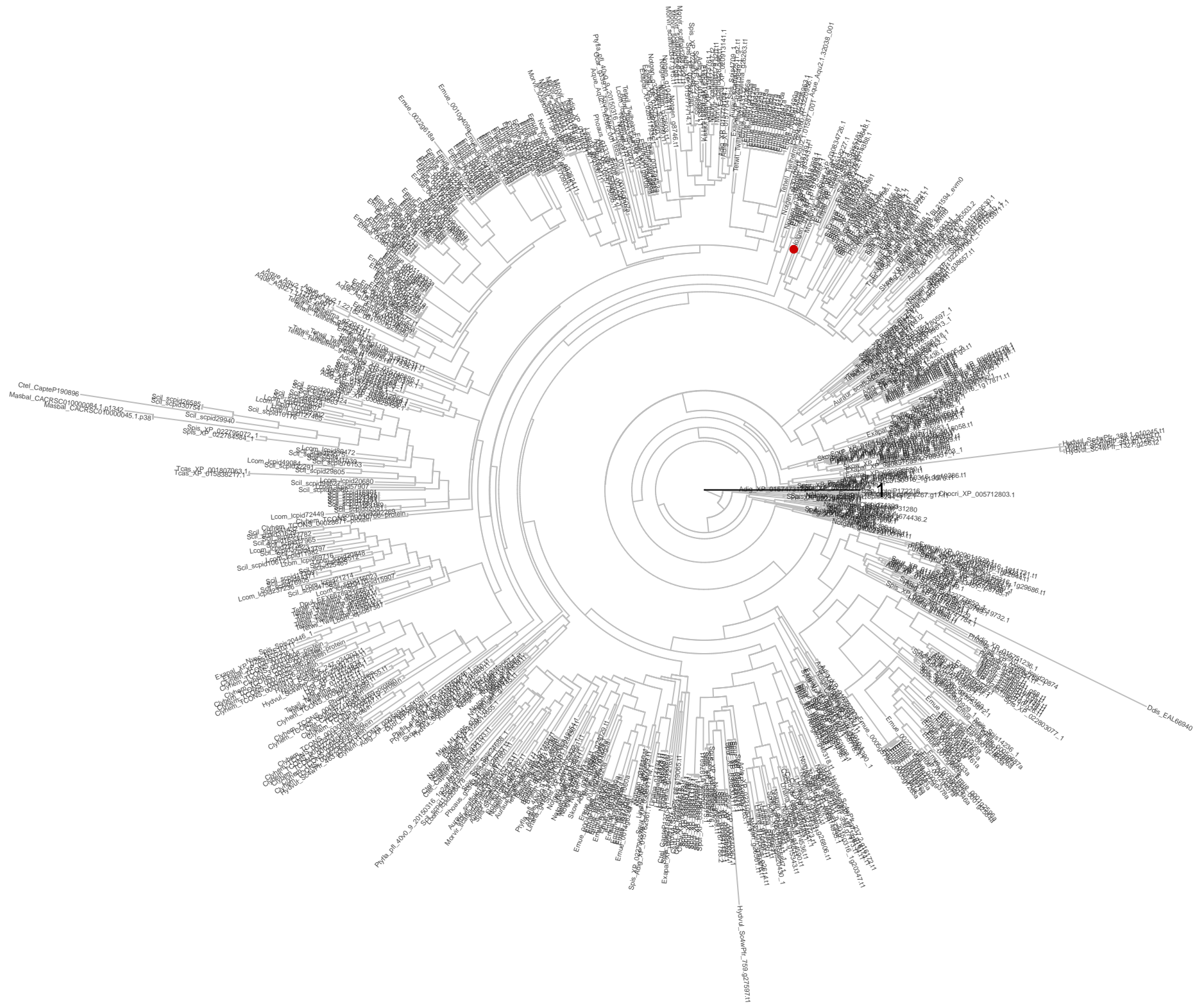
Integrase_H2C2.HG2



● SAM.HG3.14:like:ANKS1A/ANKS1B/CASKIN1... n = 1

1119 genes
n=1 fusion(s) with SAM domains
(1 pass filters)

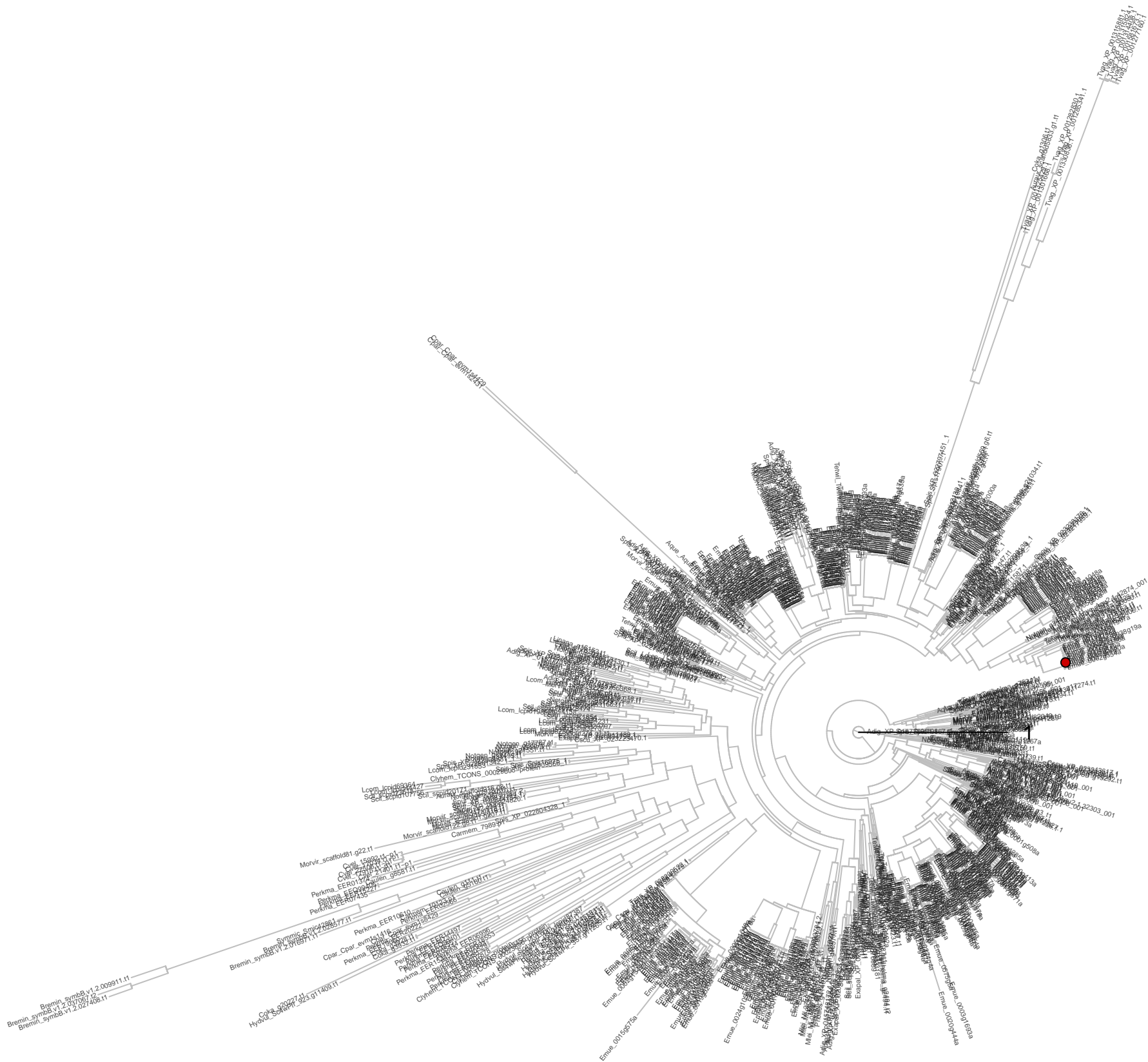
Integrase_H2C2.HG4



- SAM.HG3.1:like:ANKS1A/ANKS1B/CASKIN1/... n = 1

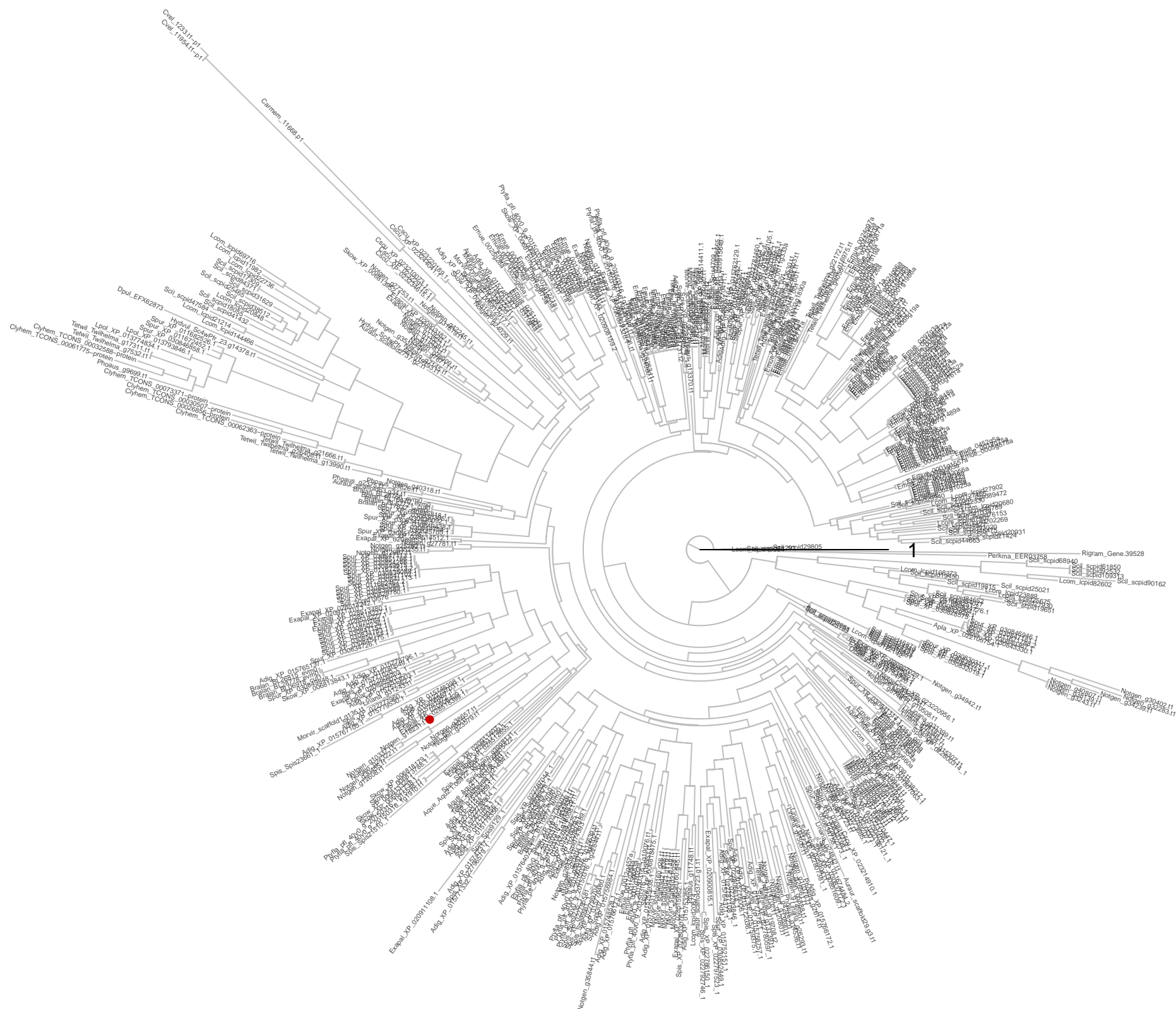
786 genes
n=1 fusion(s) with SAM domains
(0 pass filters)

rve.HG1



● SAM.HG3.14:like:ANKS1A/ANKS1B/CASKIN1... n = 1

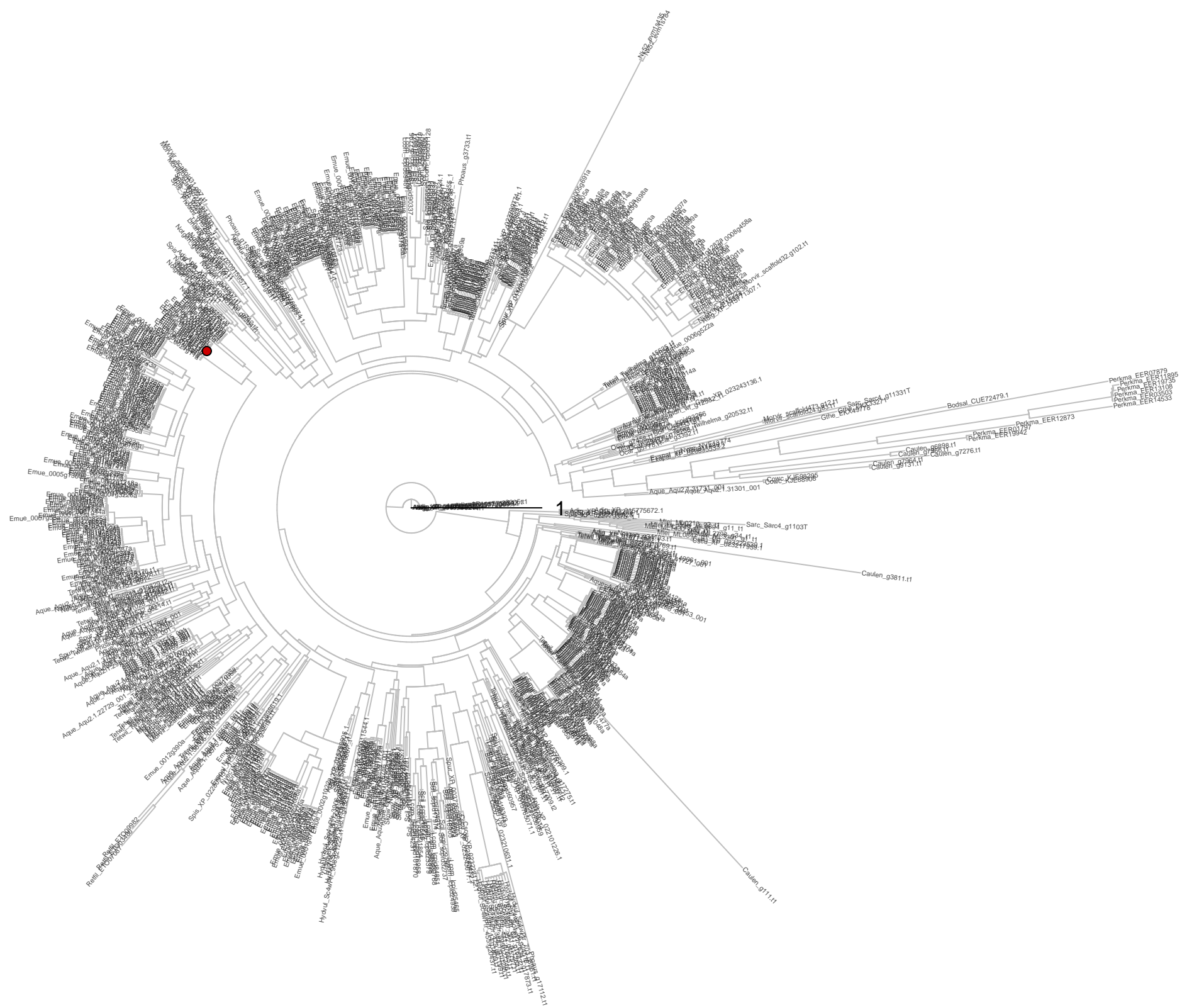
908 genes
n=1 fusion(s) with SAM domains
(1 pass filters)

rve.HG3

- SAM.HG3.1:like:ANKS1A/ANKS1B/CASKIN1/... n = 1

582 genes
n=1 fusion(s) with SAM domains
(0 pass filters)

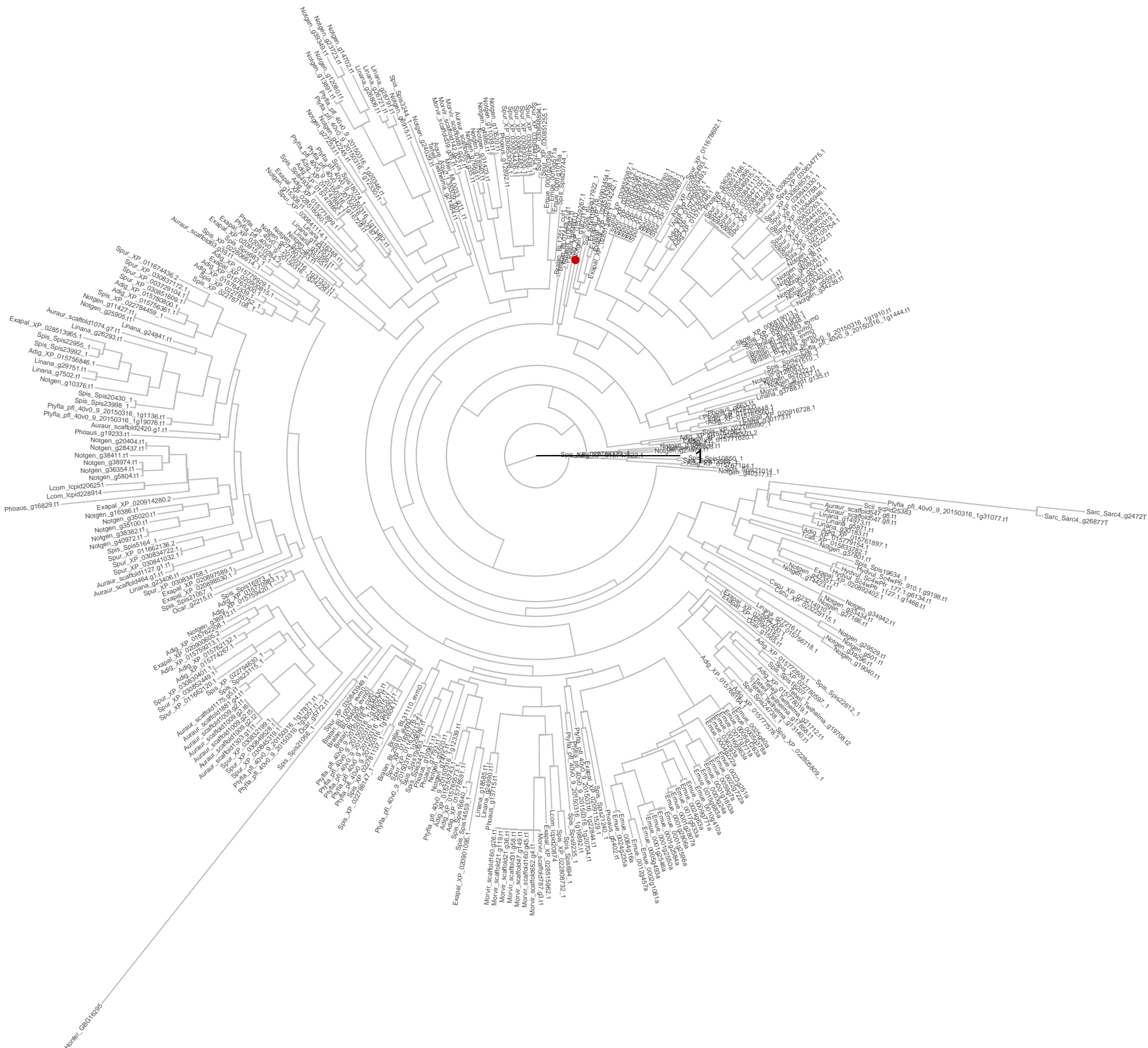
RVT.HG2



● SAM.HG3.14:like:ANKS1A/ANKS1B/CASKIN1... n = 1

874 genes
n=1 fusion(s) with SAM domains
(1 pass filters)

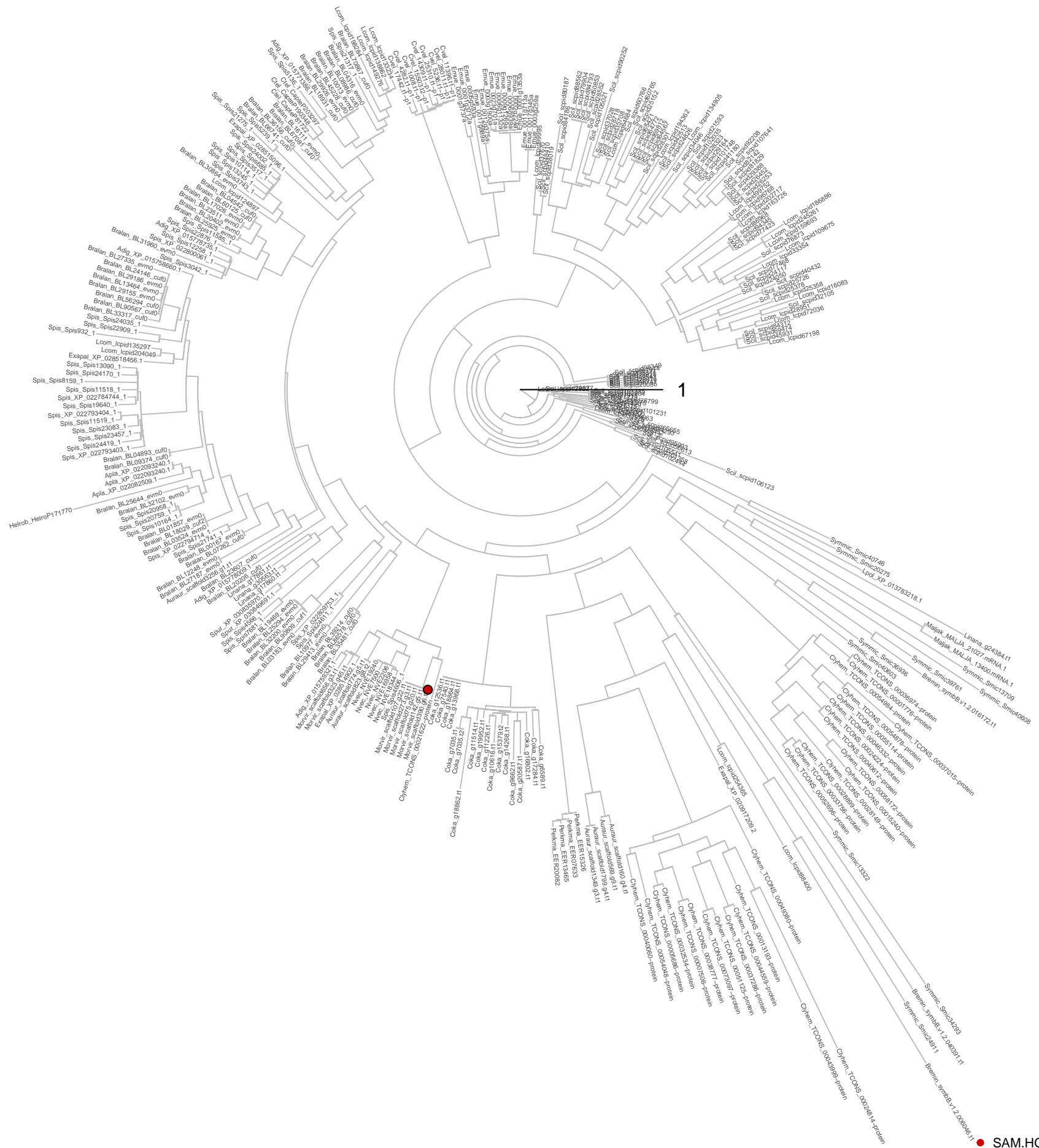
RVT.HG23



● SAM.HG3.1:like:ANKS1A/ANKS1B/CASKIN1/... n = 1

370 genes
n=1 fusion(s) with SAM domains
(0 pass filters)

RVT.HG27



336 genes
n=1 fusion(s) with SAM domains
(1 pass filters)