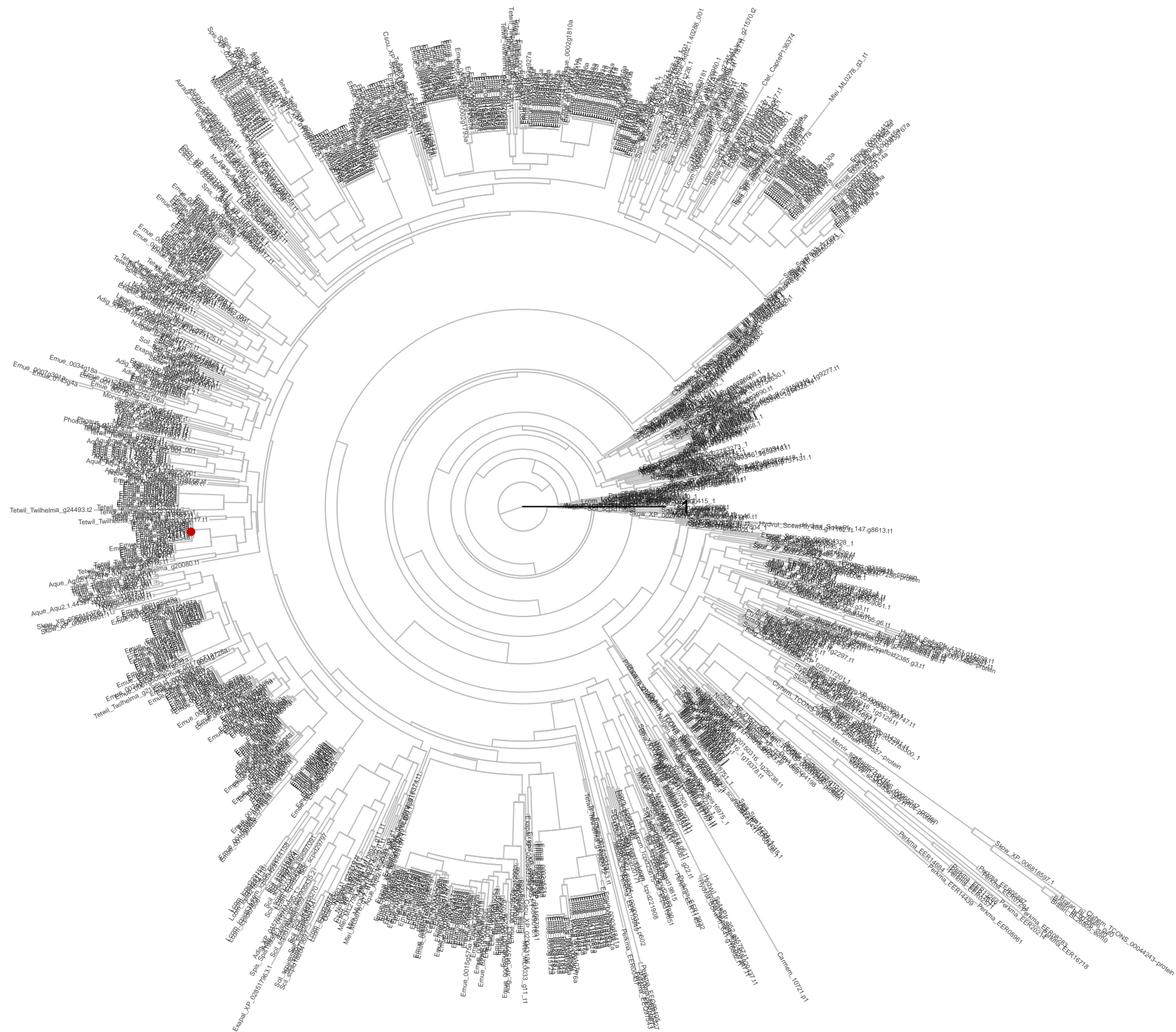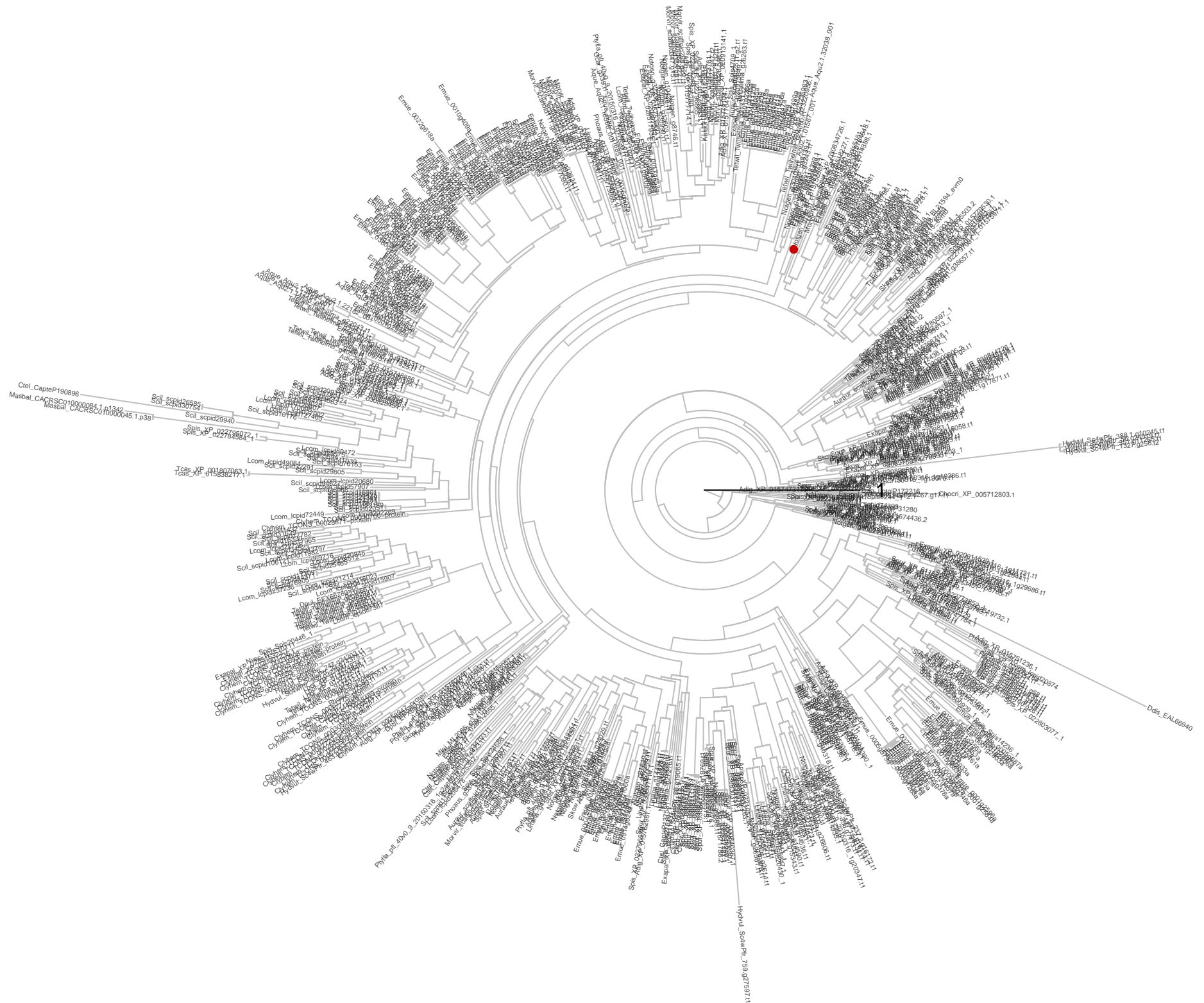# Integrase_H2C2.HG2

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

1119 genes
n=1 fusion(s) with SAM domains
(0 pass filters)
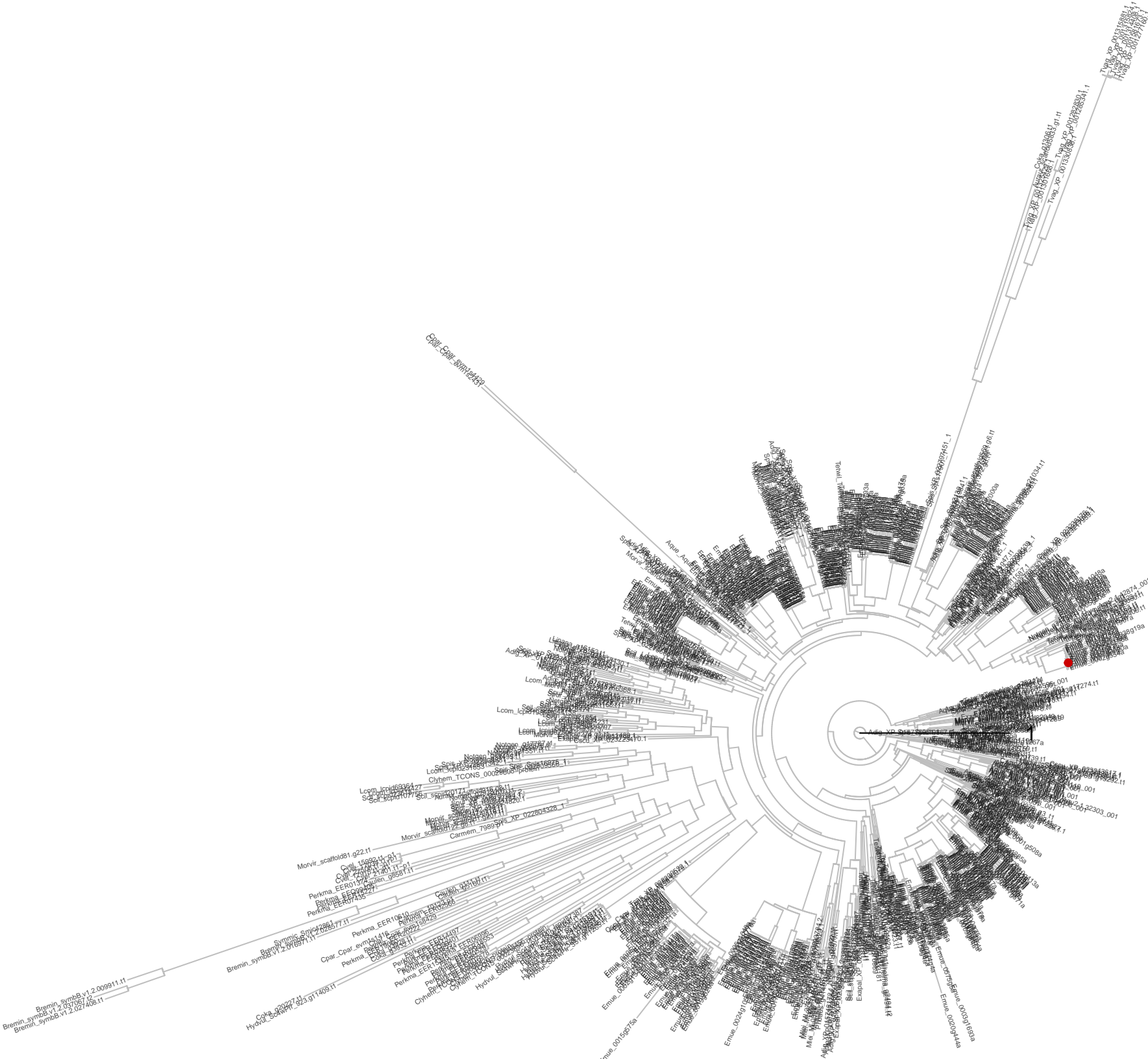
Integrase_H2C2.HG4
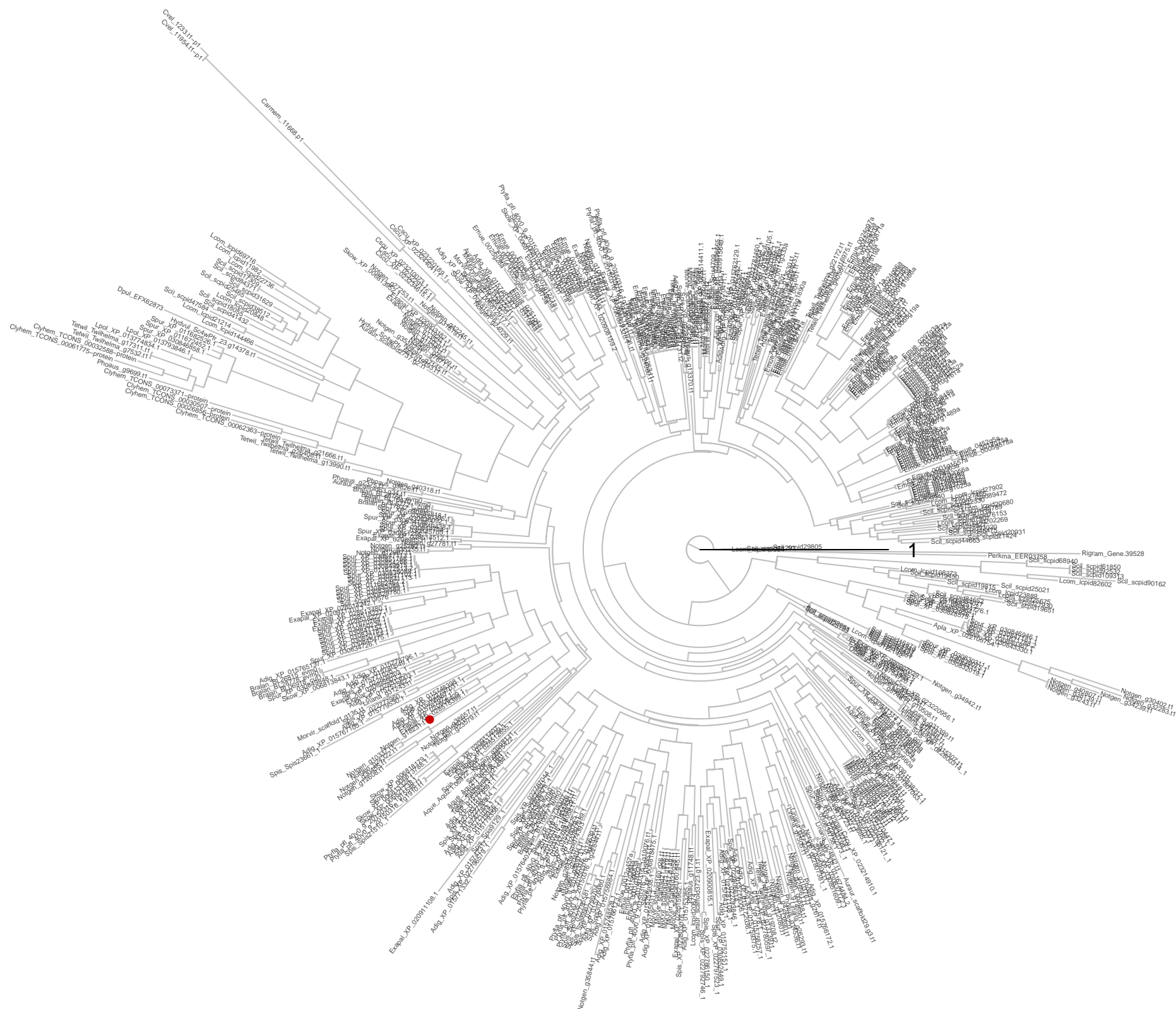

● SAM.HG3.1:like:ANKS1A/ANKS1B/CASKIN1/... n = 1

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

rve.HG1

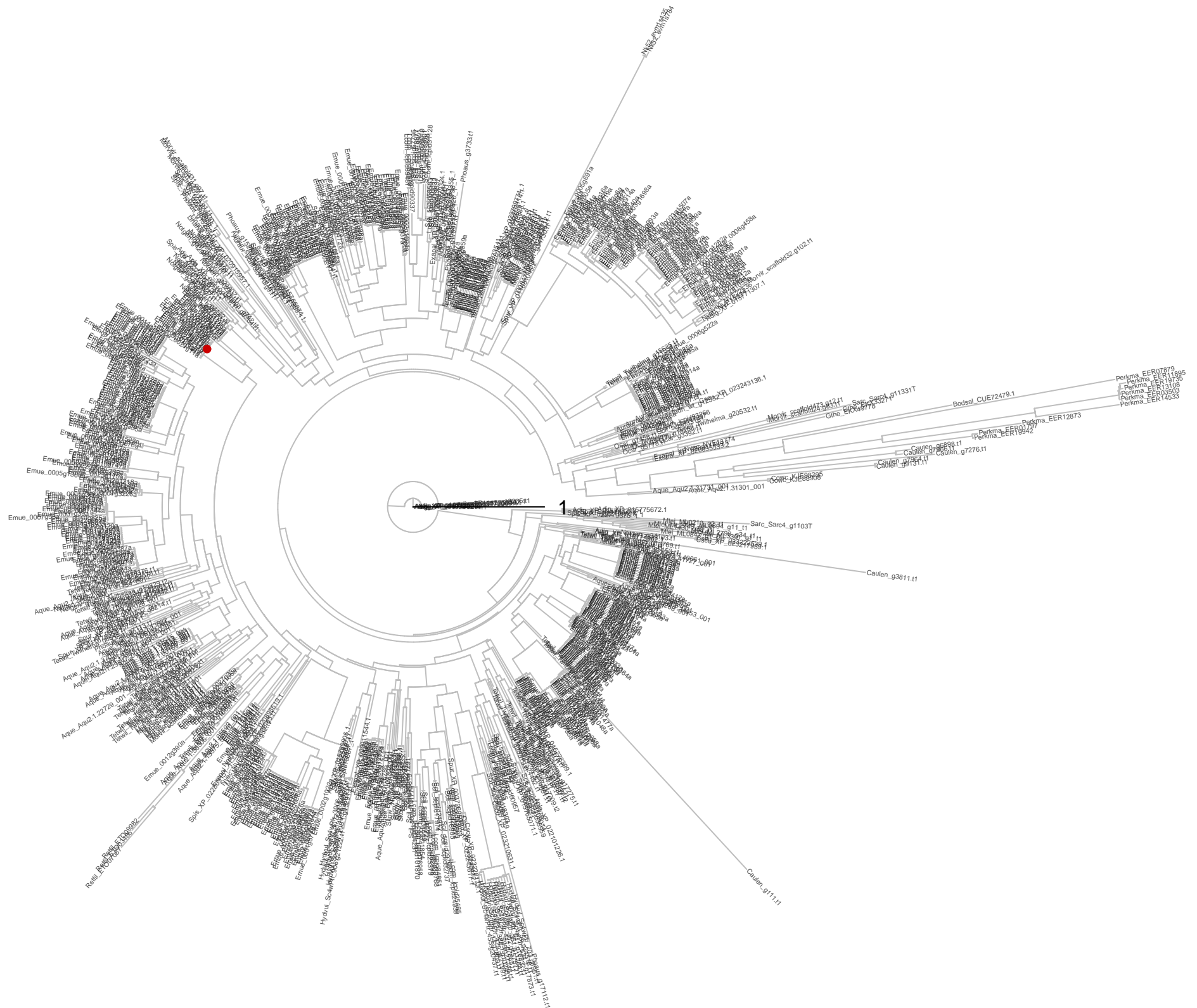● SAM.HG3.14:like:ANKS1A/ANKS1B/CASKIN1... n = 1

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

rve.HG3

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

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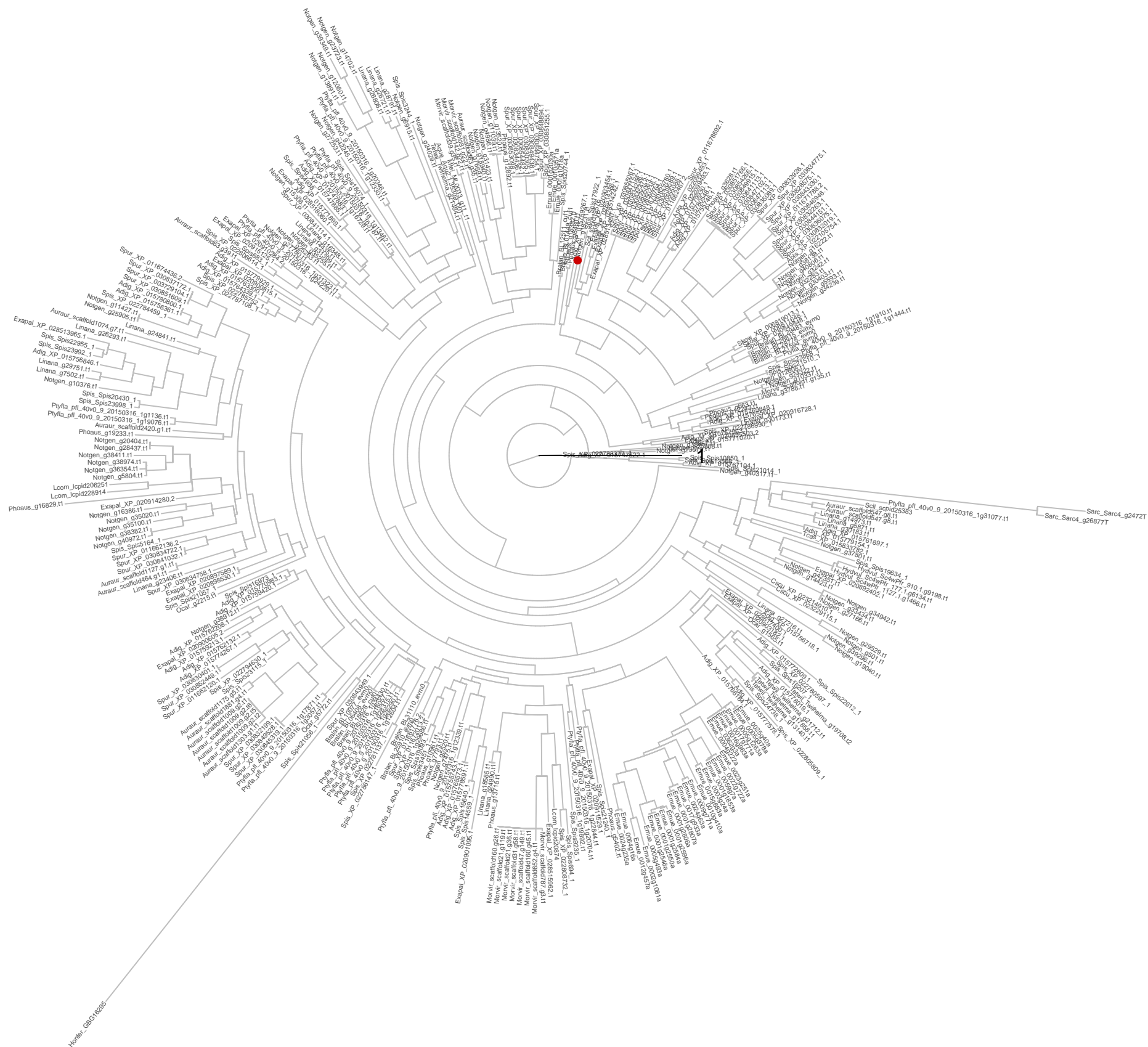n=1 fusion(s) with SAM domains
(0 pass filters)

RVT.HG2

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

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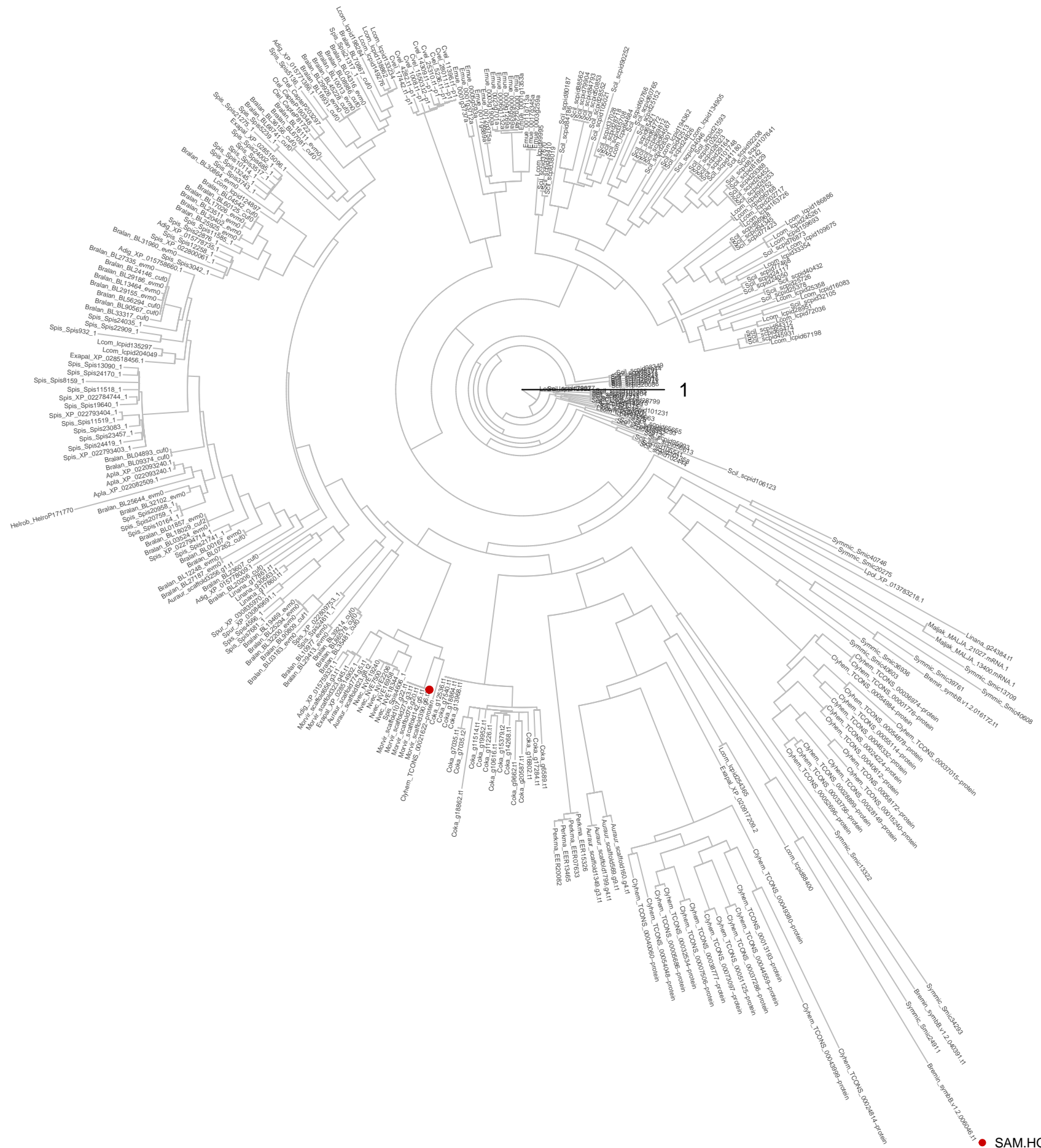n=1 fusion(s) with SAM domains
(0 pass filters)

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

● SAM.HG1.3:SAMHD1 n = 1

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