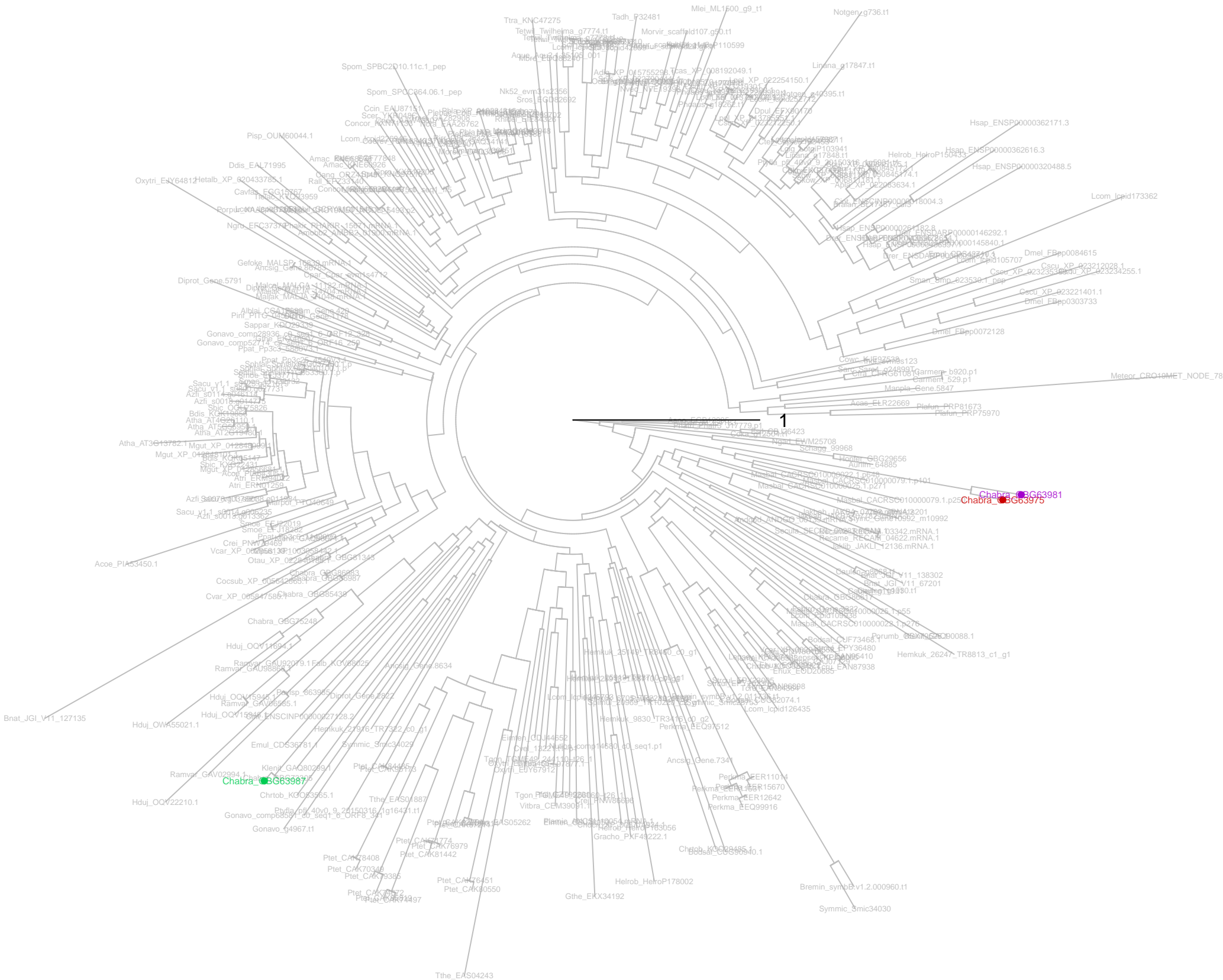


NAP.HG1



- Integrase_H2C2.HG11.0,ve.HG13.2 n = 1
- Integrase_H2C2.HG6.0 n = 1
- Integrase_H2C2.HG8.1 n = 1

342 genes
n=3 fusion(s) with RT domains
(0 pass filters)