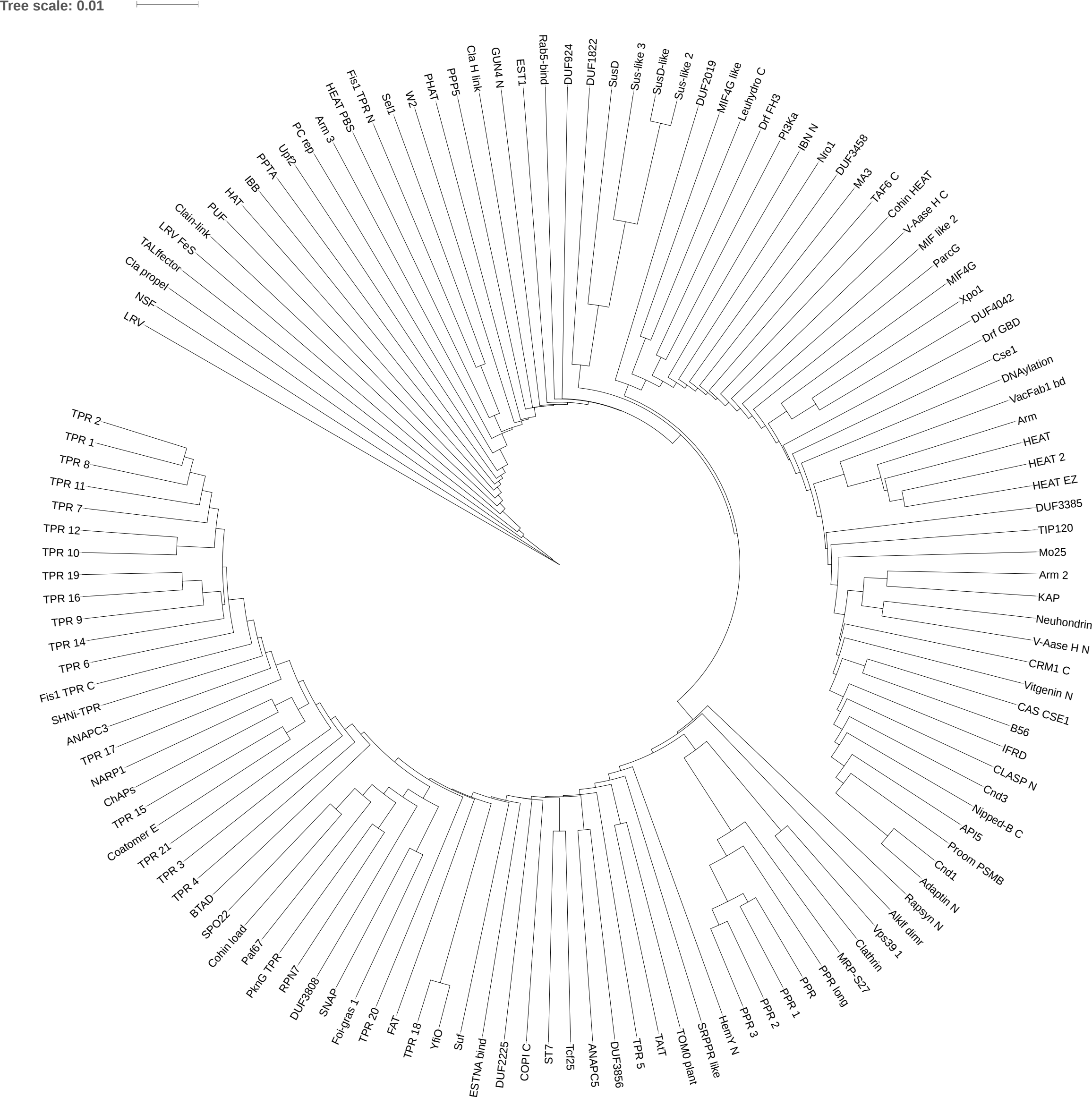
Tree scale: 0.01
Rab5-bind
DUF924
DUF1822
SusD
Sus-like 3
SusD-like
Sus-like 2
DUF2019
MIF4G like
Leuhydro C
Drf FH3
PI3Ka
IBN N
Nro1
DUF3458
MA3
TAF6 C
Cohin HEAT
V-Aase H C
MIF like 2
ParcG
MIF4G
Xpo1
DUF4042
Drf GBD
Cse1
DNAylation
VacFab1 bd
Arm
HEAT
HEAT 2
HEAT EZ
DUF3385
TIP120
Mo25
Arm 2
KAP
Neuhondrin
V-Aase H N
CRM1 C
Vitgenin N
CAS CSE1
B56
IFRD
CLASP N
Cnd3
Nipped-B C
API5
Proom PSMB
Cnd1
Adaptin N
Rapsyn N
Alkdf dimr
Vps39 1
Clathrin
MRP-S27
PPR long
PPR
PPR 1
PPR 2
PPR 3
HemY N
SRPPR like
TOM0 plant
TAtT
TPR 5
DUF3856
ANAPC5
Tcf25
ST7
COPI C
DUF2225
ESTNA bind
Suf
YfiO
TPR 18
FAT
TPR 20
Foi-gras 1
SNAP
DUF3808
RPN7
PknG TPR
Paf67
Cohin load
SPO22
BTAD
TPR 4
TPR 3
TPR 21
Coatomer E
TPR 15
ChAPs
NARP1
TPR 17
ANAPC3
SHNi-TPR
Fis1 TPR C
TPR 6
TPR 14
TPR 9
TPR 16
TPR 19
TPR 10
TPR 12
TPR 7
TPR 11
TPR 8
TPR 1
TPR 2
LRV
NSF
Cla propel
TALffector
LRV FeS
Clain-link
PUF
HAT
IBB
PPTA
Upf2
PC rep
Arm 3
HEAT PBS
Fis1 TPR N
Sel1
W2
PHAT
PPP5
Cla H link
GUN4 N
EST1