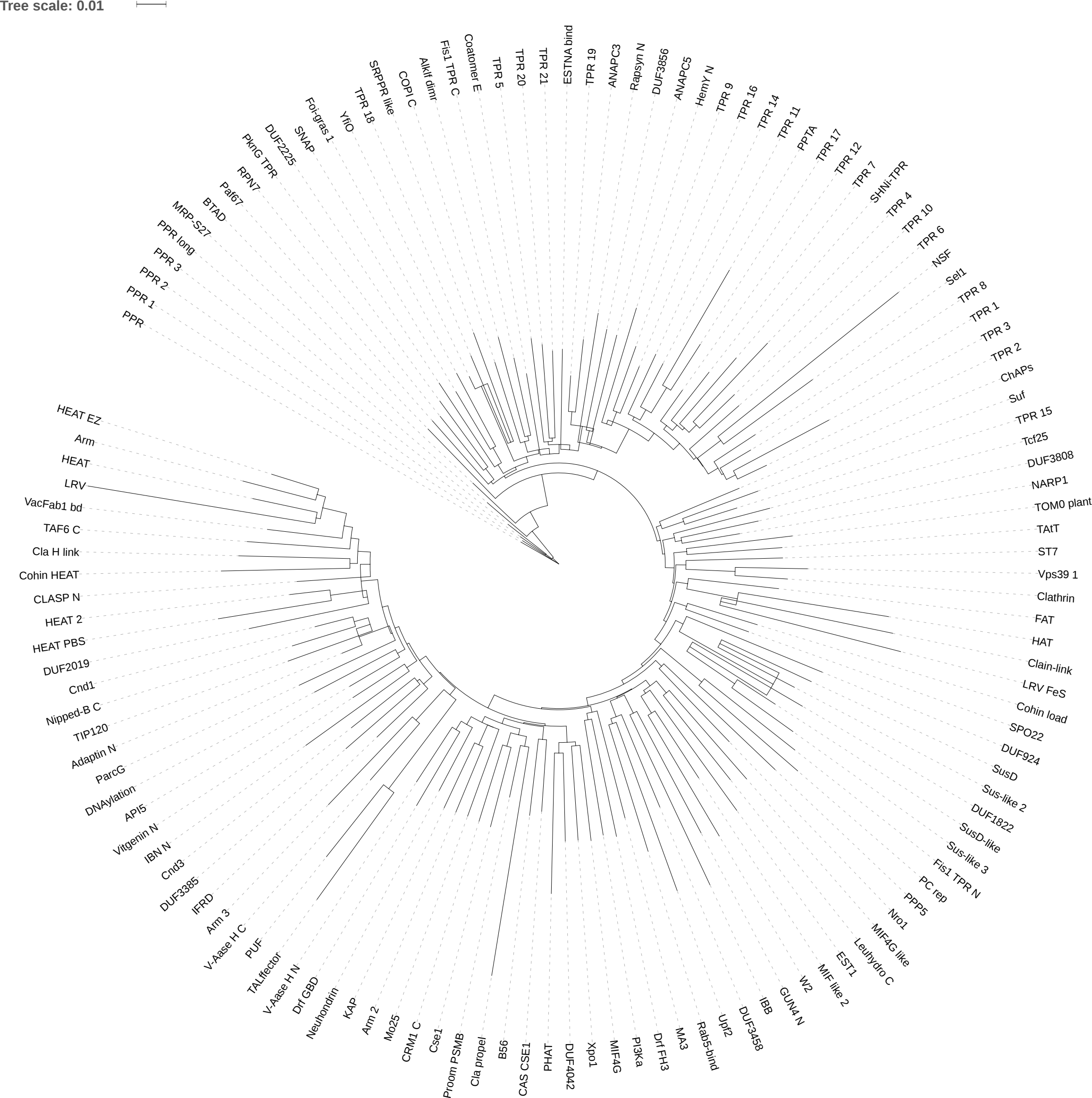
Tree scale: 0.01
ESTNA bind
TPR 19
ANAPC3
Rapsyn N
DUF3856
ANAPC5
HemY N
TPR 9
TPR 16
TPR 14
TPR 11
PPTA
TPR 17
TPR 12
TPR 7
SHNi-TPR
TPR 4
TPR 10
TPR 6
NSF
Sel1
TPR 8
TPR 1
TPR 3
TPR 2
ChAPs
Suf
TPR 15
Tcf25
DUF3808
NARP1
TOM0 plant
TAtT
ST7
Vps39 1
Clathrin
FAT
HAT
Clain-link
LRV FeS
Cohin load
SPO22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