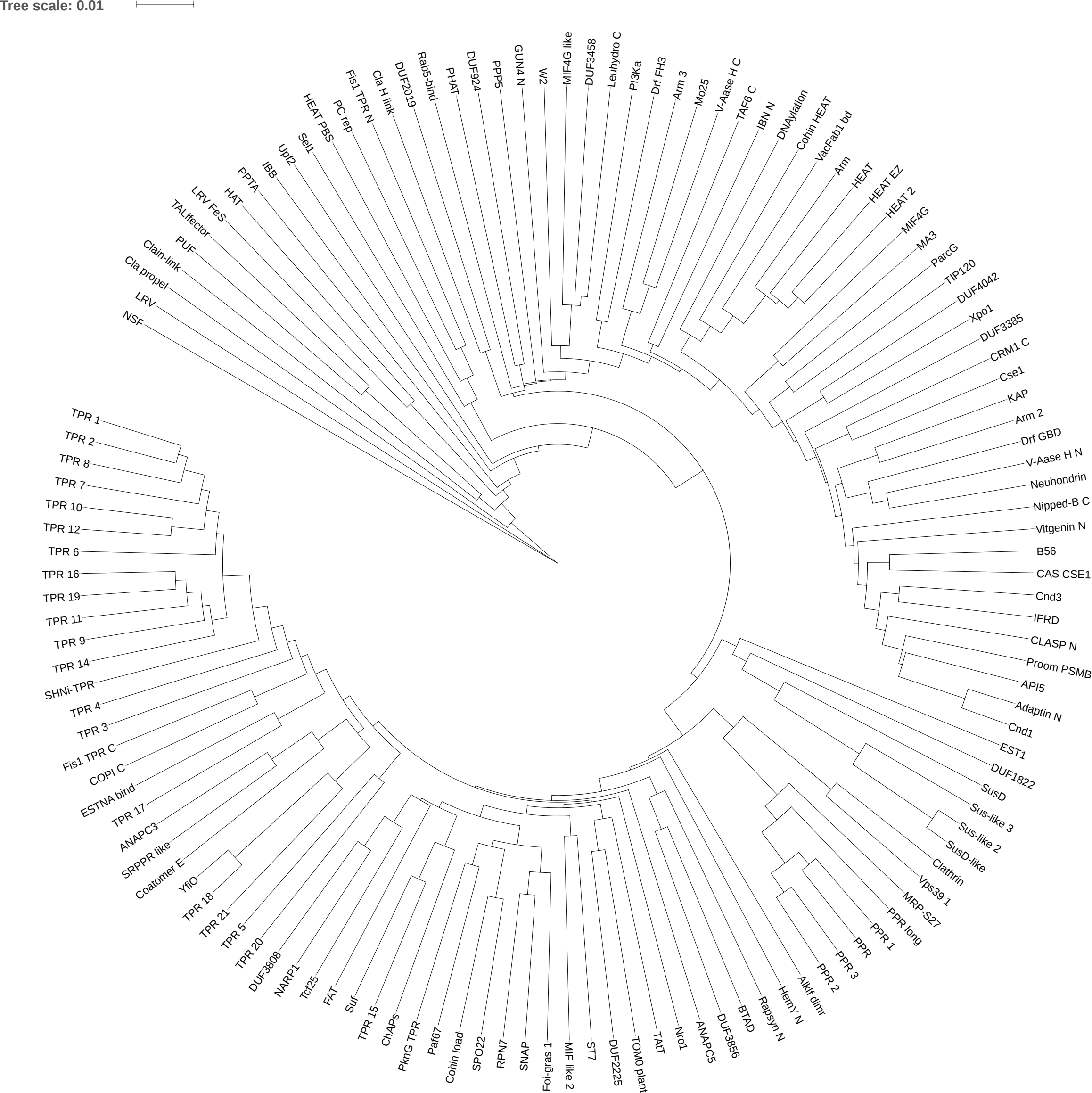
Tree scale: 0.01
NSF
LRV
Cla propel
Clain-link
PUF
TALffector
LRV FeS
HAT
PPTA
IBB
Upf2
Sel1
HEAT PBS
PC rep
Fis1 TPR N
Cla H link
DUF2019
Rab5-bind
PHAT
DUF924
PPP5
GUN4 N
W2
MIF4G like
DUF3458
Leuhydro C
PI3Ka
Drf FH3
Arm 3
Mo25
V-Aase H C
TAF6 C
IBN N
DNAylation
Cohin HEAT
VacFab1 bd
Arm
HEAT
HEAT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