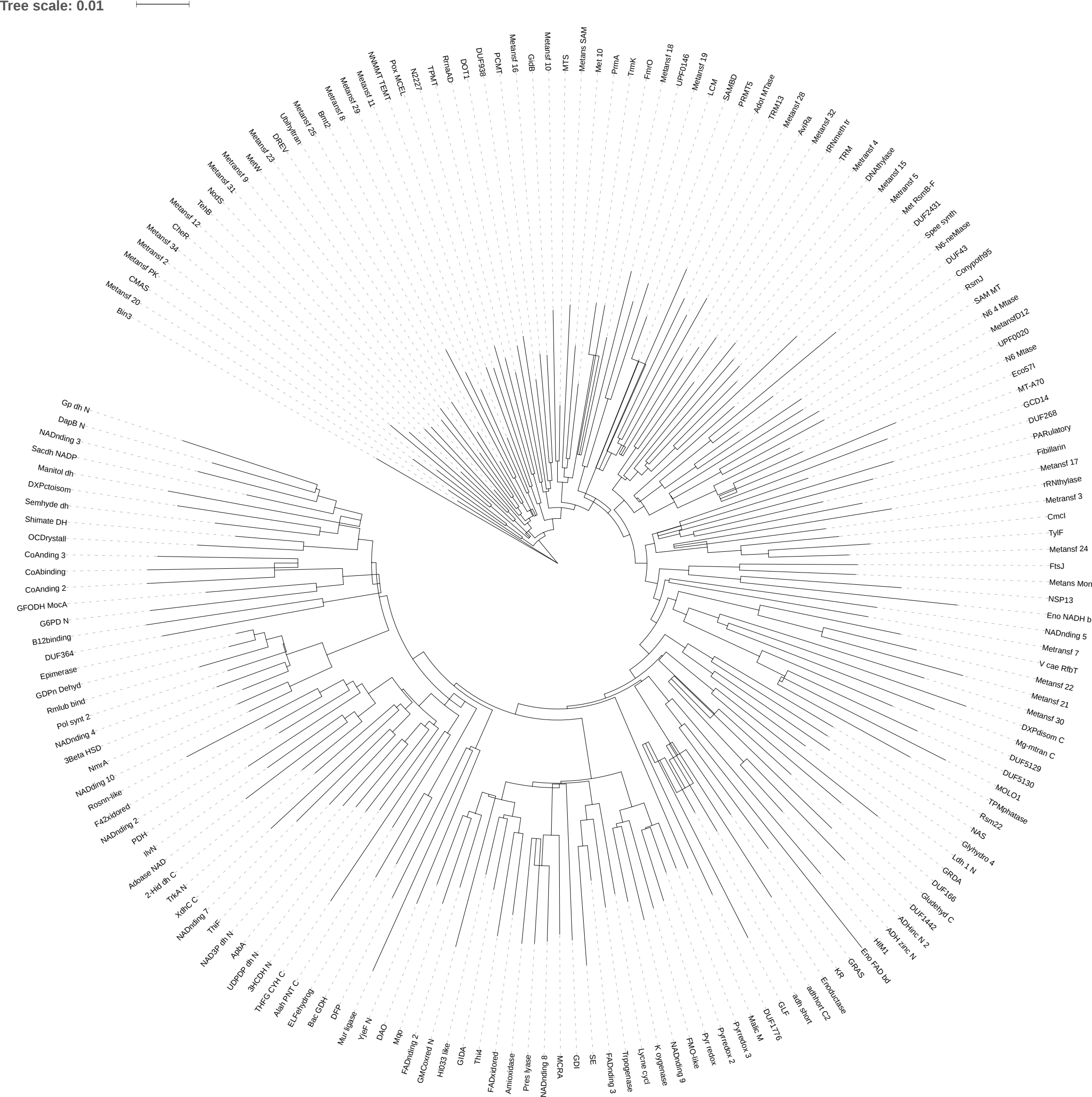
Tree scale: 0.01
Metans SAM
Met 10
PrmA
TrmK
FmrO
Metansf 18
UPF0146
Metansf 19
LCM
SAMBD
PRMT5
Adot MTase
TRM13
Metansf 28
AviRa
Metansf 32
tRNmeth tr
TRM
Metransf 4
DNAthylase
Metansf 15
Metransf 5
Met RsmB-F
DUF2431
Spee synth
N6-neMtase
DUF43
Conypothi95
RsmJ
SAM MT
N6 4 Mtase
MetansfD12
UPF0020
N6 Mtase
Eco57I
MT-A70
GCD14
DUF268
PARulatory
Fibillarin
Metansf 17
rRNthylase
Metransf 3
CmcI
TylF
Metansf 24
FtsJ
Metans Mon
NSP13
Eno NADH b
NADnding 5
Metransf 7
V cae RfbT
Metansf 22
Metansf 21
Metansf 30
DXPdisom C
Mg-mtran C
DUF5129
DUF5130
MOLO1
TPMphatase
Rsm22
NAS
Glyhydro 4
Ldh 1 N
GRDA
DUF166
Gludehyd C
DUF1442
ADHinc N 2
ADH zinc N
HIM1
Eno FAD bd
GRAS
KR
Enoductase
adhhort C2
adh short
GLF
DUF1776
Malic M
Pyrredox 3
Pyrredox 2
Pyr redox
FMO-like
NADnding 9
K oygenase
Lycne cycl
Trpogenase
FADnding 3
SE
GDI
MCRA
NADnding 8
Pres lyase
Amioxidase
FADxidored
Thi4
GIDA
HI033 like
GMCoxred N
FADnding 2
Mqo
DAO
YjeF N
Mur ligase
DFP
Bac GDH
ELFehydrog
Alah PNT C
THFG CYH C
3HCDH N
UDPDP dh N
ApbA
NAD3P dh N
ThiF
NADnding 7
XdhC C
TrkA N
2-Hid dh C
Adoase NAD
IlvN
PDH
NADnding 2
F42xidored
Rosnn-like
NADding 10
NmrA
3Beta HSD
NADnding 4
Pol synt 2
Rmlub bind
GDPn Dehyd
Epimerase
DUF364
B12binding
G6PD N
GFODH MocA
CoAnding 2
CoAbinding
CoAnding 3
OCDrystall
Shimate DH
Semhyde dh
DXPctoisom
Manitol dh
Sacdh NADP
NADnding 3
DapB N
Gp dh N
Bin3
Metansf 20
CMAS
Metansf PK
Metransf 2
Metansf 34
CheR
Metansf 12
TehB
NodS
Metansf 31
Metransf 9
MetW
Metansf 23
DREV
Ubihyltran
Metansf 25
Bmt2
Metransf 8
Metansf 29
Metansf 11
NNMMT TEMT
Pox MCEL
N2227
TPMT
RrnaAD
DOT1
DUF938
PCMT
Metansf 16
GidB
Metansf 10
MTS