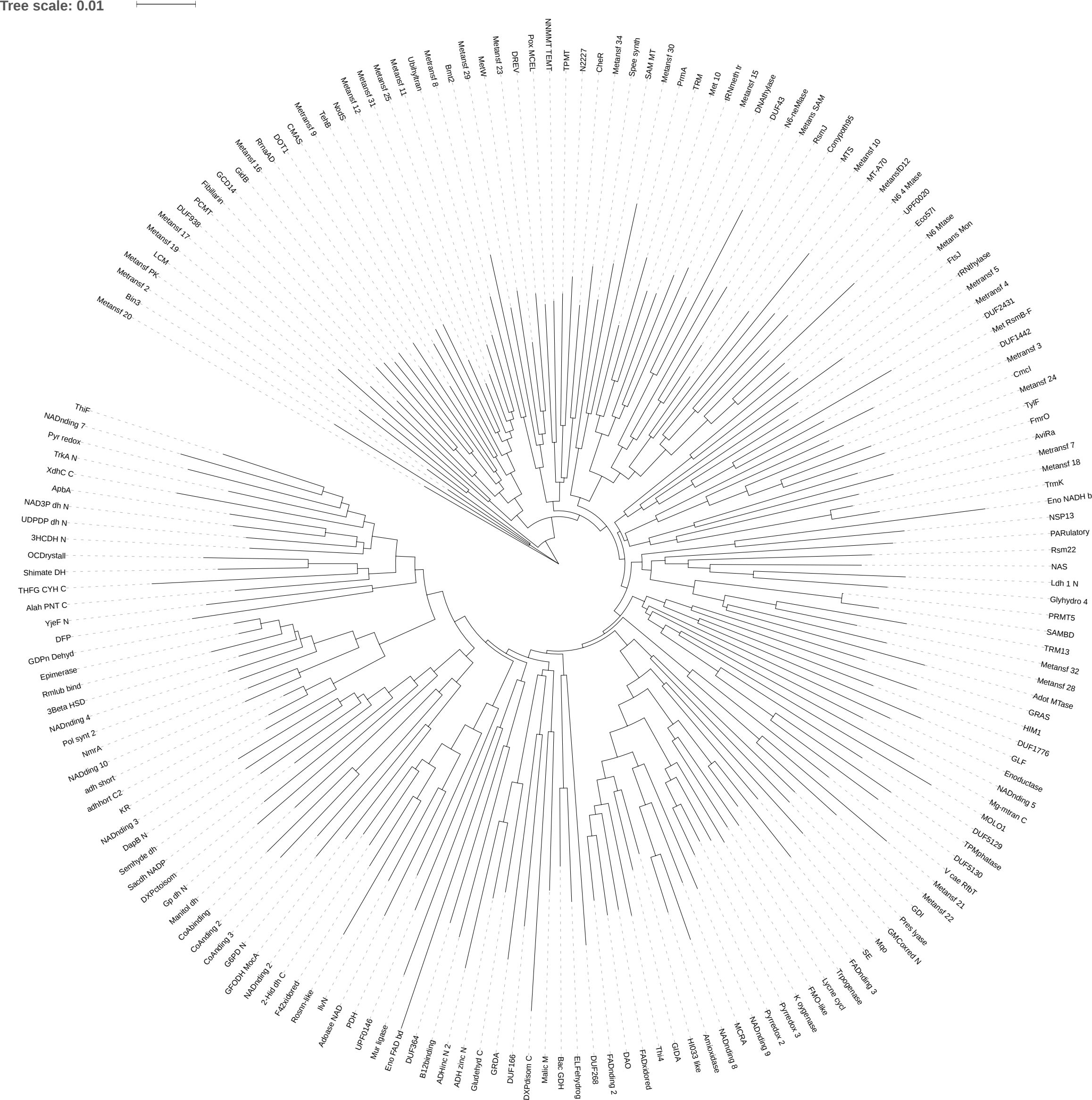
Tree scale: 0.01
NNMMT TEMT
TPMT
N2227
CheR
Metansf 34
Spee synth
SAM MT
Metansf 30
PrmA
TRM
Met 10
tRNmeth tr
Metansf 15
DNAthylase
DUF43
N6-neMtase
Metans SAM
RsmJ
Conypothi95
MTS
Metansf 10
MT-A70
MetansfD12
N6 4 Mtase
UPF0020
Eco57I
N6 Mtase
Metans Mon
FtsJ
rRNthylase
Metransf 5
Metransf 4
DUF2431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