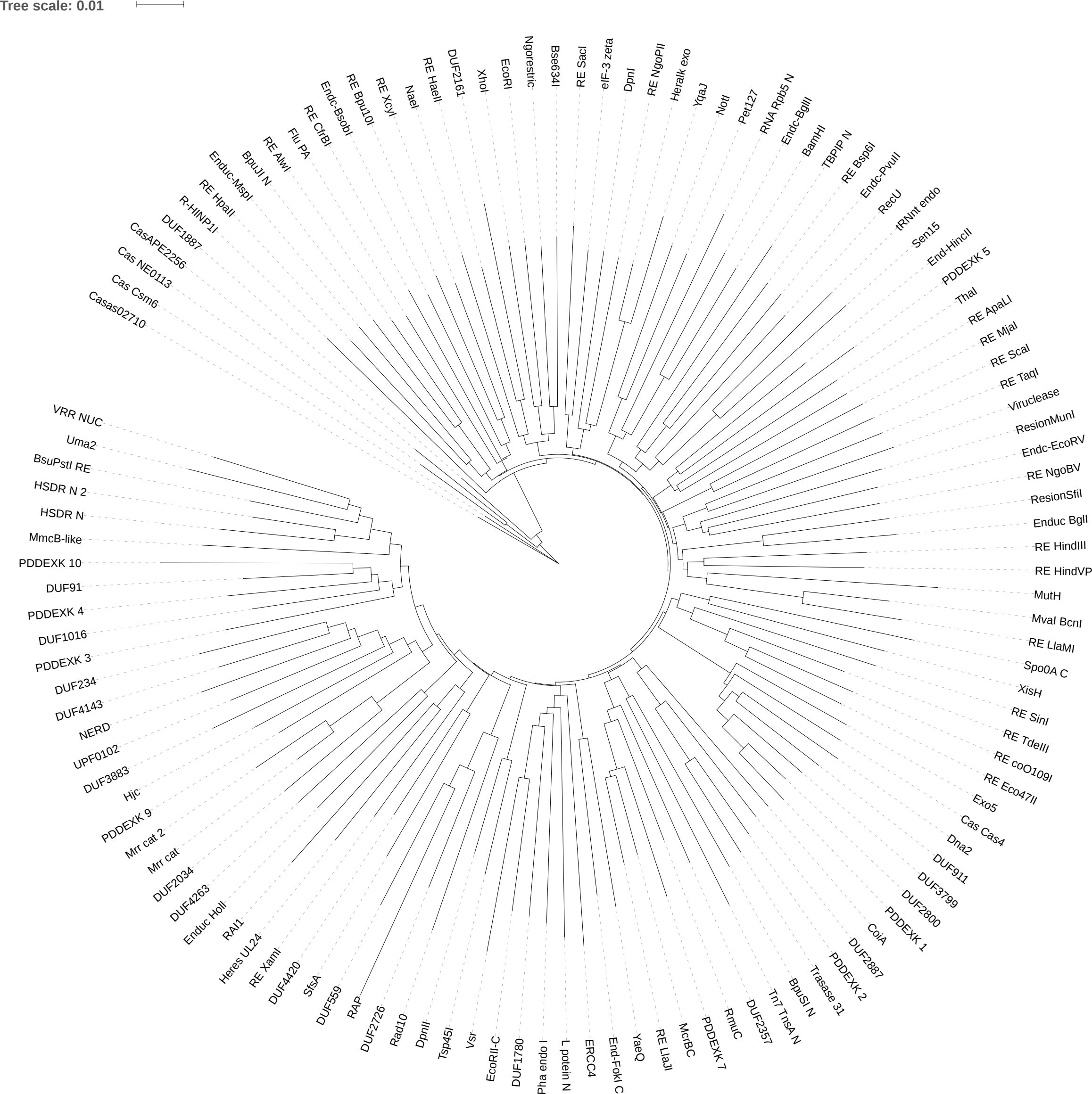
Tree scale: 0.01
Ngorestric
Bse634I
RE SacI
eIF-3 zeta
DpnI
RE NgoPII
Heralk exo
YqaJ
NotI
Pet127
RNA Rpb5 N
Endc-BglII
BamHI
TBPIP N
RE Bsp6I
Endc-PvuII
RecU
tRNnt endo
Sen15
End-HincII
PDDEXK 5
ThaI
RE ApaLI
RE MjaI
RE ScaI
RE TaqI
Viruclease
ResionMunI
Endc-EcoRV
RE NgoBV
ResionSfiI
Enduc BglI
RE HindIII
RE HindVP
MutH
MvaI BcnI
RE LlaMI
Spo0A C
XisH
RE SinI
RE TdeIII
RE coO109I
RE Eco47II
Exo5
Cas Cas4
Dna2
DUF911
DUF3799
DUF2800
PDDEXK 1
CoiA
DUF2887
PDDEXK 2
Trasase 31
BpuSI N
Tn7 TnsA N
DUF2357
RmuC
PDDEXK 7
McrBC
RE LlaJI
YaeQ
End-FokI C
ERCC4
L potein N
Pha endo I
DUF1780
EcoRII-C
Vsr
Tsp45I
DpnII
Rad10
DUF2726
R4P
DUF559
SfsA
DUF4420
RE XamI
Heres UL24
RAI1
Enduc HolI
DUF4263
DUF2034
Mrr cat
Mrr cat 2
PDDEXK 9
Hjc
DUF3883
UPF0102
NERD
DUF4143
DUF234
PDDEXK 3
DUF1016
PDDEXK 4
DUF91
PDDEXK 10
MmcB-like
HSDR N
HSDR N 2
BsuPstI RE
Uma2
VRR NUC
Casas02710
Cas Csm6
Cas NE0113
CasAPE2256
DUF1887
R-HINP1I
RE HpaII
Enduc-MspI
BpuJI N
RE AlwI
Flu PA
RE CfrBI
Endc-BsobI
RE Bpu10I
RE XcyI
NaeI
RE HaeII
DUF2161
XhoI
EcoRI