Tree scale: 0.01

HTHB IS605
HSFNA-bind
DUF3908
YokU
La
DUF3116
Exc
RFXbinding
Pox D5
Ets
IRF
Trai 2 C
HTHTnp Tc5
ARID
RapNA-bind
Myb-bind 2
SLIDE
SANT-1 like
Myb-bind 7
Mybbinding
Myb-bind 6
Myb-bind 3
Myb-bind 5
Myb-bind 4
MADSNA bdg
HTH 16
Sigma70 r2
MotA activ
DUF1836
CUT
OST-HTH
Forkhead
DEP
RTP
PhaItermQ
Viralpha C
HTHTnp IS1
IBD
HTHnp Mu 2
ESCRT-II
AbiEi 3 N
HPD
Fe repr C
FokI N
DUF3860
Tn916-Xis
Homobox KN
Homez
Pou
KicB
DUF739
HARE-HTH
HTH 33
HTH 21
SWIRM
DUF3253
HTHnp Mu 1
DUF1133
TFIF alpha
ELL
TFIIF beta
Mqsitoxin
DUF1870
DUF4447
RepA C
RepC
TrfA
RPA
Rep 3
DUF1153
Sel-wing 1
F-112
E2F TDP
RepA N
STN1 2
YjcQ
FokI C
Sigar70 ner
SAC3 GANP
CSN8 EIF3K
DUF480
Tn7 TnsA C
HTH 39
DsrD
Linhistone
RQC
ANAPC2
Sel-wing 2
HTH 13
AphA like
DUF1495
TFIIE beta
DUF2582
HTH 42
Sel-wing 3
HTHp Tc3 1
PAX
Cdc6 C
MerNA-bind
BrkDBD
SatD
Cro
Tras reg C
DUF433
BetR
DUF1492
DUF3853
Bac DnaA C
Culn Nedd8
Phap org N
MerR 2
Siga54 CBD
Phage CII
HTH 43
EAP30
RuvB C
Ribal S19e
DUF1441
DUF2316
Bot1p
HTH WhiA
HTH 32
Argpressor
HTHp Tc3 2
Sulbus pRN
PuR N
MRP-L20
Neugrin
GcrA
Mor
kORA
Dimsation
Dimsation2
z-alpha
HemN C
HTH 46
FUR
Mnd1
TBPIP
Rio2 N
HTH 12
HTH 45
DUF4364
Repc Relax
HxlR
PadR
DUF977
KorB
HTH 47
DUF7a2
DUF2250
Ribmal S25
Ftsk gamma
DDRGK
PCI
Put-bind N
HTH 15
SMC ScpB
IF2 N
DUF134
Homeobox
Siga54 DBD
HTH 35
HTH AraC
HTH 18
DUF1323
DUF1804
DUF1670
SpoIIID
Tnpressor
NUMOD1
PhaCI repr
YdaItoxin
P22 Cro
LacI
HNF-1 N
HTH 37
HTH 25
HTH 26
HTH 31
HTH 19
HTH 3
HTHp IS630
PaaX
DUF2089
Sigma70 r3
Pharminase
HTHTnp 1 2
AbiEi 4
Phage AlpA
MerR 1
HTH 17
MerR
ReplL
TFIE alpha
HTH micro
RNAI Rpc34
B.b TFIIIC
HTH CodY
HTH 41
Mga
TetR N
UPF0122
HTH 40
HTH 10
HTH 22
HTH 1
HTH 8
HTH 30
Siga70 ECF
HTH psq
LZ p IS481
HTH Tnp 4
GerE
Sig70 r4 2
Sigma70 r4
HTHnp ISL3
CENP-B N
HTH Tnp 1
HTH 38
HTH 7
Teninase 5
HTH 23
HTH 29
HTH 28
HTH Mga
SgrR N
RPA C
FeoC
FaeA
HrcDNA-bdg
RP-C
Peninase R
HTH 6
HTH 34
LexNA bind
HTH 36
Crp
HTH Crp 2
Rrf2
GntR
HTH 11
HTH DeoR
Fe repress
HTH 27
TrmB
HTH 5
HTH 20
HTH IclR
MarR
MarR 2
HTH 24
HTHnC-type