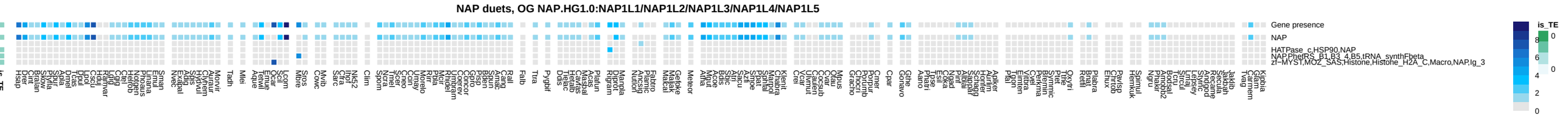
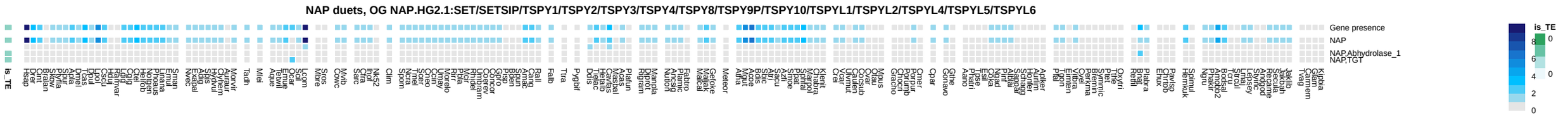
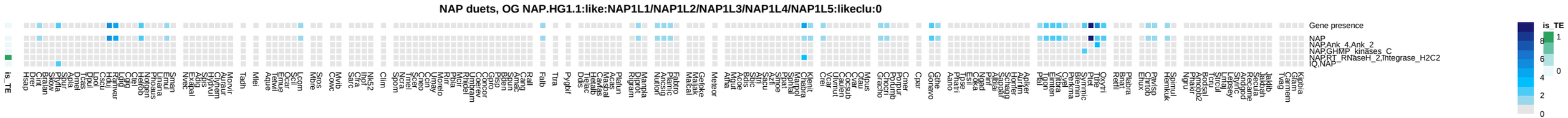
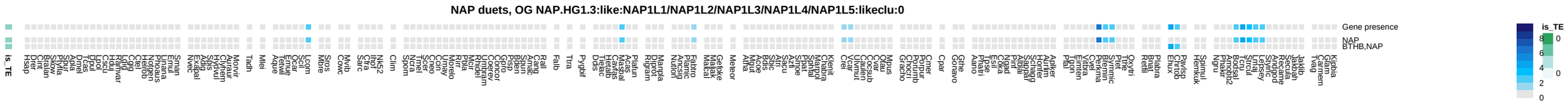
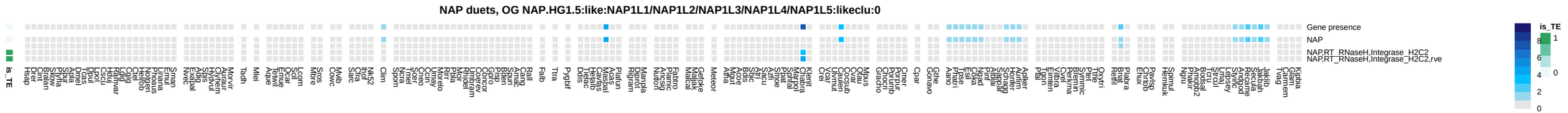
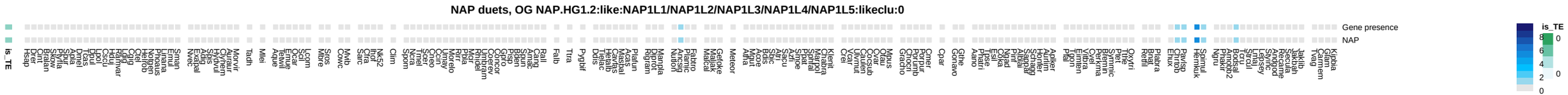

NAP duets, OG NAP.HG1.0:NAP1L1/NAP1L2/NAP1L3/NAP1L4/NAP1L5
Gene presence
NAP
HATPase_c,HSP90,NAP
NAP,PhefRS_B1,B3_4,B5,tRNA_synthFbeta
zf−MYST,MOZ_SAS,Histone,Histone_H2A_C,Macro,NAP,Ig_3
is_TE
is_TE
8
6
4
2
0
0
0


NAP duets, OG NAP.HG2.1:SET/SETSIP/TSPY1/TSPY2/TSPY3/TSPY4/TSPY8/TSPY9P/TSPY10/TSPYL1/TSPYL2/TSPYL4/TSPYL5/TSPYL6
Gene presence
NAP
NAP,Abhydrolase_1
NAP,TGT
is_TE
is_TE
8
6
4
2
0
0
0


NAP duets, OG NAP.HG1.1:like:NAP1L1/NAP1L2/NAP1L3/NAP1L4/NAP1L5:likeclu:0
Gene presence
NAP
NAP,Ank_4,Ank_2
NAP,GHMP_kinases_C
NAP,RT_RNaseH_2,Integrase_H2C2
IQ,NAP
is_TE
1
0
8
6
4
2
0
is_TE
Hsap
Drer
Ornan
Bralan
Skow
Bflo
Bfla
Apla
Dmel
Dpul
Lpol
Lgig
Cgig
Cteli
Helrob
Nvec
Sman
Emul
Lhana
Hvulgas
Nvitr
Ccsu
Ccsu
Rsanvar
Lgig
Cgig
Cteli
Helrob
Nvec
Exapal
Adig
Aflis
Hdivul
Clvnem
Mpovir
Aqcoar
Mlei
Tadh
Mlei
Aqfve
Aqfve
Emue
Ocar
Ocar
Loom
Loom
Spos
More
Mvib
Cowc
Nvit52
Ihcf
Cfra
Sarc
Clim
Spom
Ncra
Tmel
Cneo
Sscr
Ccin
Umay
Umaydo
Rirr
Mpla
Mlel
Mlel
Umbram
Coerev
Coerev
Gpro
Pisp
Bbren
Bbren
Bmin
Amac
Amac
Rall
Cang
Falb
Falb
Ttra
Pygbif
Tralec
Dds
Hetalb
Cassp
Platun
Acas
Acas
Rigram
Dipot
Mampla
Nuriof
Ancsig
Pfamic
Pfamic
Fabtro
Gallke
Gallke
Malcal
Meteor
Atha
Atha
Acose
Acose
Mgut
Sbic
Sbic
Atin
Atin
Sacu
Sacu
Smoe
Aphal
Phat
Sarhal
Marpol
Klenit
Klenit
Chabra
Vcar
Crei
Vcar
Crei
Umuut
Umuut
Cocsub
Cvar
Cvar
Olus
Mnus
Gracho
Gracho
Porpur
Porpur
Porumb
Porumb
Cmer
Cpar
Cpar
Gthe
Gthe
Gonavo
Aano
Aano
Pbiser
Esil
Esil
Ostka
Ostka
Nvad
Pnr
Nvad
Ostka
Abdal
Abdal
Sphag
Sphag
Sphag
Aurlim
Aurlim
Honter
Honter
Ankker
Ankker
Pal
Pal
Tpon
Tpon
Ellilen
Ellilen
Vibra
Vibra
Cvel
Cvel
Petkma
Petkma
Symmic
Symmic
Pter
Pter
Ovvri
Ovvri
Rettil
Rettil
Blabra
Blabra
Elux
Elux
Pavisp
Pavisp
Ovvri
Ovvri
Hemkuk
Hemkuk
Spimul
Spimul
Amodpb2
Amodpb2
Ngru
Ngru
Bodsal
Bodsal
Tcru
Tcru
Tcru
Lmaj
Lmaj
Lepsey
Lepsey
Sscrofa
Sscrofa
Acyrosp
Acyrosp
Recanne
Recanne
Secula
Secula
Jaldla
Jaldla
Nvlp
Nvlp
Klpbia
Klpbia
Glam
Glam
Carmhem
Carmhem
Tvag
Tvag

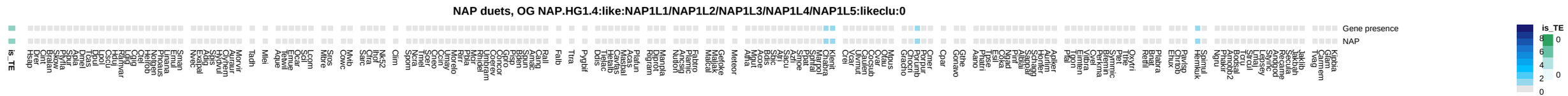

NAP duets, OG NAP.HG1.3:like:NAP1L1/NAP1L2/NAP1L3/NAP1L4/NAP1L5:likeclu:0
Gene presence
NAP
BTHB,NAP
is_TE
is_TE
8
6
4
2
0
0
0


NAP duets, OG NAP.HG1.5:like:NAP1L1/NAP1L2/NAP1L3/NAP1L4/NAP1L5:likeclu:0
is_TE
Gene presence
NAP
NAP,RT_RNaseH,Integrase_H2C2
NAP,RT_RNaseH,Integrase_H2C2,rve
is_TE
8
6
4
2
0
1
0


NAP duets, OG NAP.HG1.2:like:NAP1L1/NAP1L2/NAP1L3/NAP1L4/NAP1L5:likeclu:0
is_TE
Gene presence
NAP
is_TE
8
6
4
2
0
0
0

Gene presence
NAP
NAP,TTL

is_TE
8
6
4
2
0
0
0


NAP duets, OG NAP.HG1.4:like:NAP1L1/NAP1L2/NAP1L3/NAP1L4/NAP1L5:likeclu:0
Gene presence
NAP
is_TE
is_TE
8
6
4
2
0
0
0