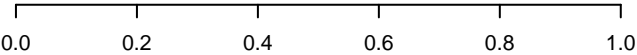


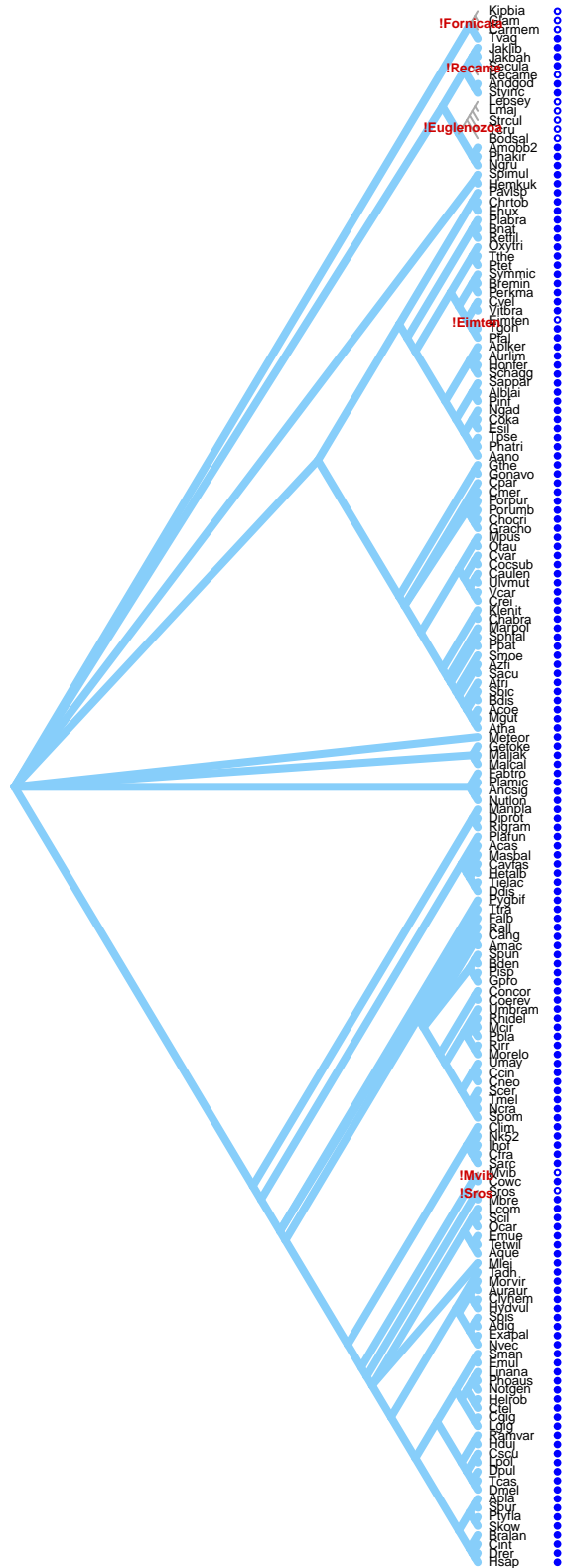
absent absent | n=1



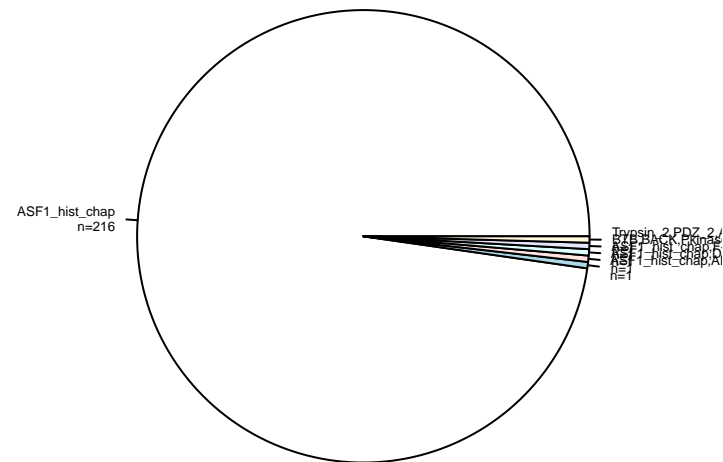
Frequency

Gain: Bnat,Euglenozoa,Fornicata,NA
Presence Eukaryota prob = 0.00
Present: 8
Losses: NA

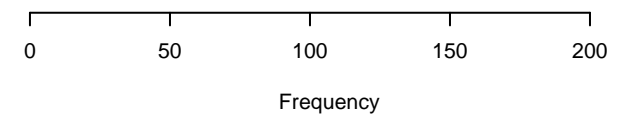
ASF1_hist_chap.HG1.1
ASF1A/ASF1B



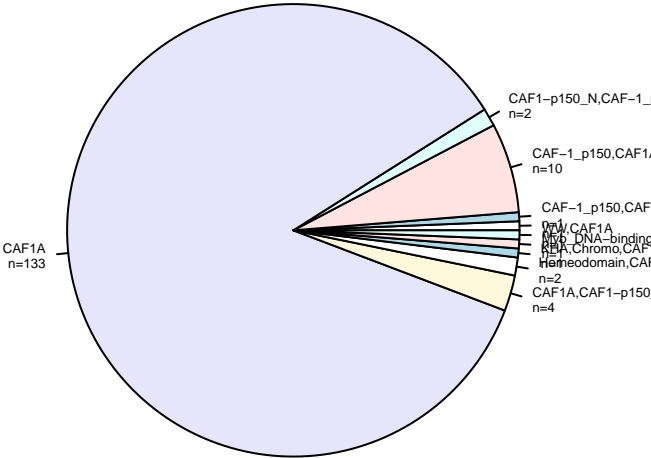
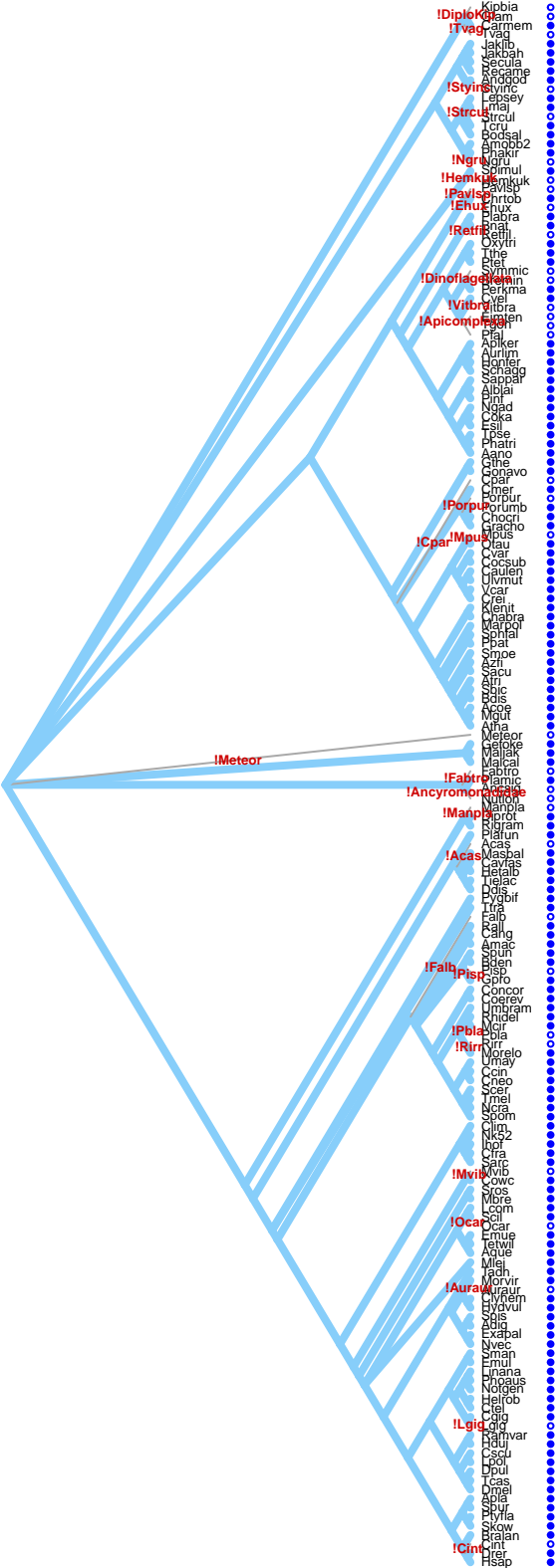
Gain: NA
Presence Eukaryota prob = 1.00
Present: 160
Losses: NA



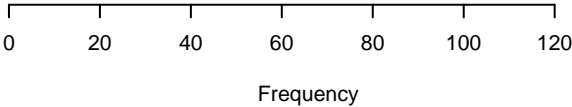
Trypsin_2_PDZ_2_ASF1_hist_chap	Trypsin_2_PDZ_2_ASF1_hist_chap n=1
BTB,BACK,Pkinase,ASF1_hist_chap	BTB,BACK,Pkinase,ASF1_hist_chap n=1
ASF1_hist_chap,F-box	ASF1_hist_chap,F-box n=1
ASF1_hist_chap,Drf_GBD,Drf_FH3	ASF1_hist_chap,Drf_GBD,Drf_FH3 n=1
ASF1_hist_chap,ADIP	ASF1_hist_chap,ADIP n=1
ASF1_hist_chap	ASF1_hist_chap n=216



CAF1A.HG1.0
CHAF1A

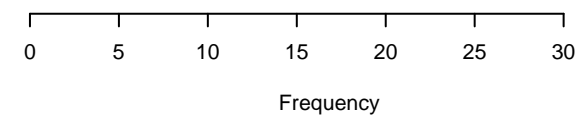
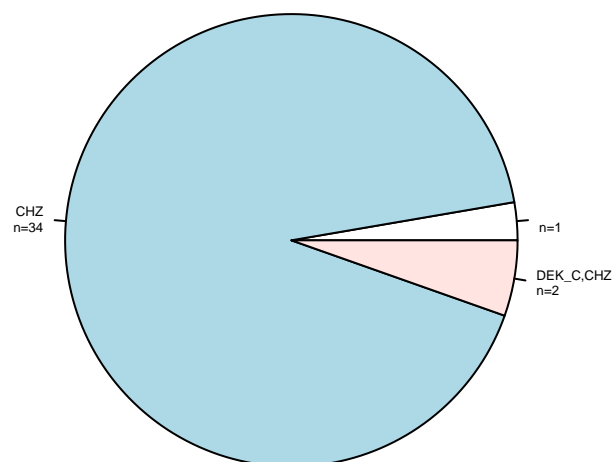
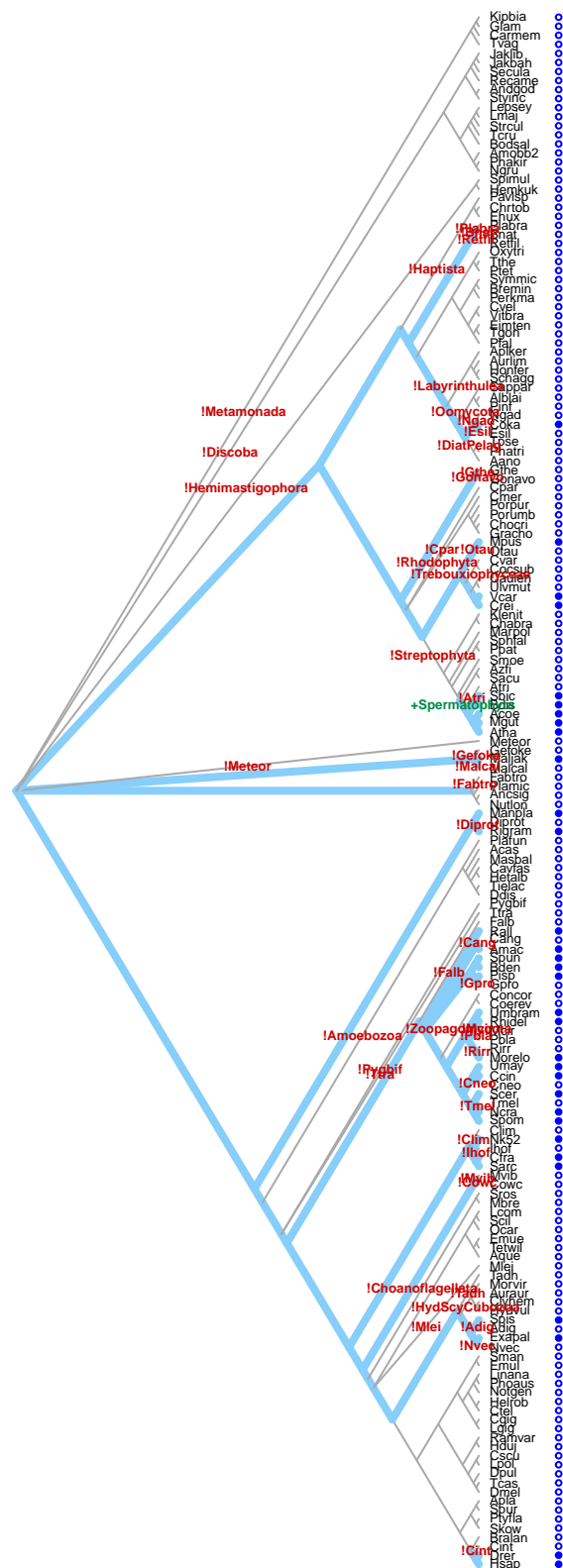


WW,CAF1A	WW,CAF1A n=1
Myb_DNA-binding,CAF1A	Myb_DNA-binding,CAF1A n=1
KHA,Chromo,CAF1A,PWWP	KHA,Chromo,CAF1A,PWWP n=1
Homeodomain,CAF1A	Homeodomain,CAF1A n=2
CAF1A,CAF1-p150_C2	CAF1A,CAF1-p150_C2 n=4
CAF1A	CAF1A n=133
CAF1-p150_N,CAF-1_p150,CAF1A,CAF1-p150_C2	CAF1-p150_N,CAF-1_p150,CAF1A,CAF1-p150_C2 n=2
CAF-1_p150,CAF1A,CAF1-p150_C2	CAF-1_p150,CAF1A,CAF1-p150_C2 n=10
CAF-1_p150,CAF1A	CAF-1_p150,CAF1A n=1
	n=1

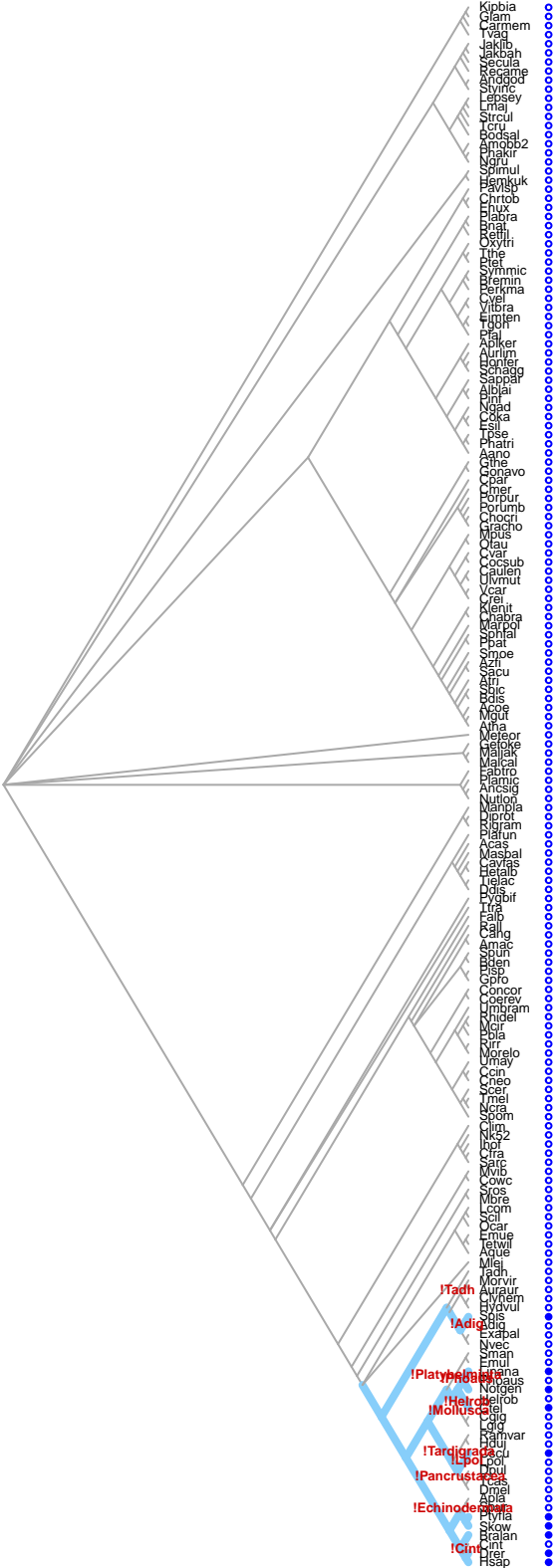


Gain: NA
Presence Eukaryota prob = 1.00
Present: 138
Losses: NA

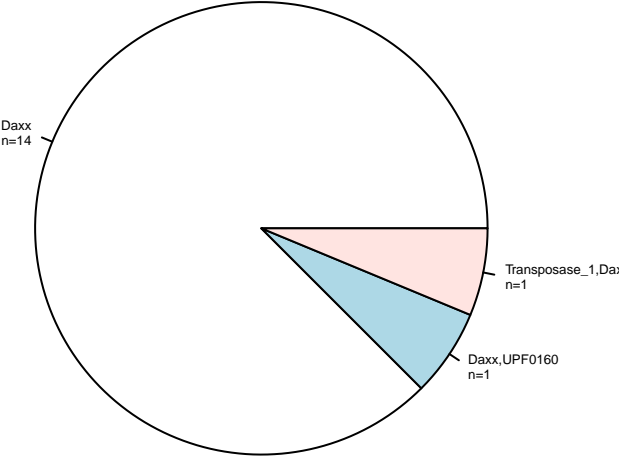
CHZ.HG1.0
HIRIP3



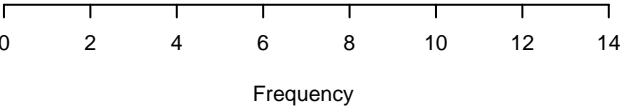
Gain: Spermatophyta,NA
Presence Eukaryota prob = 1.00
Present: 32
Losses: NA



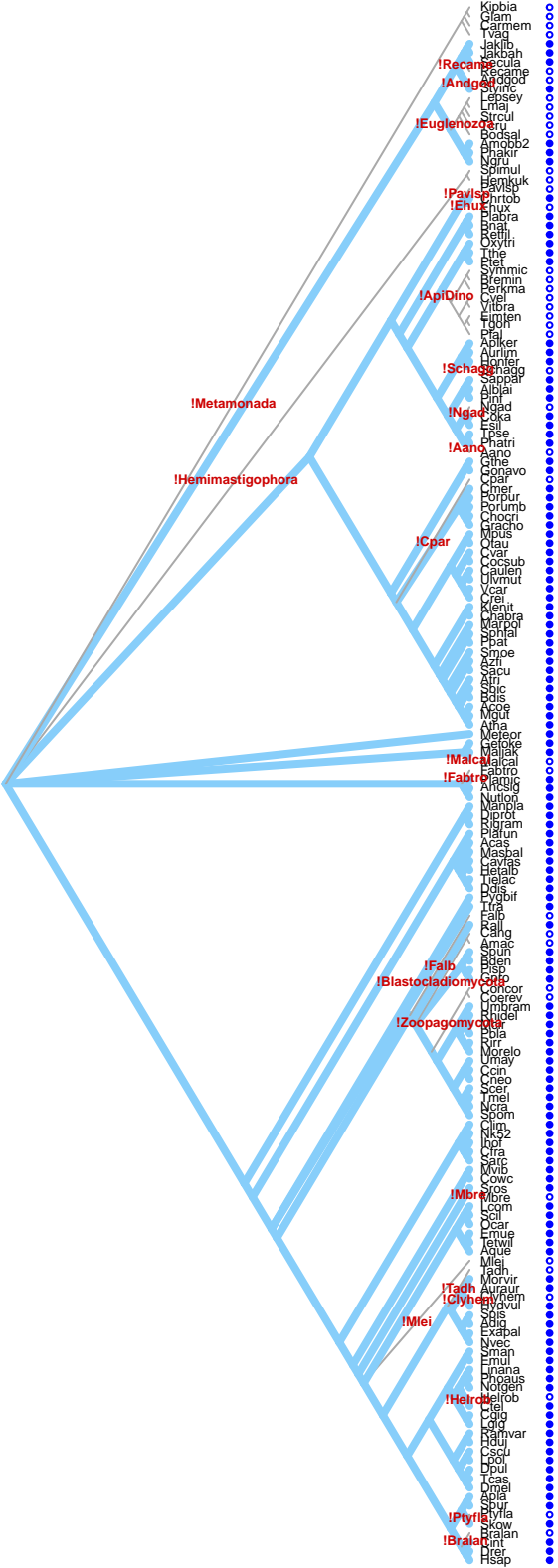
Gain: NA
Presence Eukaryota prob = 0.00
Present: 10
Losses: NA



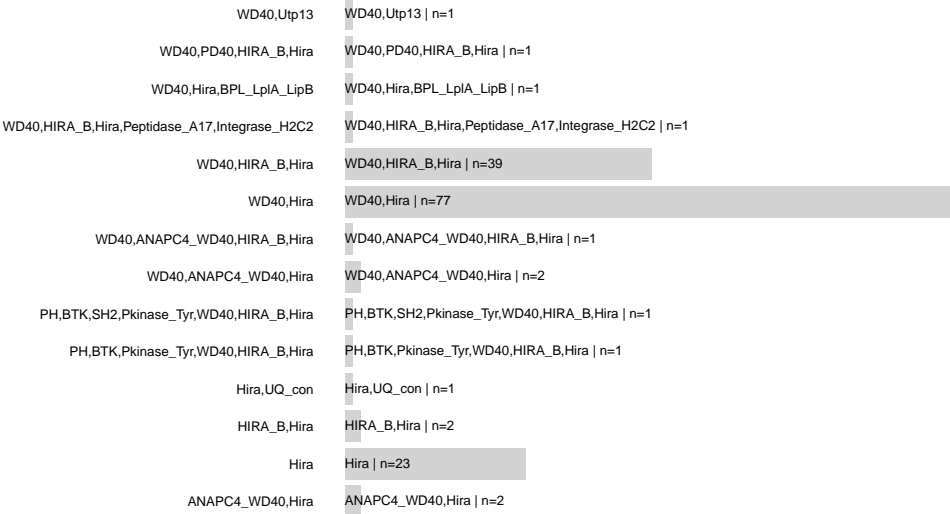
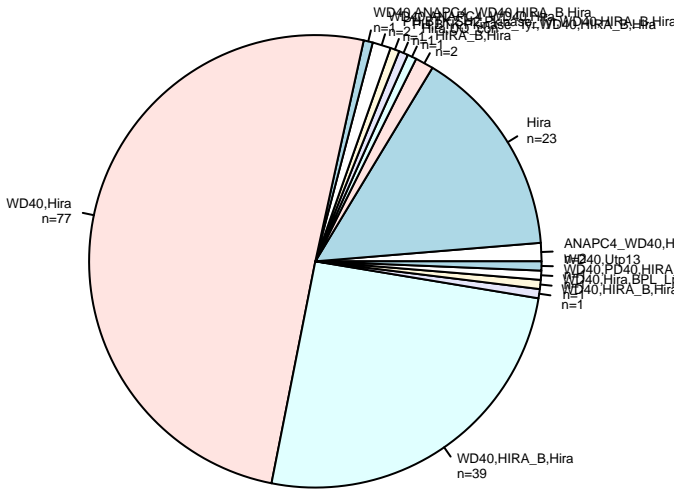
Transposase_1,Daxx | Transposase_1,Daxx | n=1
Daxx,UPF0160 | Daxx,UPF0160 | n=1
Daxx | Daxx | n=14



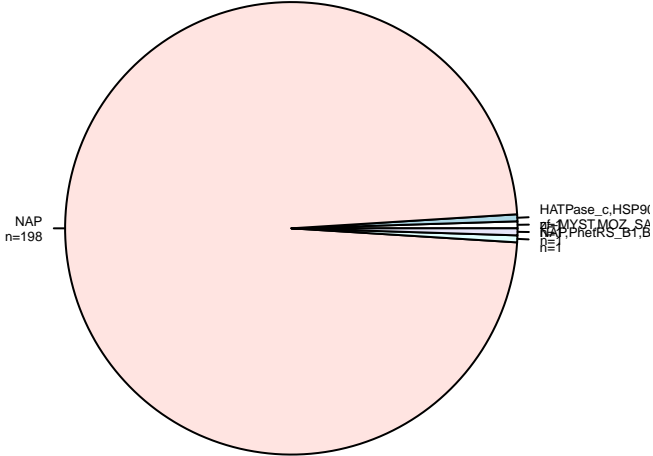
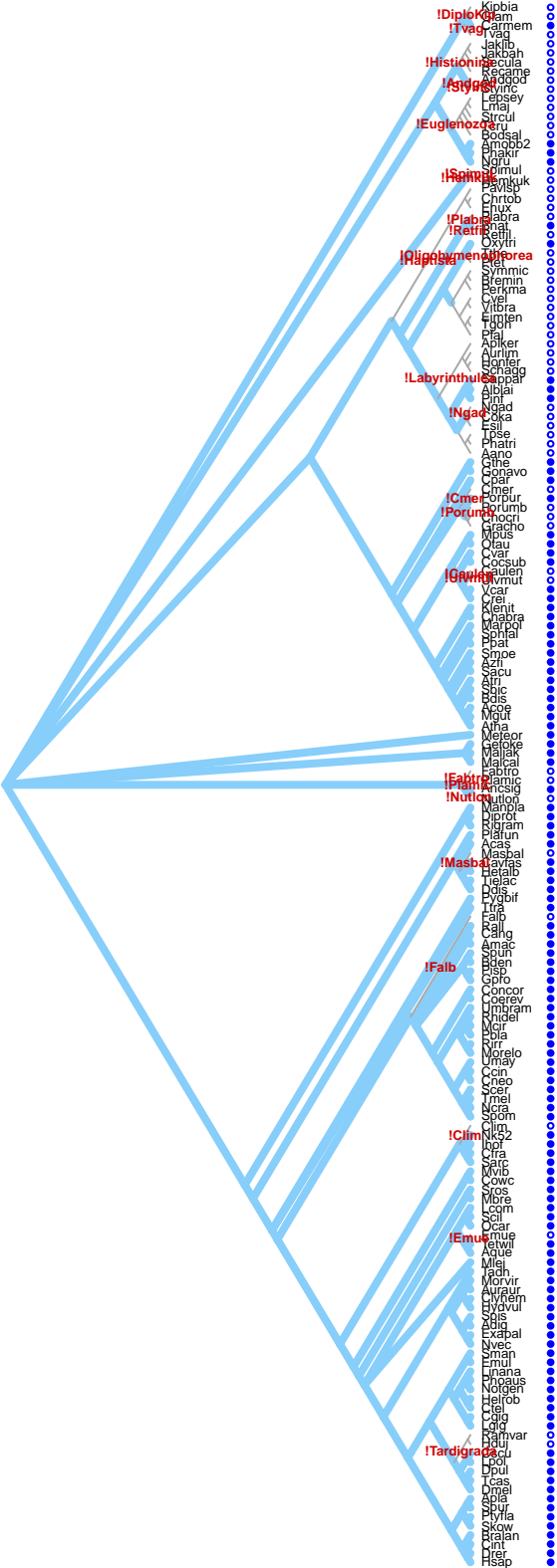
Hira.HG1.0
HIRA



Gain: NA
Presence Eukaryota prob = 1.00
Present: 131
Losses: NA

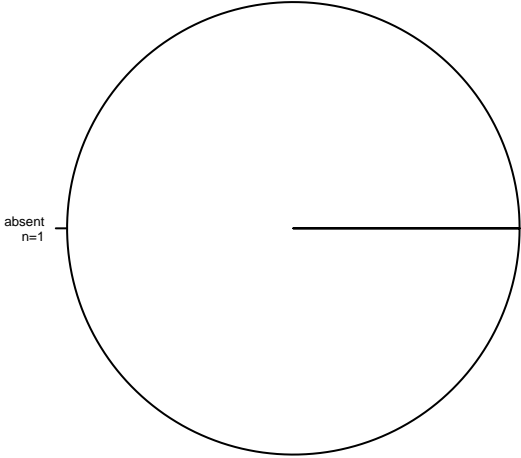
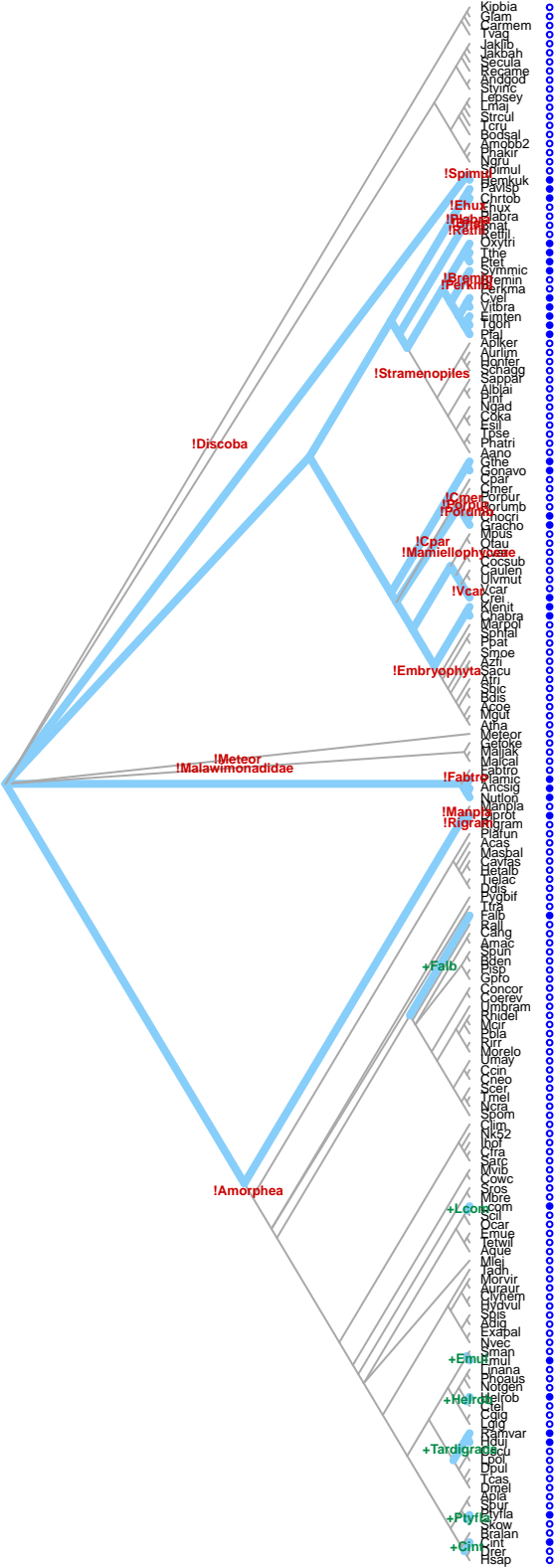


0 10 20 30 40 50 60 70
Frequency



zf-MYST,MOZ_SAS,Histone,Histone_H2A_C,Macro,NAP,Ig_3	zf-MYST,MOZ_SAS,Histone,Histone_H2A_C,Macro,NAP,Ig_3 n=1
NAP,PhetRS_B1,B3_4,B5,tRNA_synthFbeta	NAP,PhetRS_B1,B3_4,B5,tRNA_synthFbeta n=1
NAP	NAP n=198
HATPase_c,HSP90,NAP	HATPase_c,HSP90,NAP n=1
	n=1





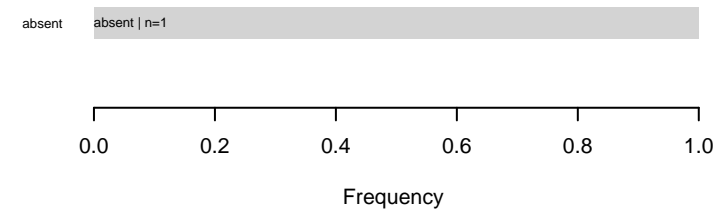
absent

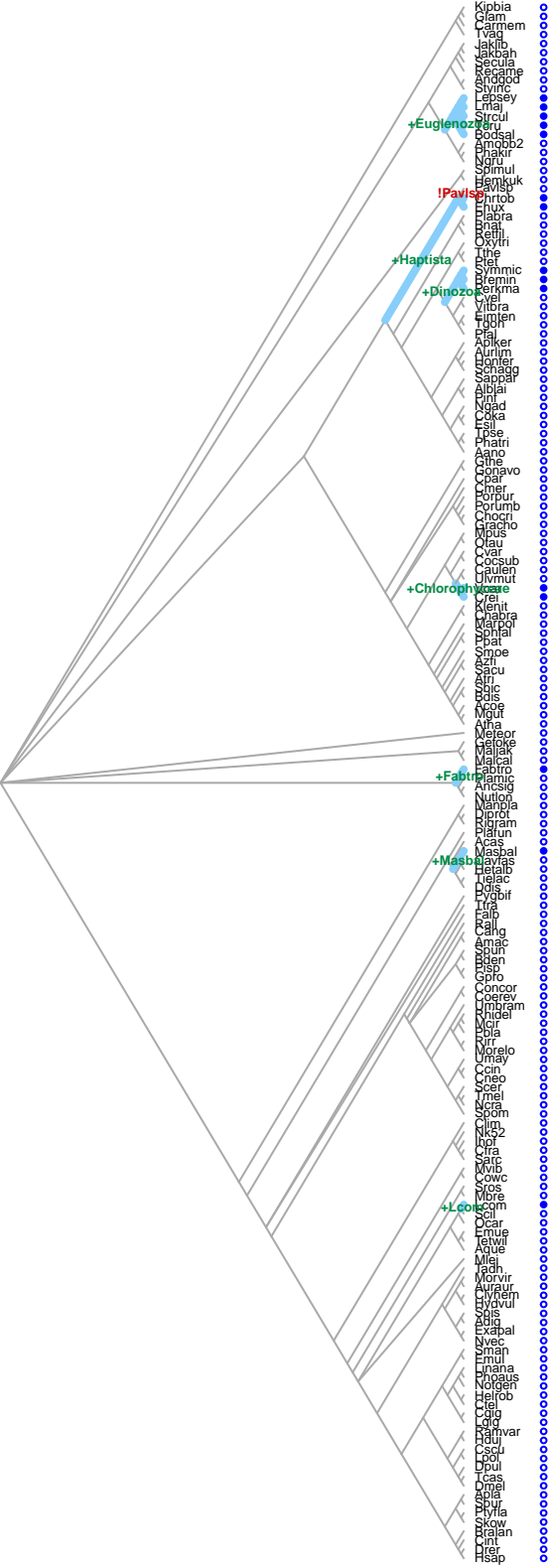
absent | n=1

0.0 0.2 0.4 0.6 0.8 1.0

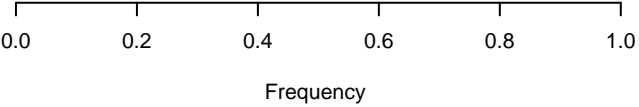
Frequency

Gain: Ancsig,Haptista,Hemimastigophora,Bodsal,NA
 Presence Eukaryota prob = 0.00
 Present: 6
 Losses: NA

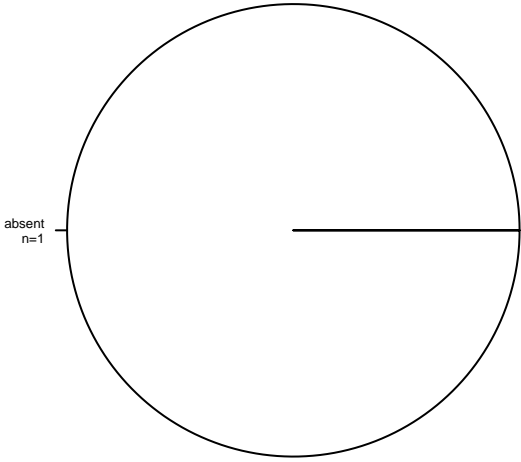
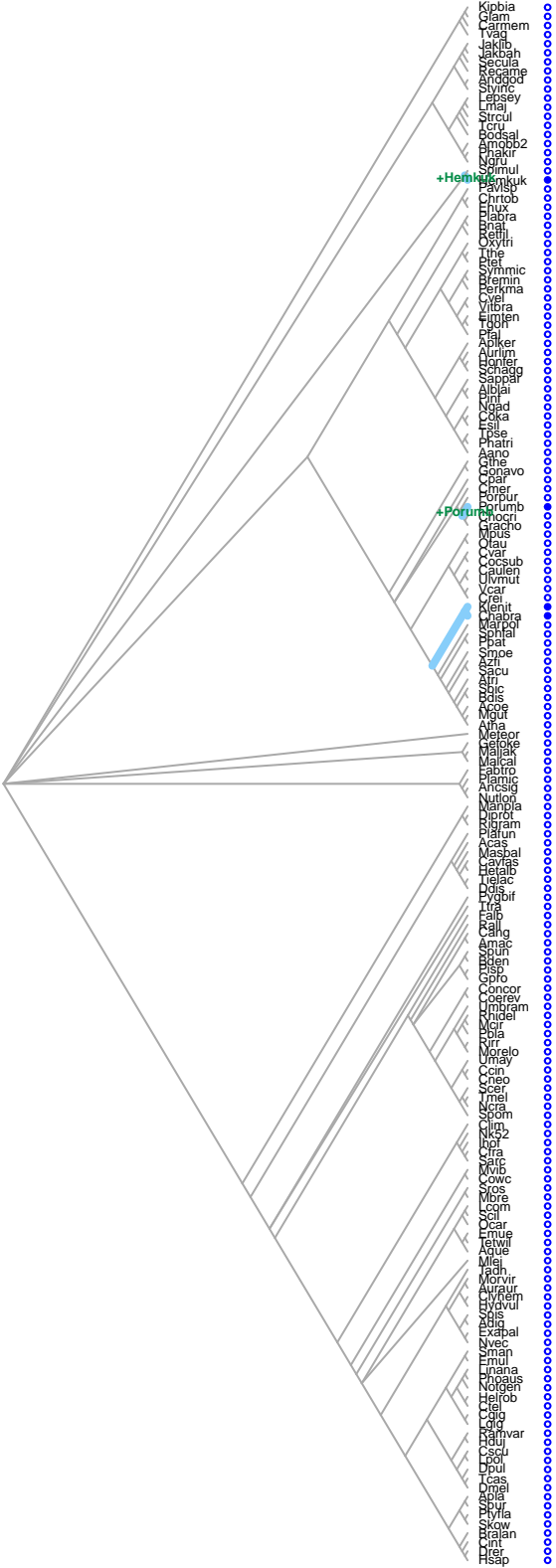




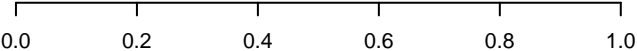
absent absent | n=1



Gain: Lcom,Masbal,Fabtro,Chlorophyceae,Dinzoa,Haptista,Euglenozoa,NA
Presence Eukaryota prob = 0.00
Present: 15
Losses: NA

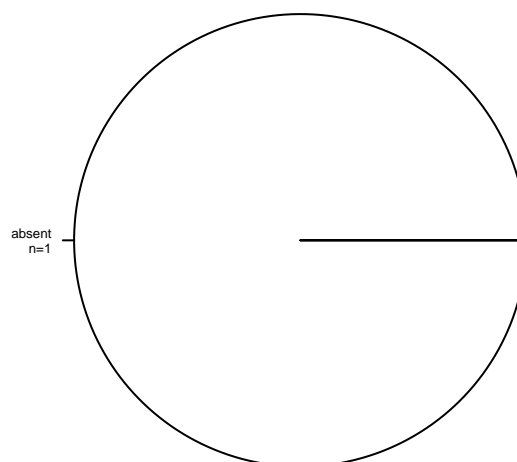
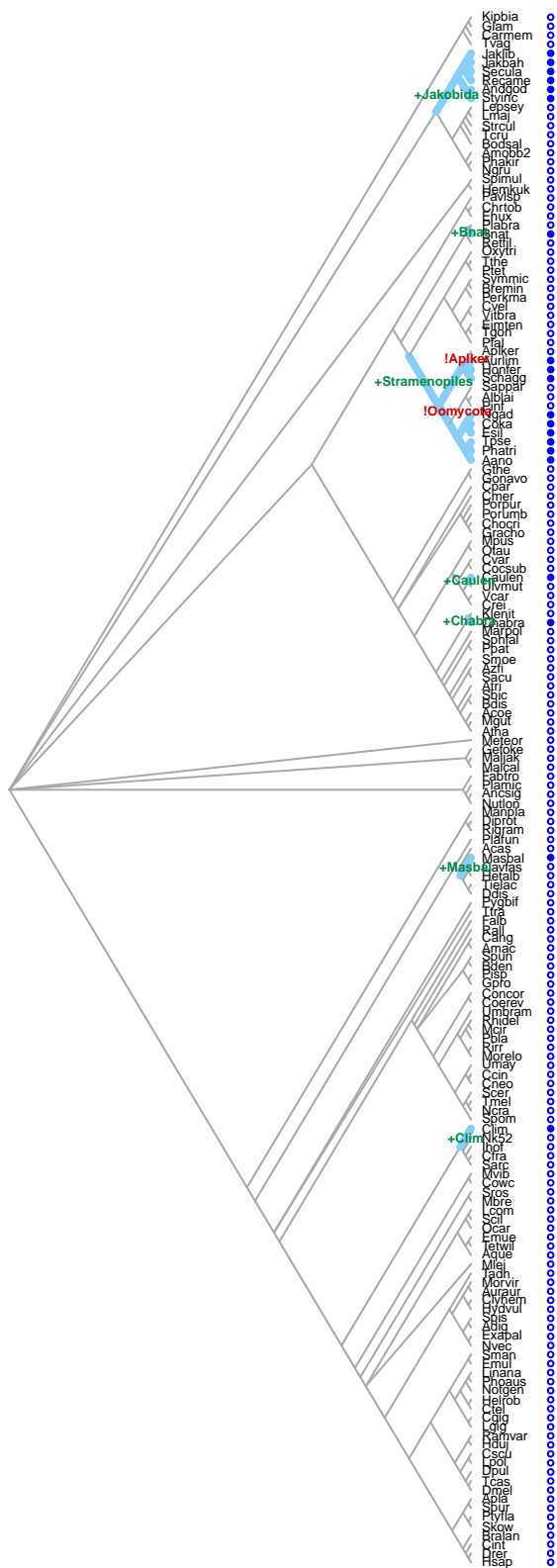


absent absent | n=1

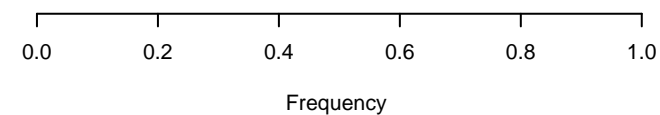


Gain: Porumb,Hemkuk,NA
Presence Eukaryota prob = 0.01
Present: 4
Losses: NA

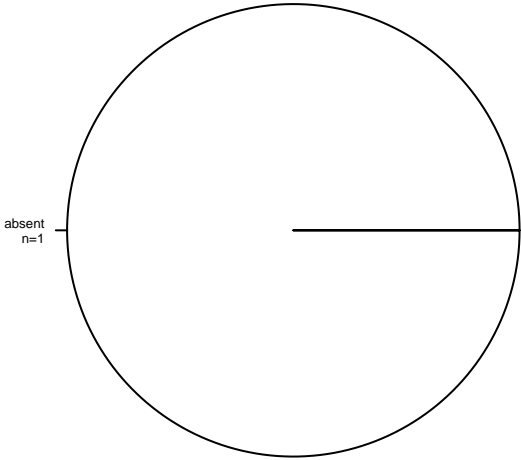
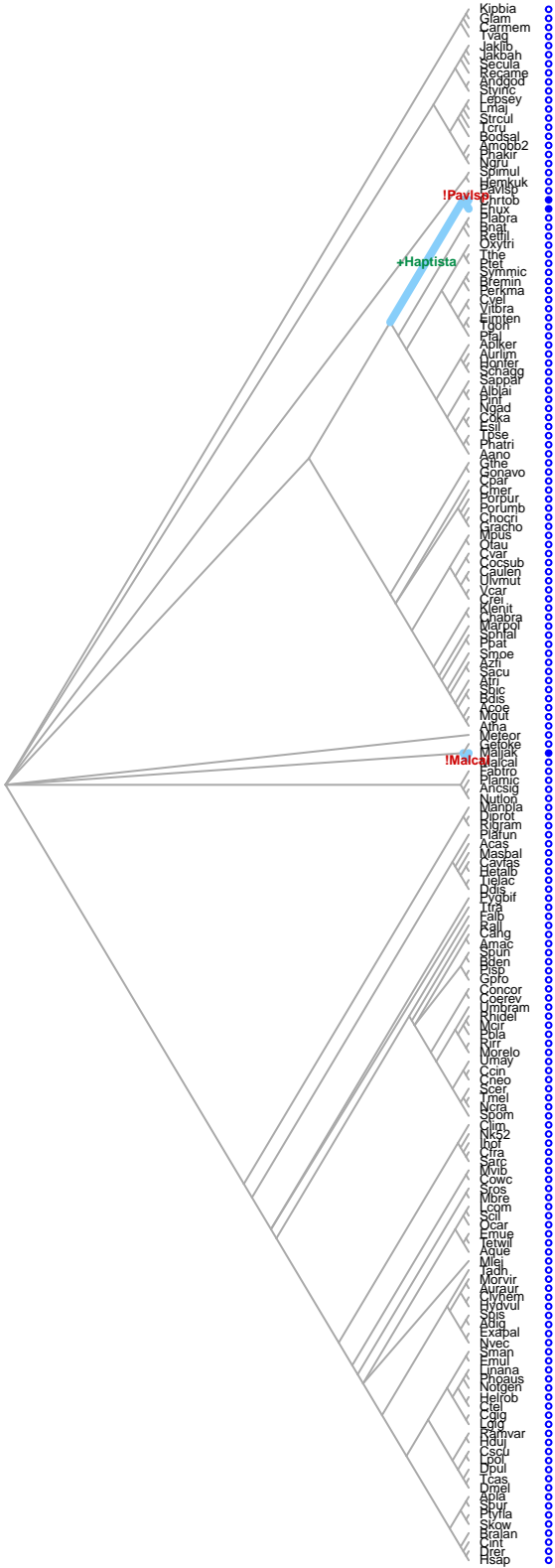
NAP.HG1.5
like·NAP1| 1/NAP1| 2/NAP1| 3/NAP1| 4/NAP1| 5



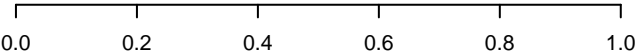
absent absent | n=1



Gain: Clm,Masbal,Chabra,Caulen,Stramenopiles,Bnat,Jakobida,NA
 Presence Eukaryota prob = 0.02
 Present: 20
 Losses: NA

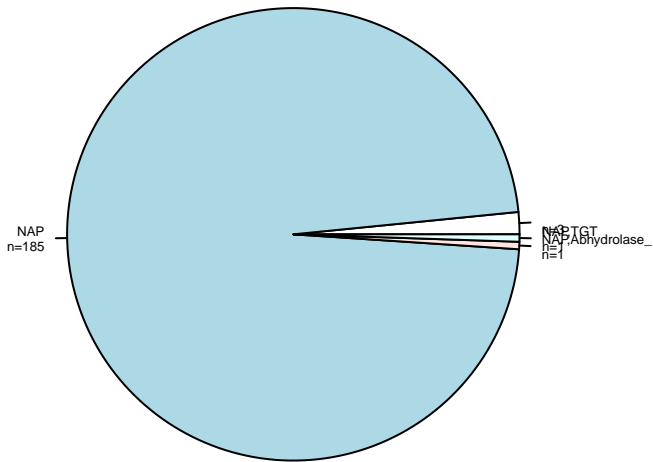
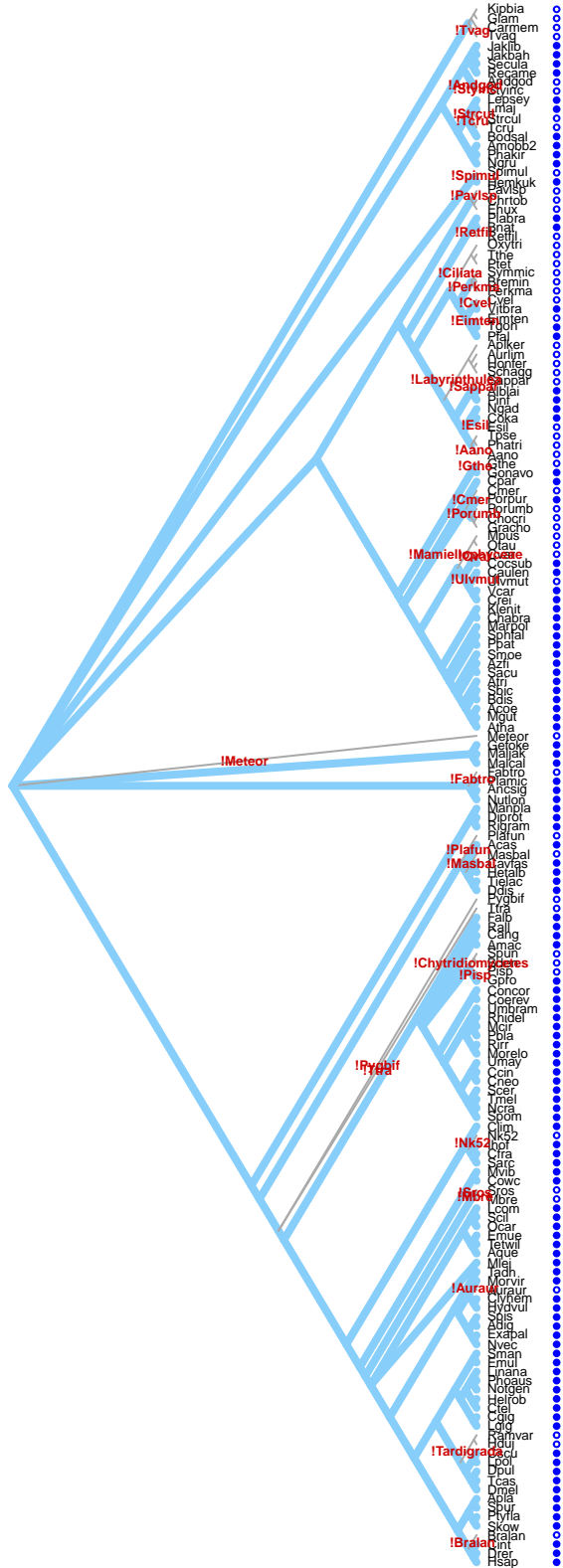


absent absent | n=1



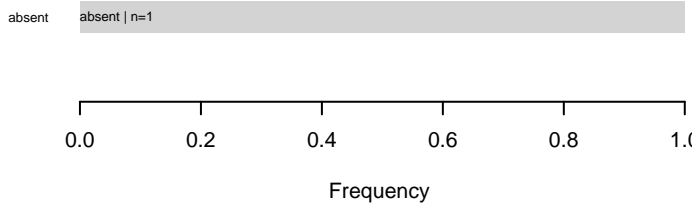
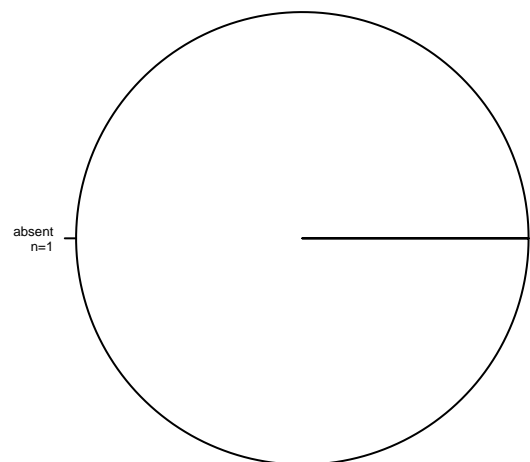
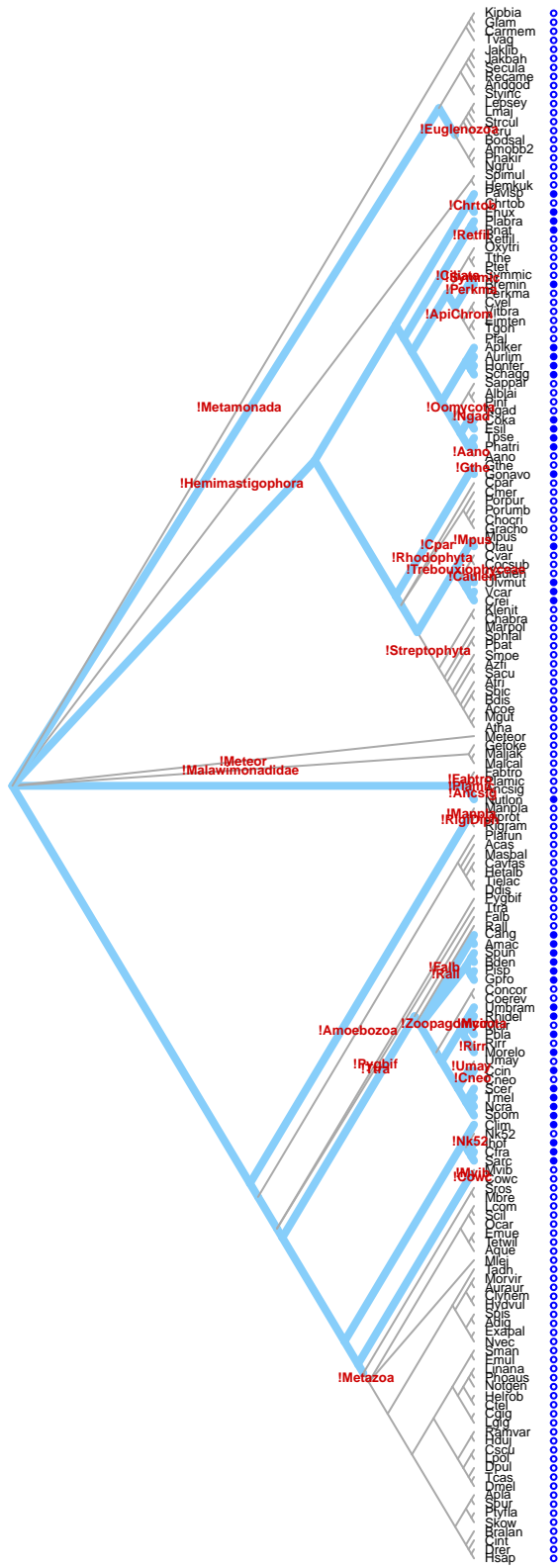
Gain: Haptista,NA
Presence Eukaryota prob = 0.01
Present: 3
Losses: NA

Frequency

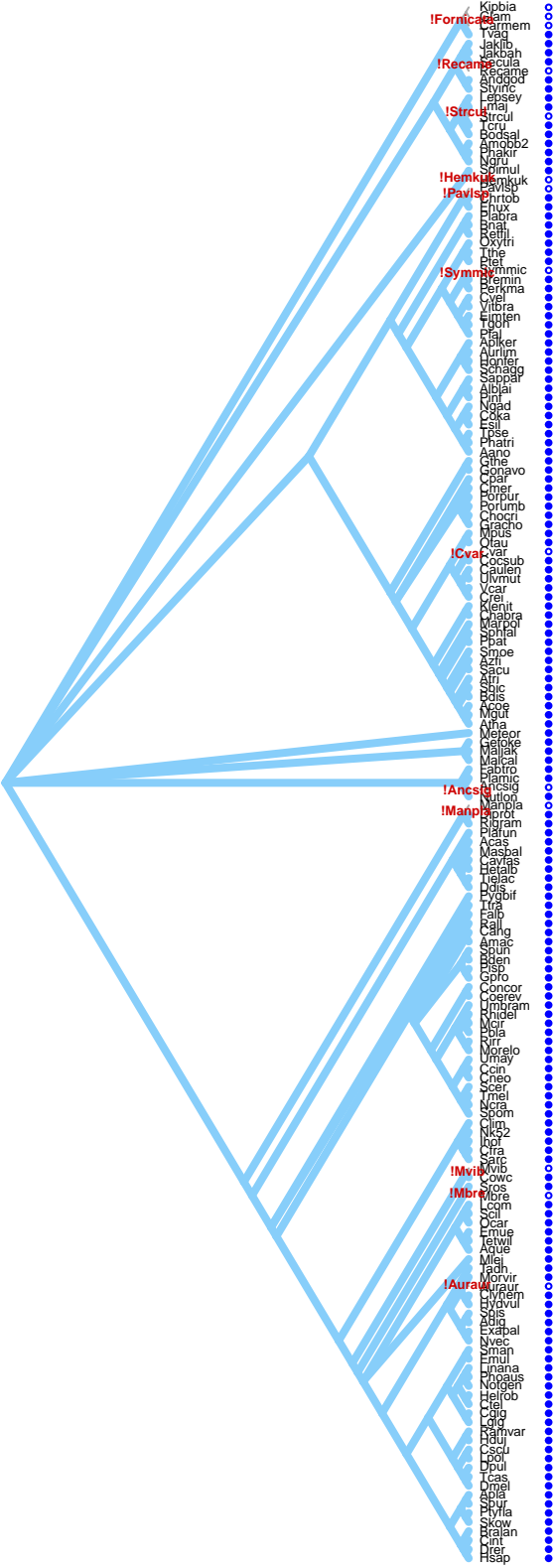


Gain: NA
Presence Eukaryota prob = 1.00
Present: 117
Losses: NA

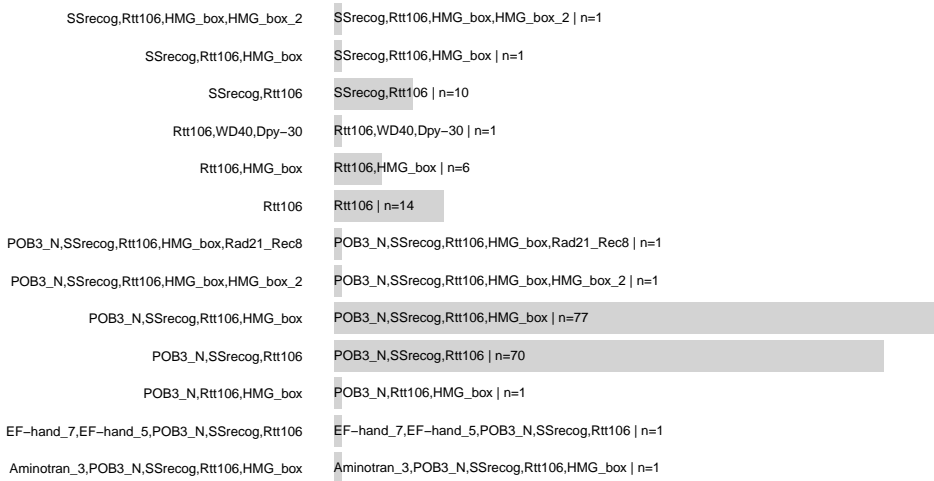
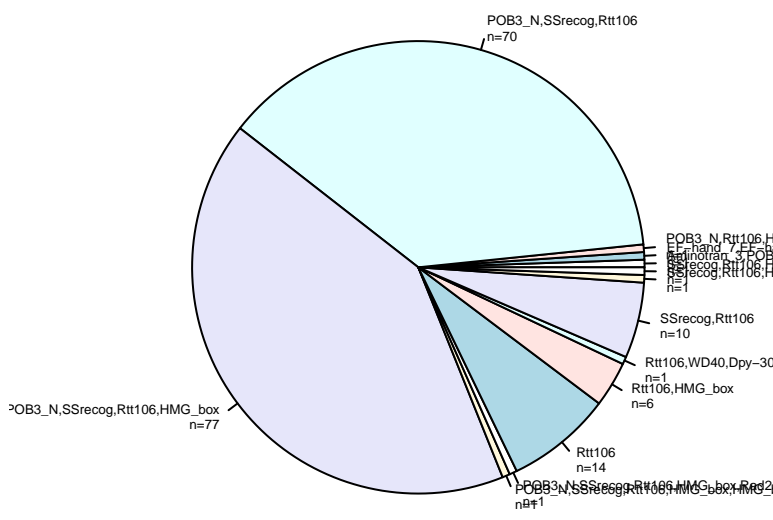
Rtt106.HG1.0
like:SSRP1



Rtt106.HG1.1
SSRP1

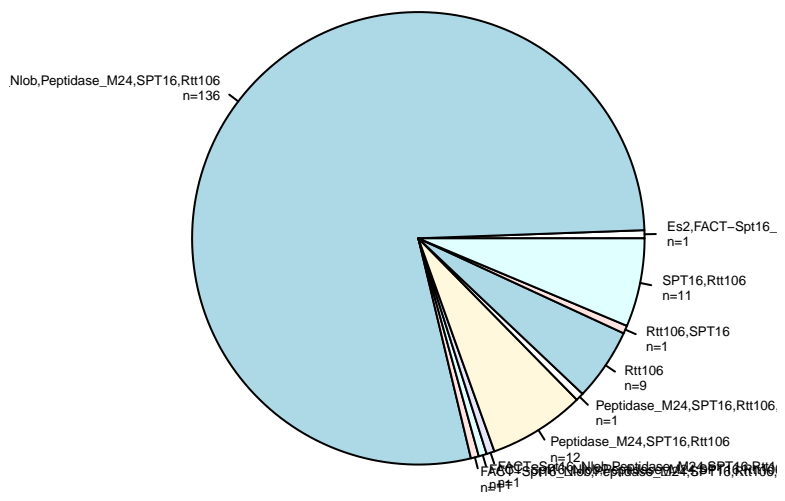
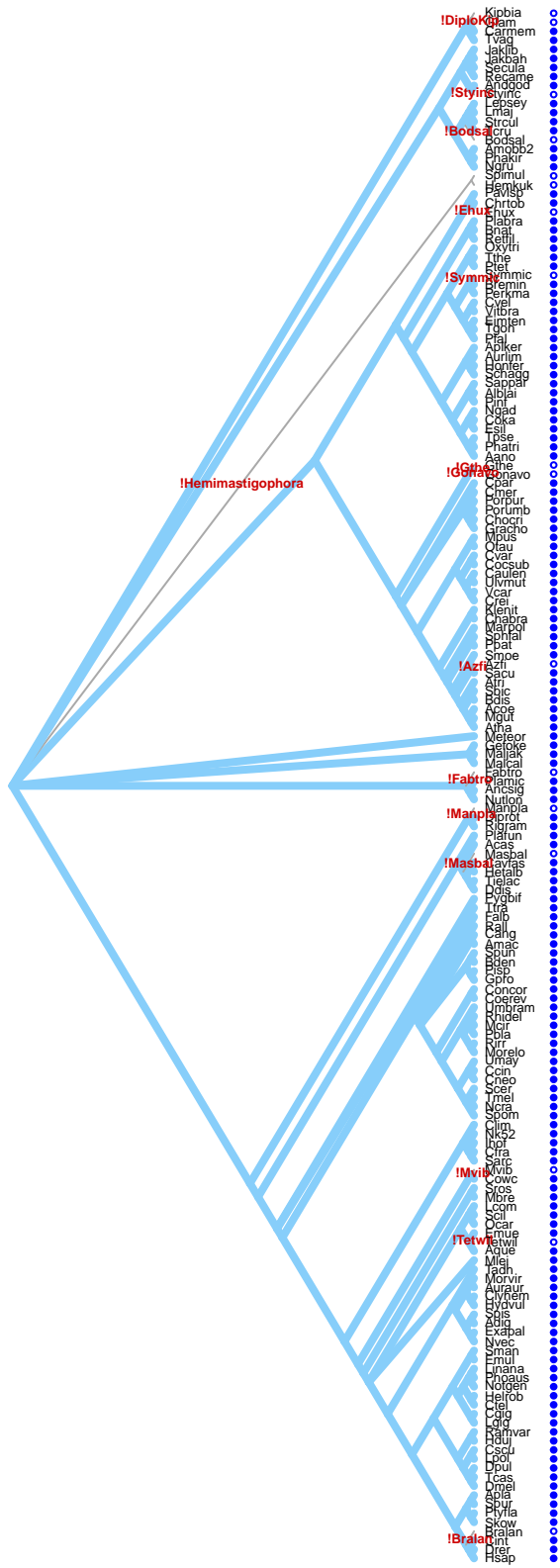


Gain: NA
Presence Eukaryota prob = 1.00
Present: 158
Losses: NA

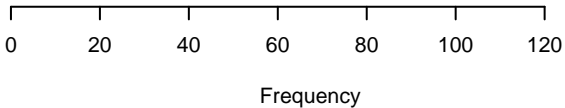


0 10 20 30 40 50 60 70
Frequency

Rtt106.HG2.0
SUPT16H

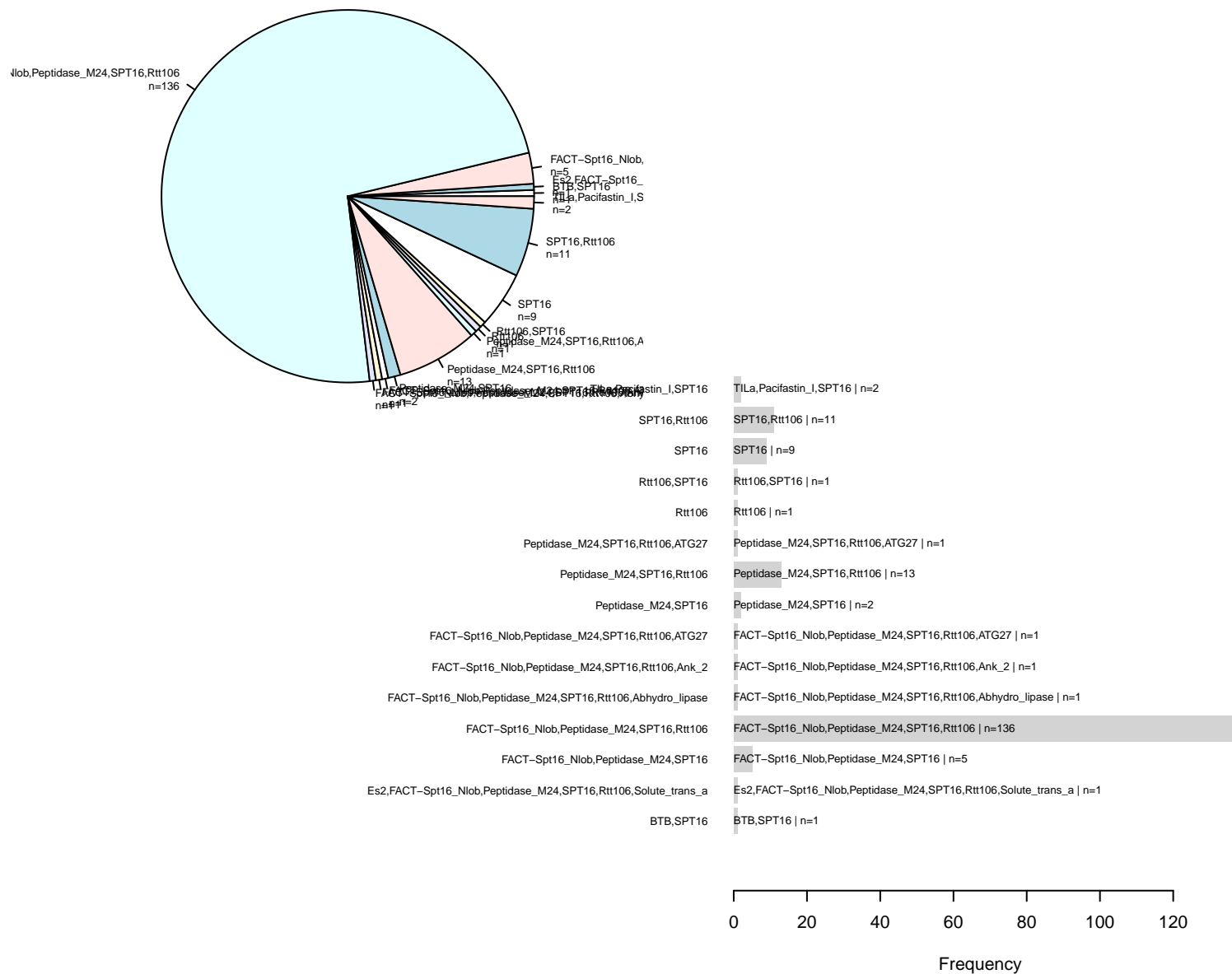


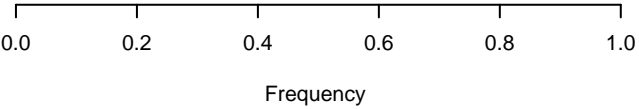
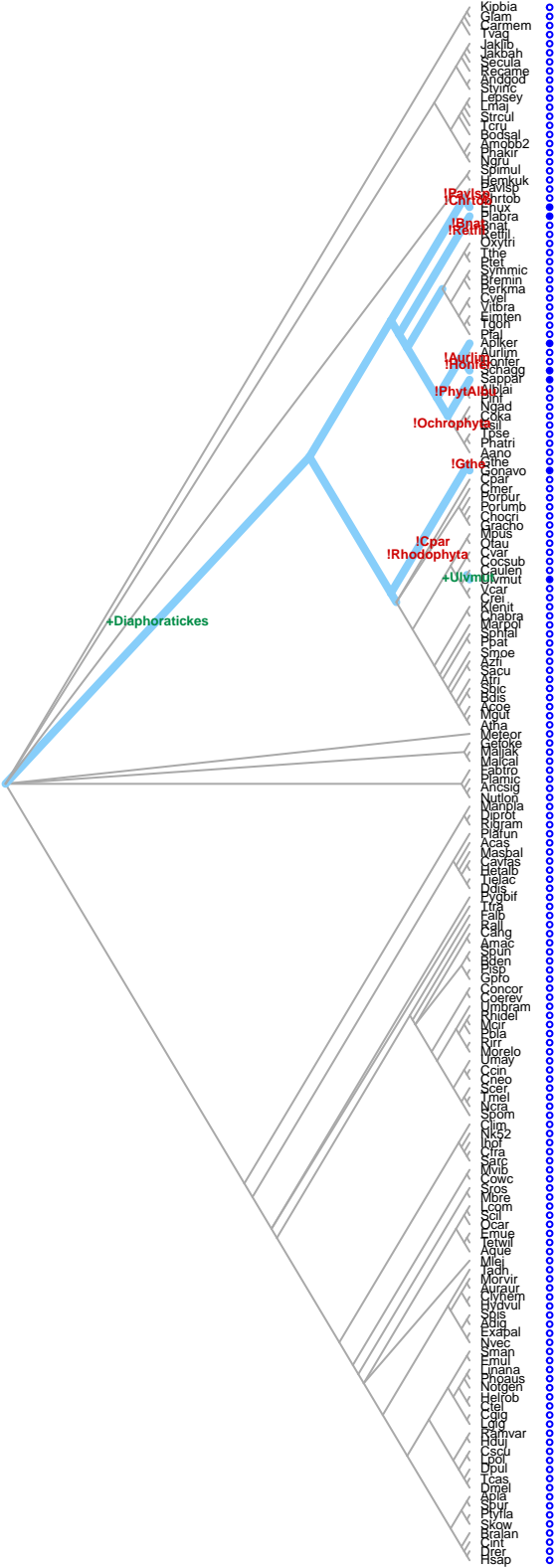
SPT16,Rtt106	SPT16,Rtt106 n=11
Rtt106,SPT16	Rtt106,SPT16 n=1
Rtt106	Rtt106 n=9
Peptidase_M24,SPT16,Rtt106,ATG27	Peptidase_M24,SPT16,Rtt106,ATG27 n=1
Peptidase_M24,SPT16,Rtt106	Peptidase_M24,SPT16,Rtt106 n=12
FACT-Spt16_Nlob,Peptidase_M24,SPT16,Rtt106,ATG27	FACT-Spt16_Nlob,Peptidase_M24,SPT16,Rtt106,ATG27 n=1
FACT-Spt16_Nlob,Peptidase_M24,SPT16,Rtt106,Ank_2	FACT-Spt16_Nlob,Peptidase_M24,SPT16,Rtt106,Ank_2 n=1
FACT-Spt16_Nlob,Peptidase_M24,SPT16,Rtt106,Abhydro_lipase	FACT-Spt16_Nlob,Peptidase_M24,SPT16,Rtt106,Abhydro_lipase n=1
FACT-Spt16_Nlob,Peptidase_M24,SPT16,Rtt106	FACT-Spt16_Nlob,Peptidase_M24,SPT16,Rtt106 n=136
Es2,FACT-Spt16_Nlob,Peptidase_M24,SPT16,Rtt106,Solute_trans_a	Es2,FACT-Spt16_Nlob,Peptidase_M24,SPT16,Rtt106,Solute_trans_a n=1



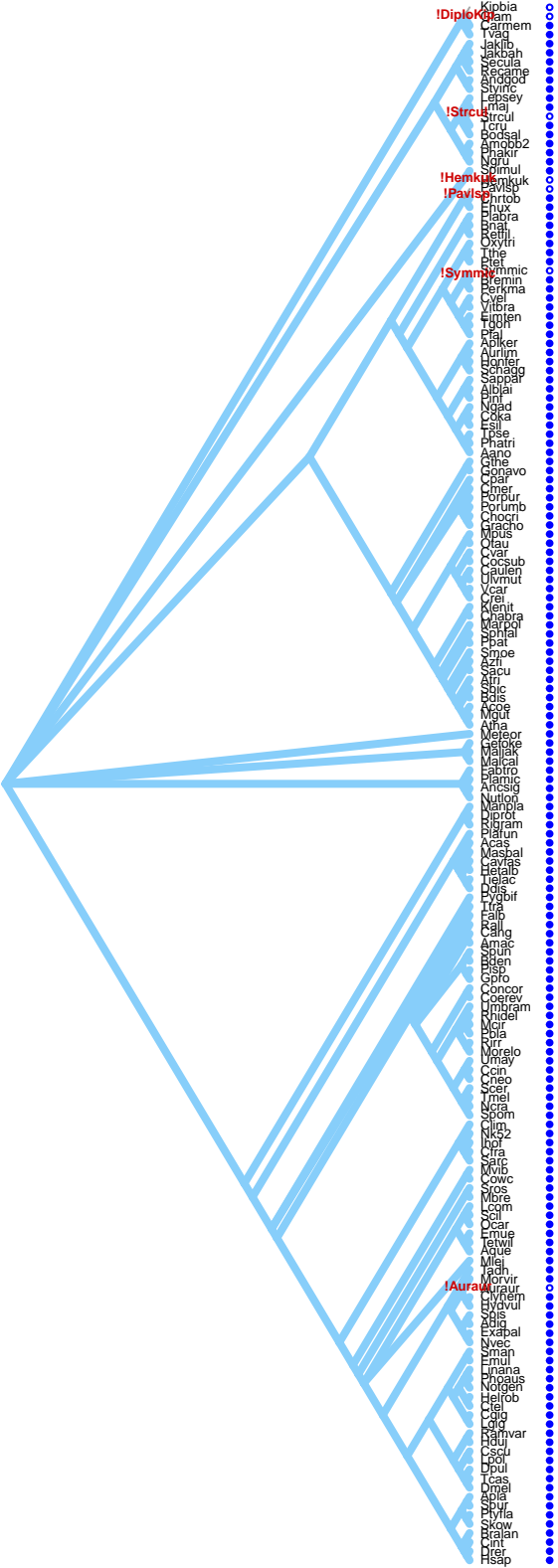
Gain: NA
Presence Eukaryota prob = 1.00
Present: 155
Losses: NA

Gain: NA
Presence Eukaryota prob = 1.00
Present: 160
Losses: NA

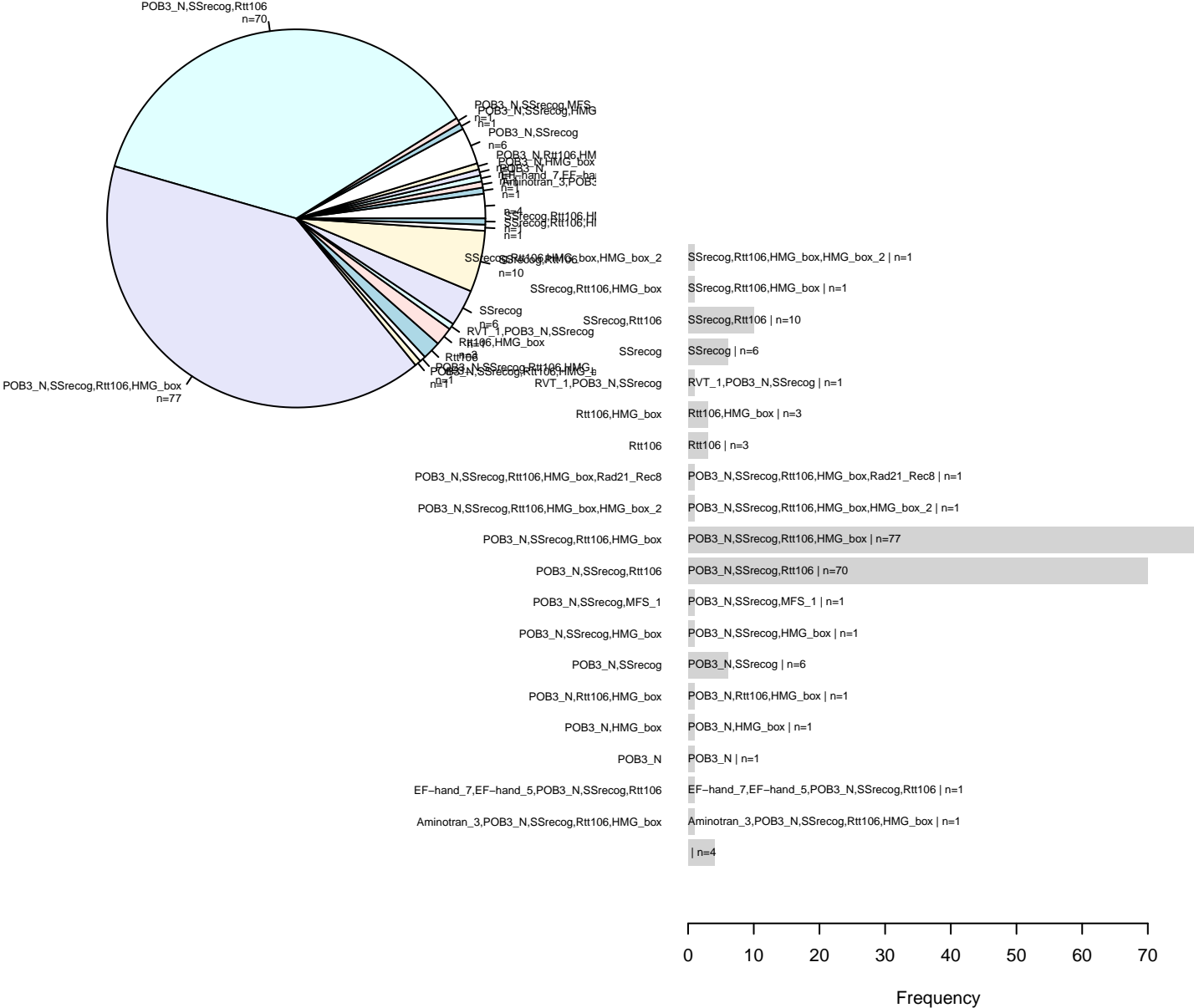


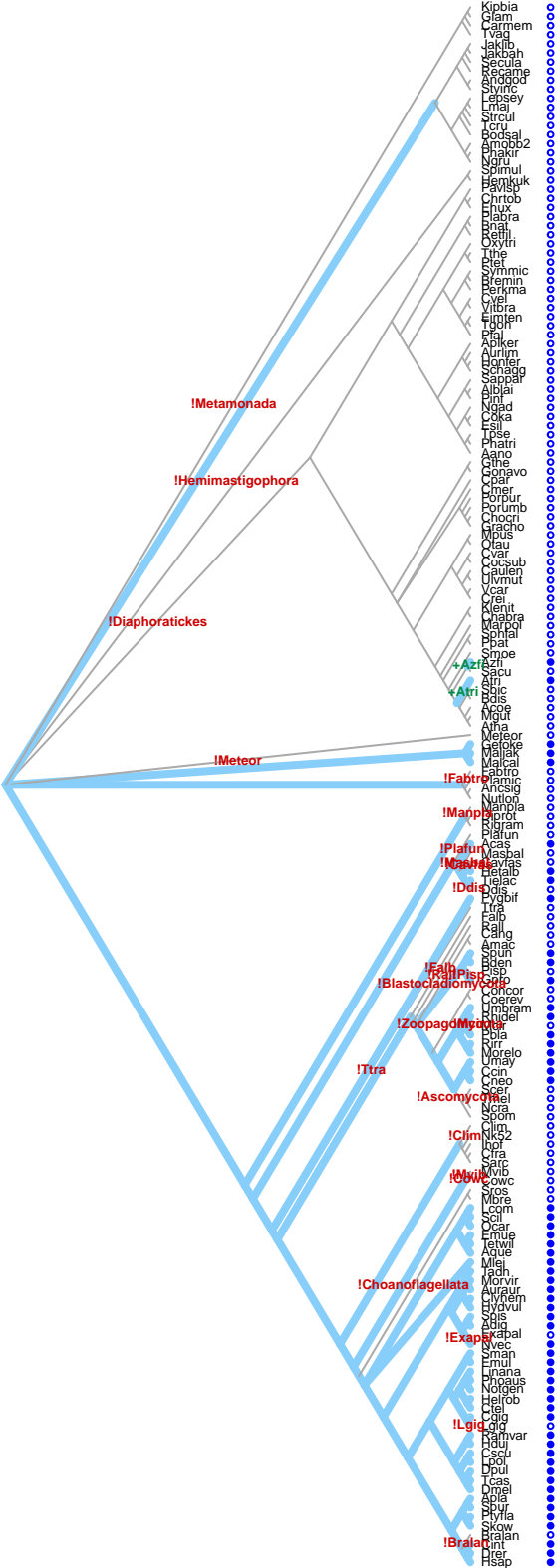


SSrecog.HG1.1
SSRP1

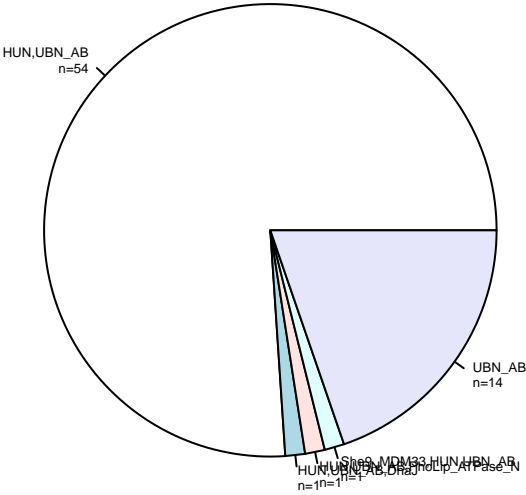


Gain: NA
Presence Eukaryota prob = 1.00
Present: 165
Losses: NA

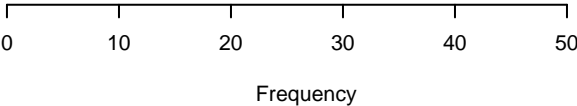




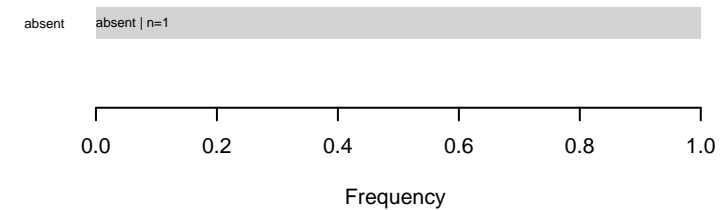
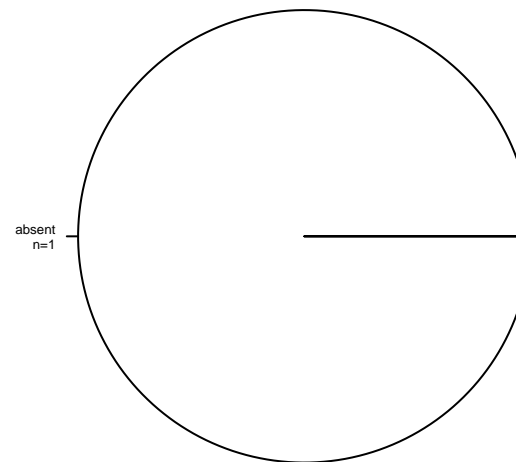
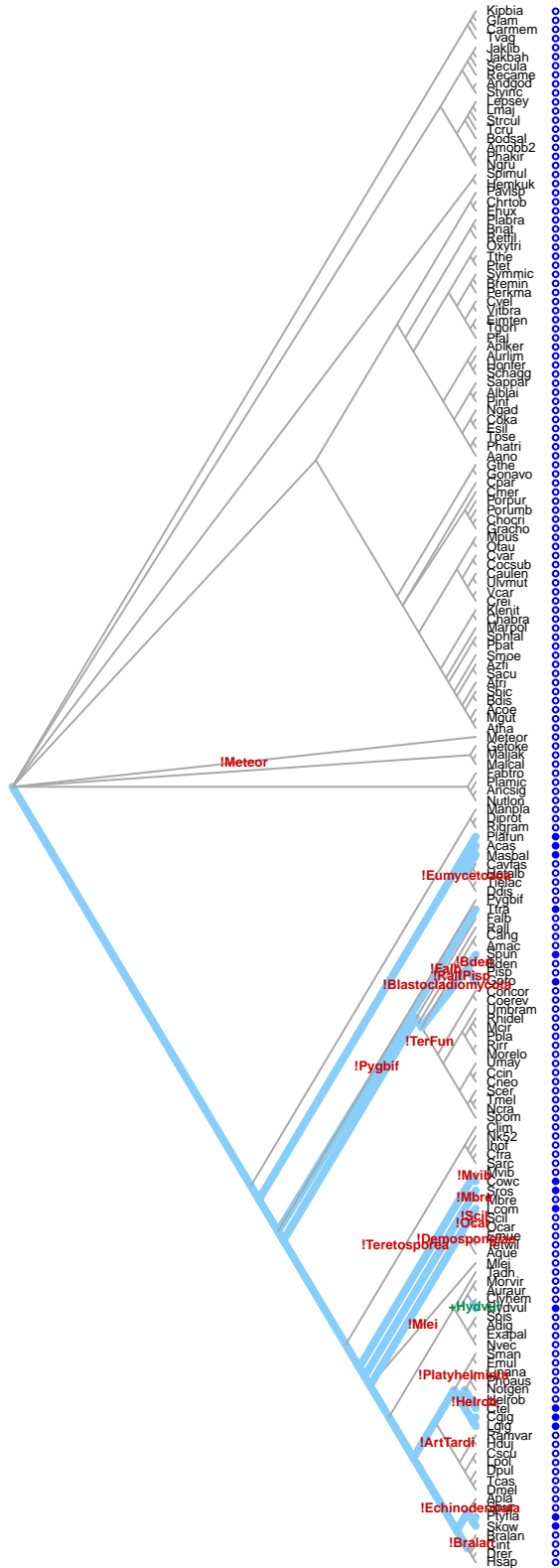
Gain: Atri,Azfi,NA
Presence Eukaryota prob = 0.99
Present: 57
Losses: NA



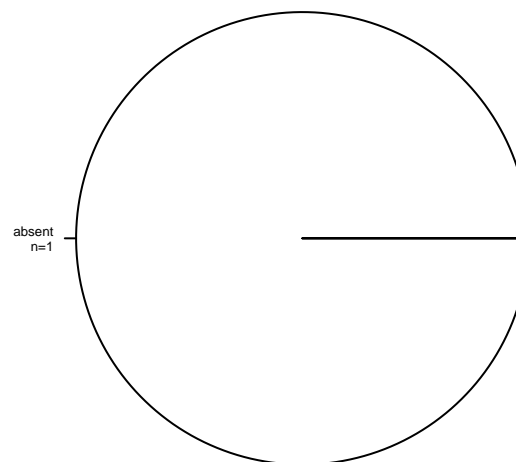
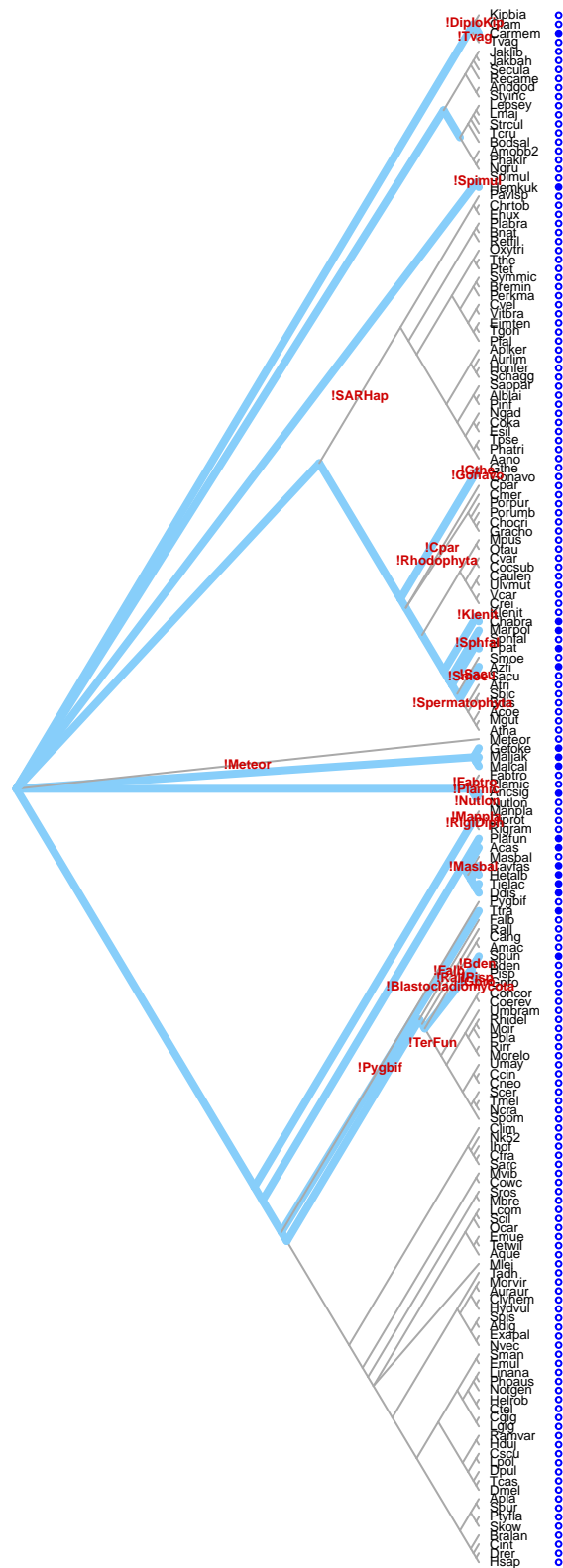
UBN_AB	UBN_AB n=14
She9_MDM33,HUN,UBN_AB	She9_MDM33,HUN,UBN_AB n=1
HUN,UBN_AB,PhoLip_ATPase_N	HUN,UBN_AB,PhoLip_ATPase_N n=1
HUN,UBN_AB,DnaJ	HUN,UBN_AB,DnaJ n=1
HUN,UBN_AB	HUN,UBN_AB n=54



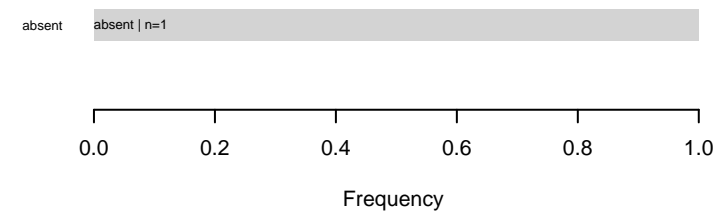
zf-CCHH.HG1.0
like:CHFR



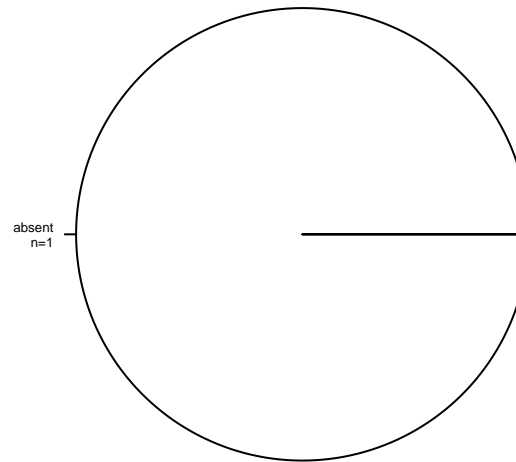
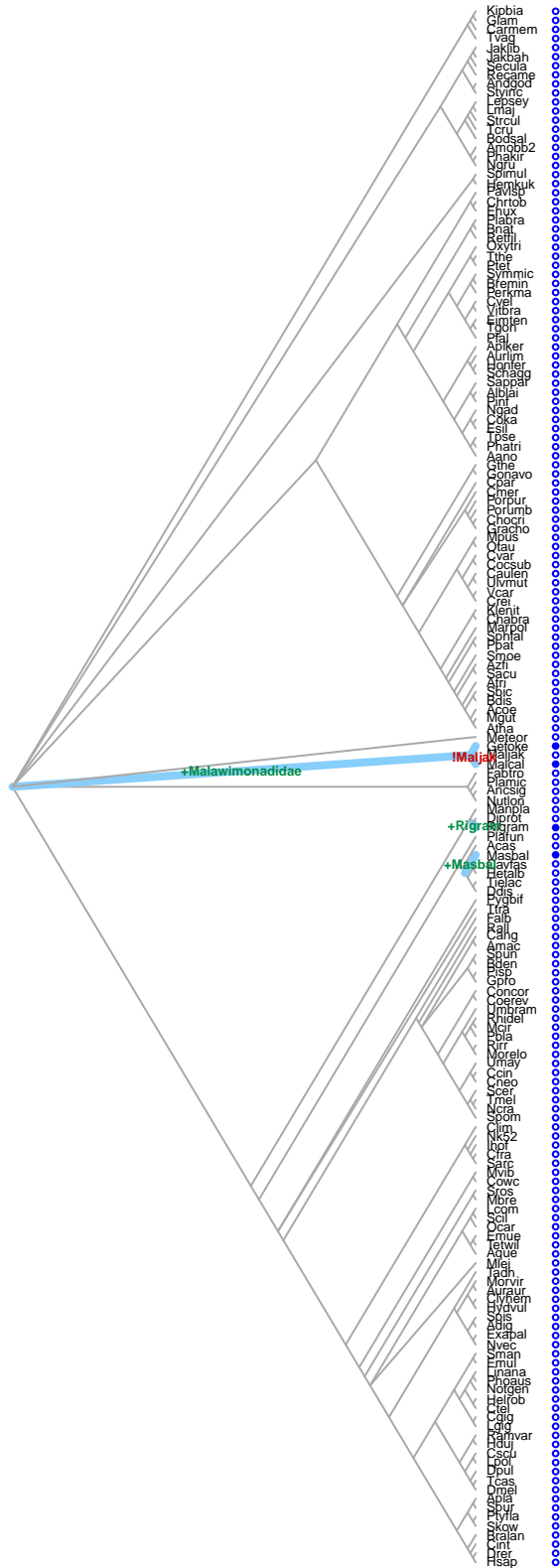
zf-CCHH.HG1.1
like:APLF



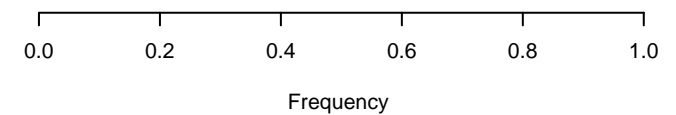
Gain: NA
Presence Eukaryota prob = 1.00
Present: 18
Losses: NA



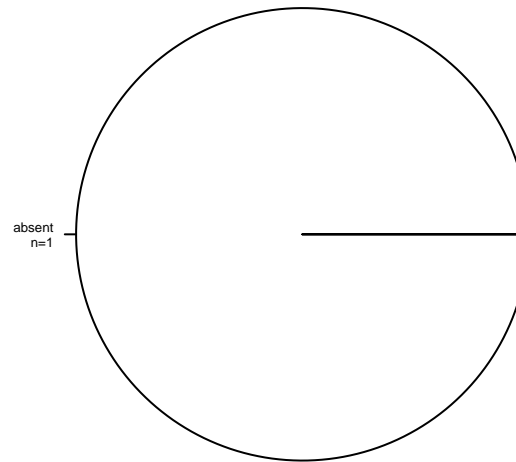
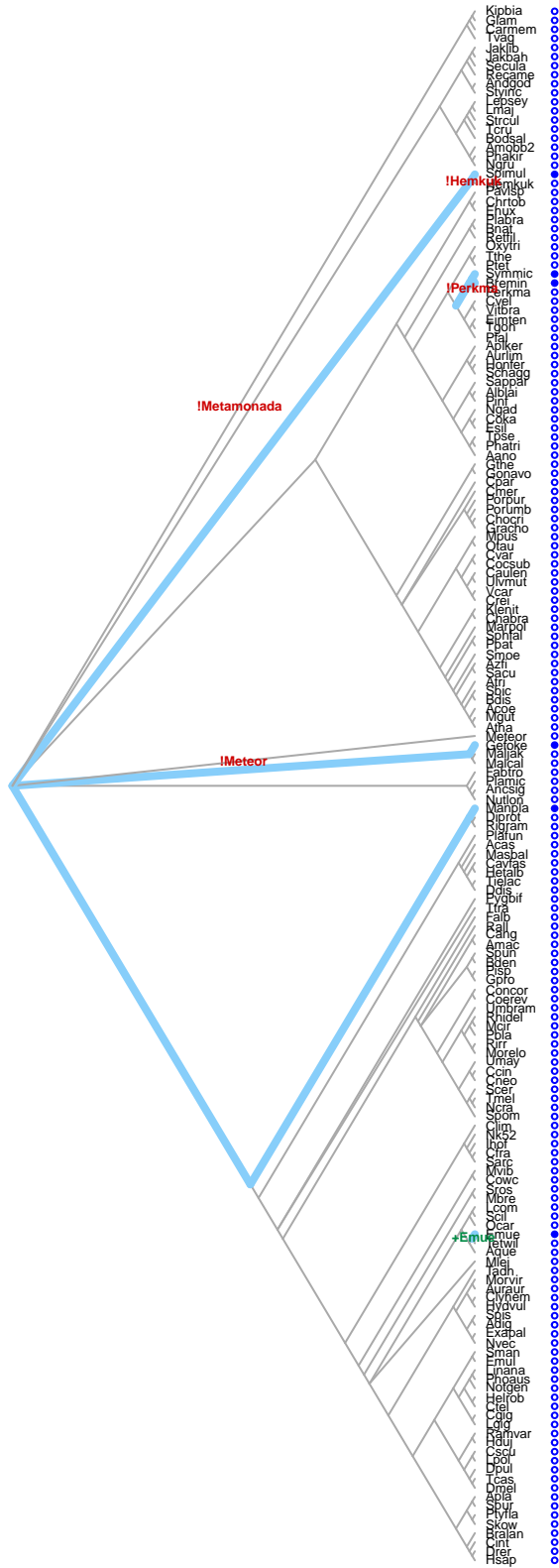
zf-CCHH.HG1.10
like:APLF



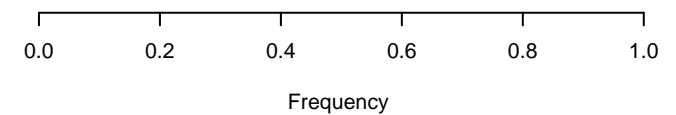
absent absent | n=1



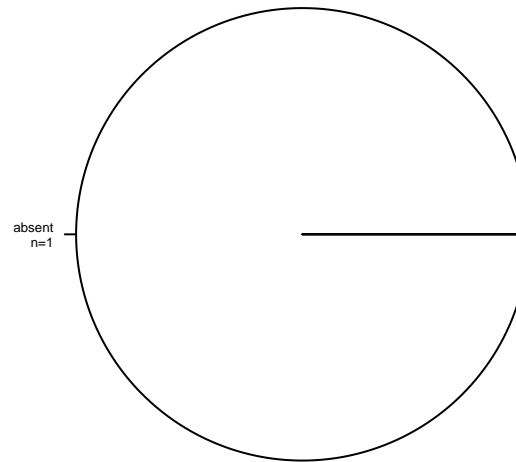
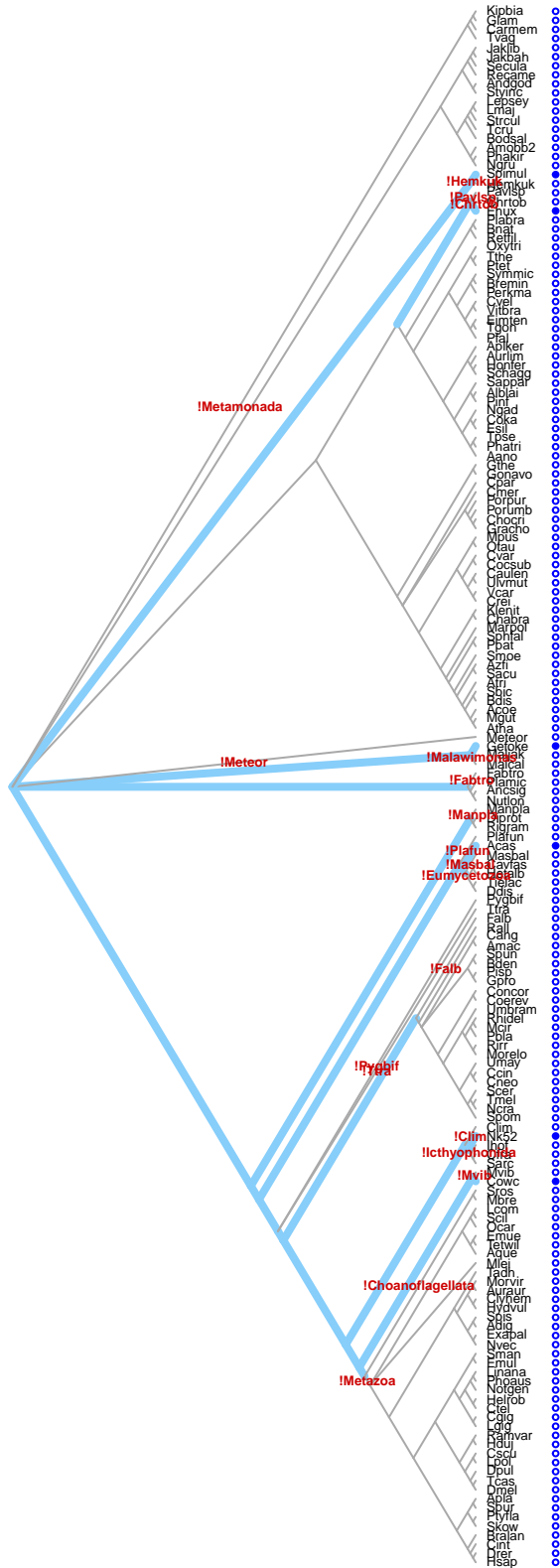
zf-CCHH.HG1.11
like:APLF



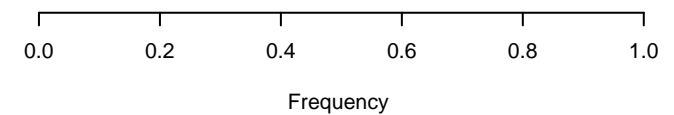
absent absent | n=1



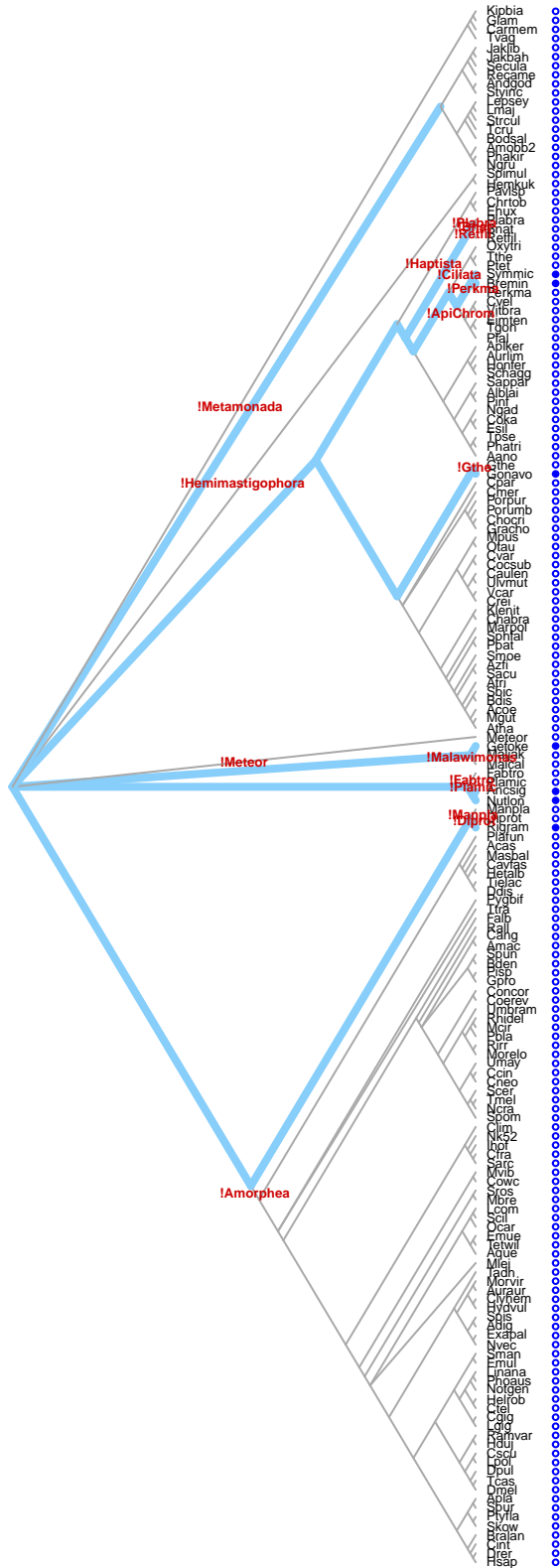
zf-CCHH.HG1.12
like:APLF



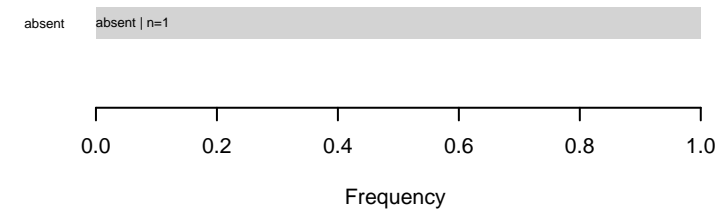
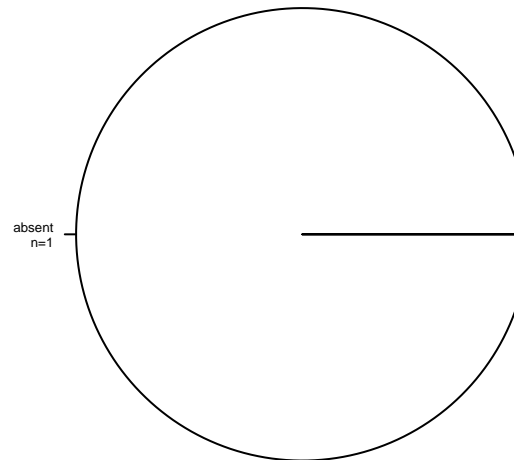
absent absent | n=1



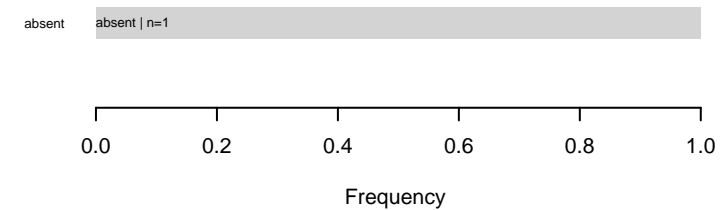
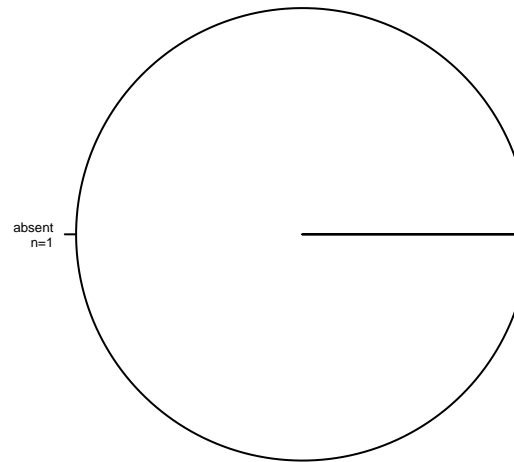
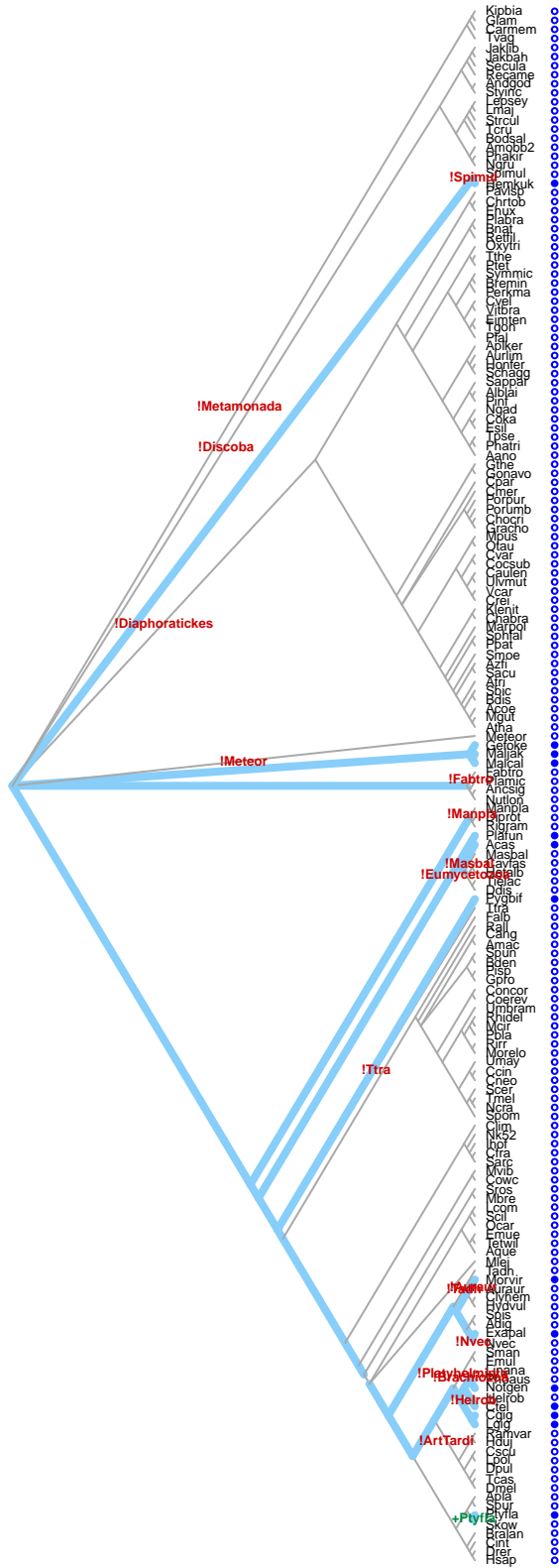
zf-CCHH.HG1.13
like:APLF



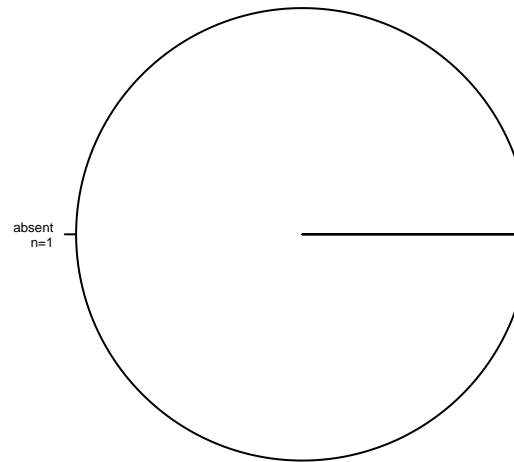
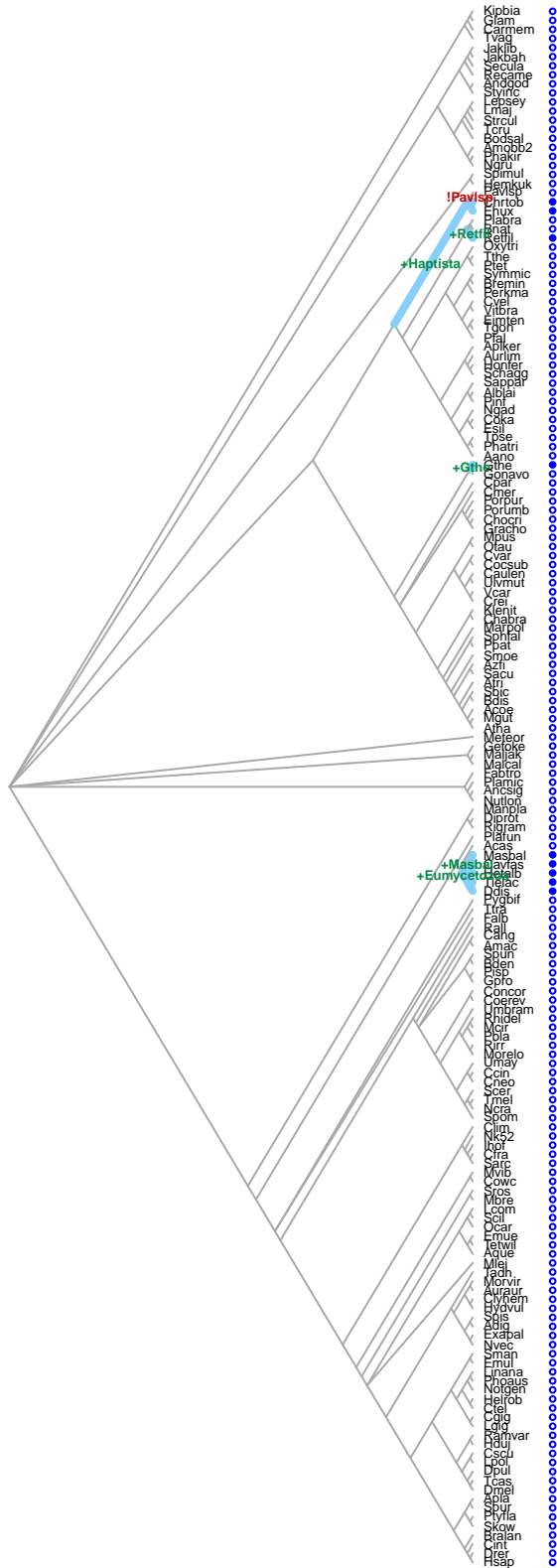
Gain: NA
Presence Eukaryota prob = 0.95
Present: 7
Losses: NA



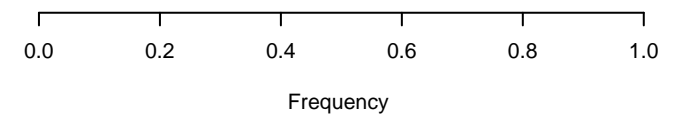
zf-CCHH.HG1.14
like:APLF



zf-CCHH.HG1.15
like:APLF

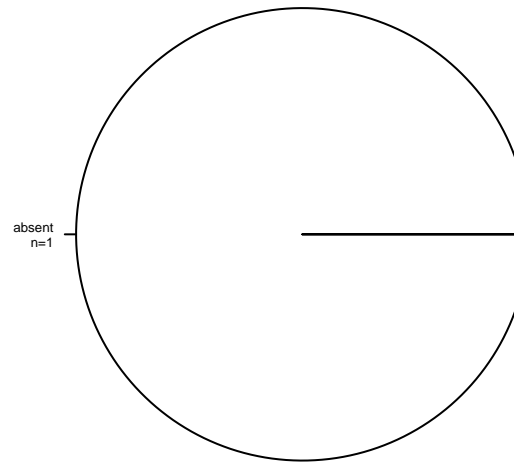
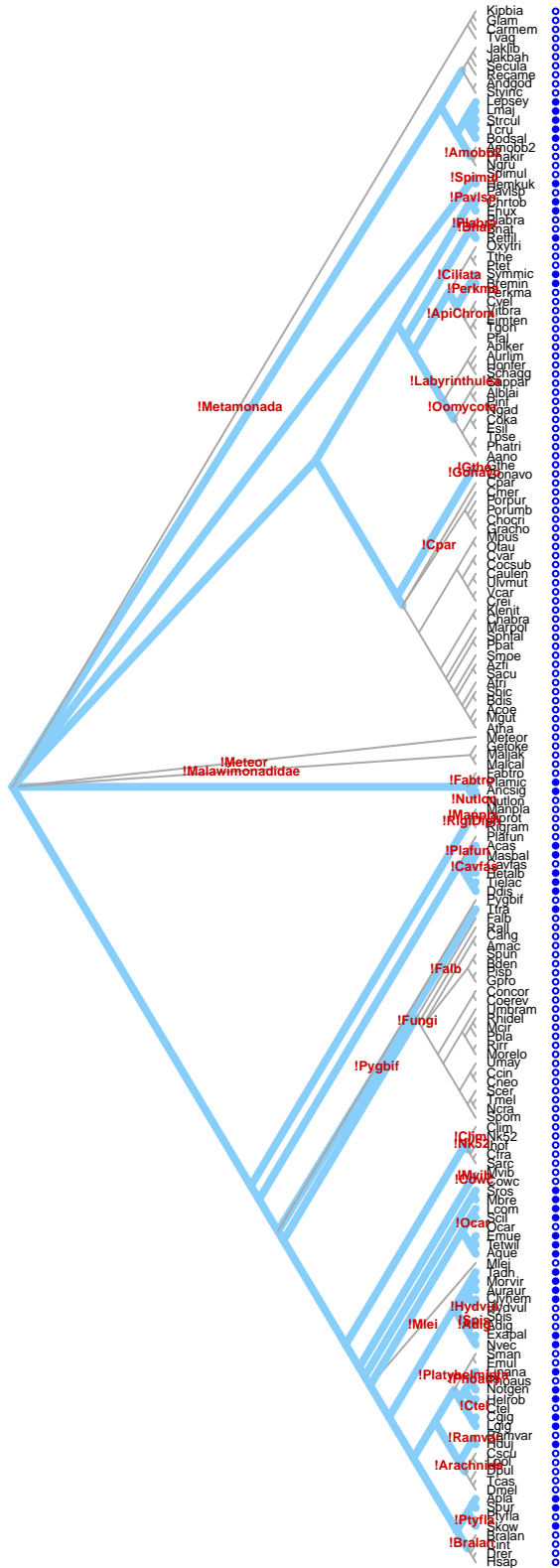


absent absent | n=1

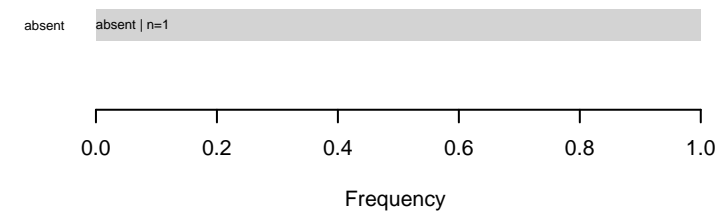


Gain: Eumycetozoa,Masbal,Gthe,Retfil,Haptista,NA
 Presence Eukaryota prob = 0.12
 Present: 9
 Losses: NA

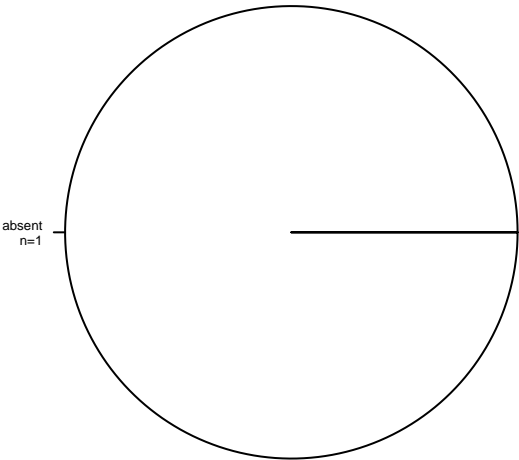
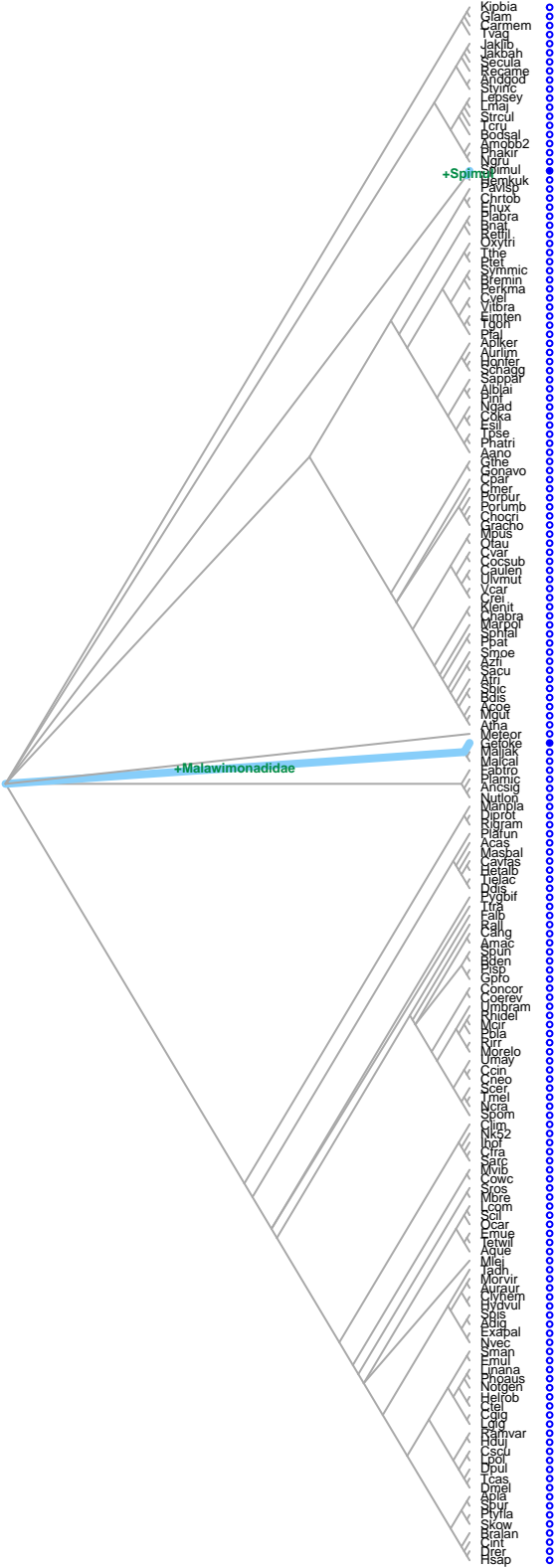
zf-CCHH.HG1.16
like:APLF



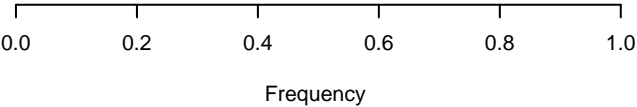
Gain: NA
Presence Eukaryota prob = 1.00
Present: 41
Losses: NA



zf-CCHH.HG1.17
like:APLF

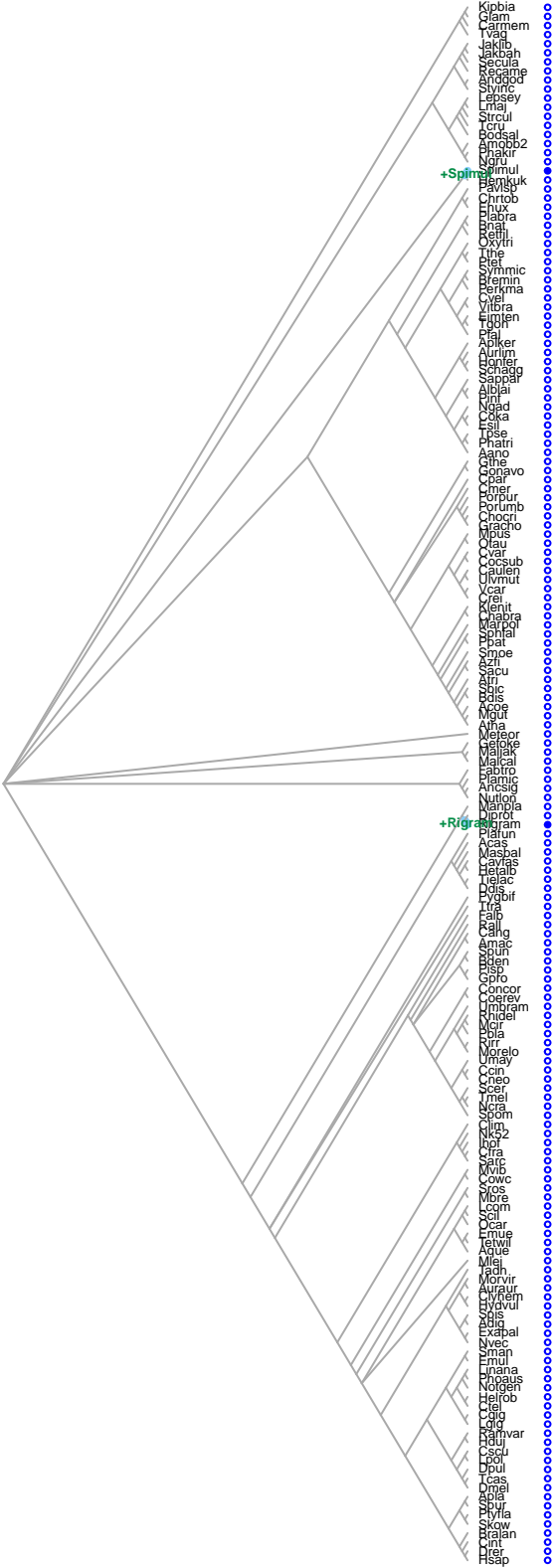


absent absent | n=1

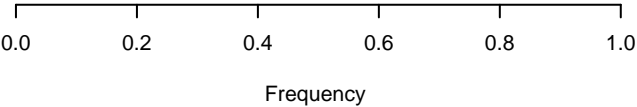


Gain: Malawimonadidae, Spimul, NA
Presence Eukaryota prob = 0.12
Present: 2
Losses: NA

zf-CCHH.HG1.18
like:APLF

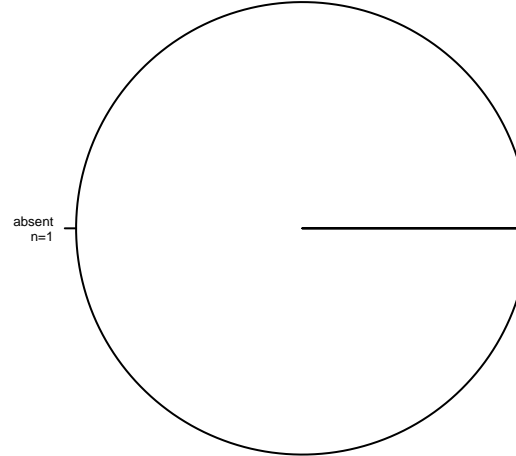
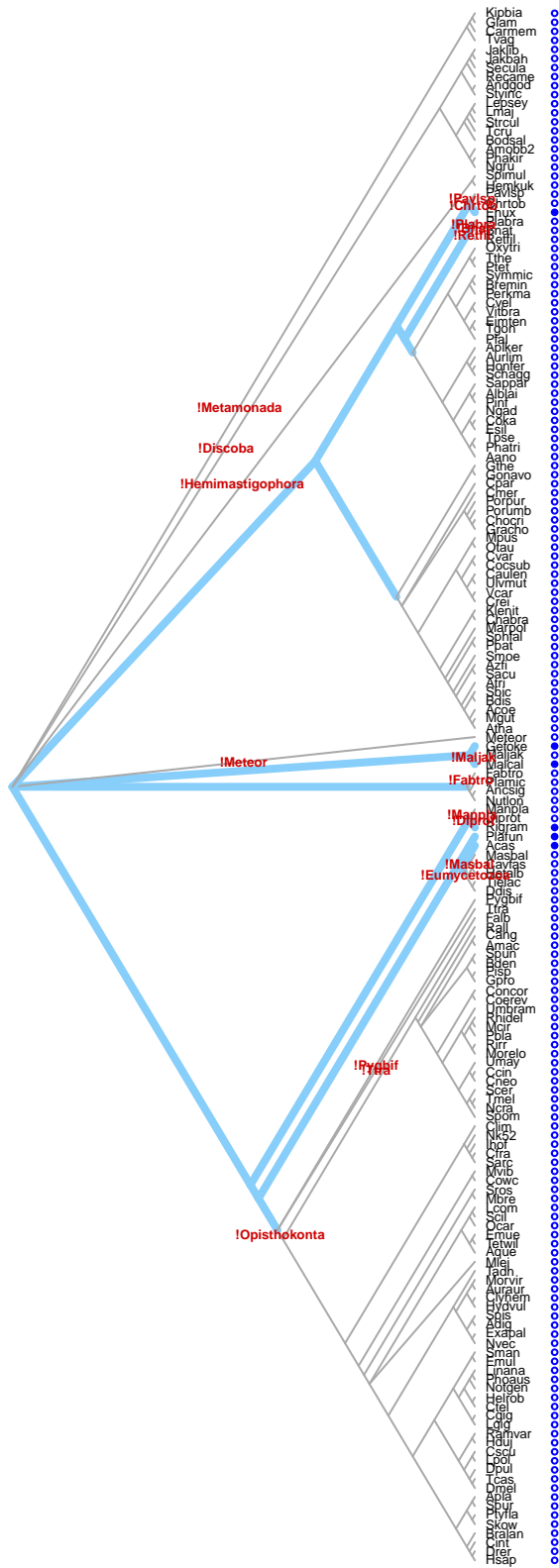


absent absent | n=1

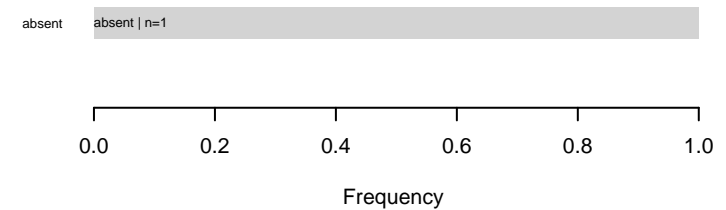


Gain: Rigram,Spimul,NA
Presence Eukaryota prob = 0.17
Present: 2
Losses: NA

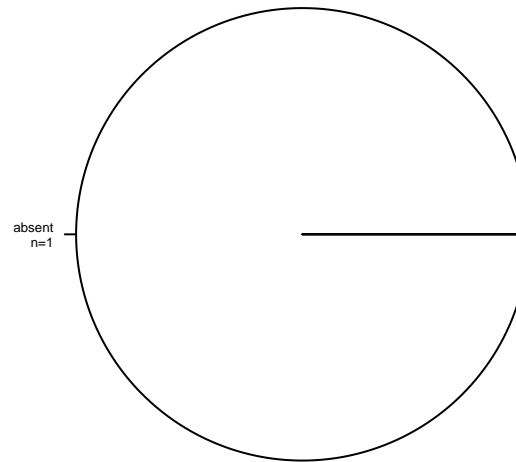
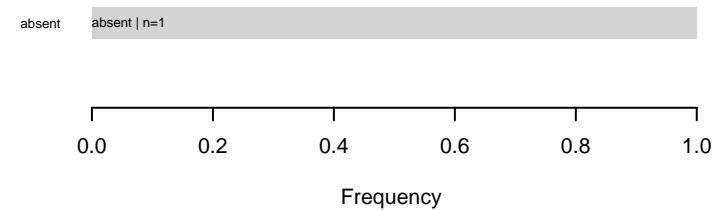
zf-CCHH.HG1.2
like:CHFR



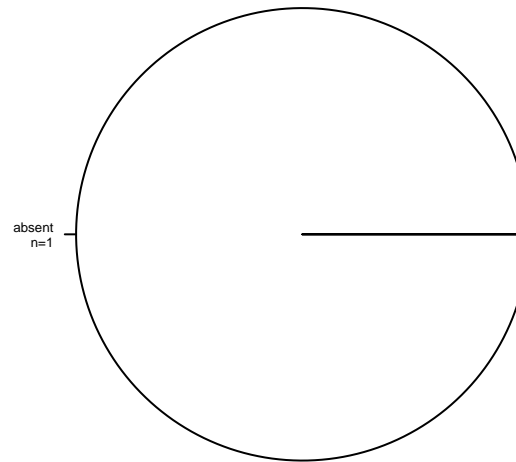
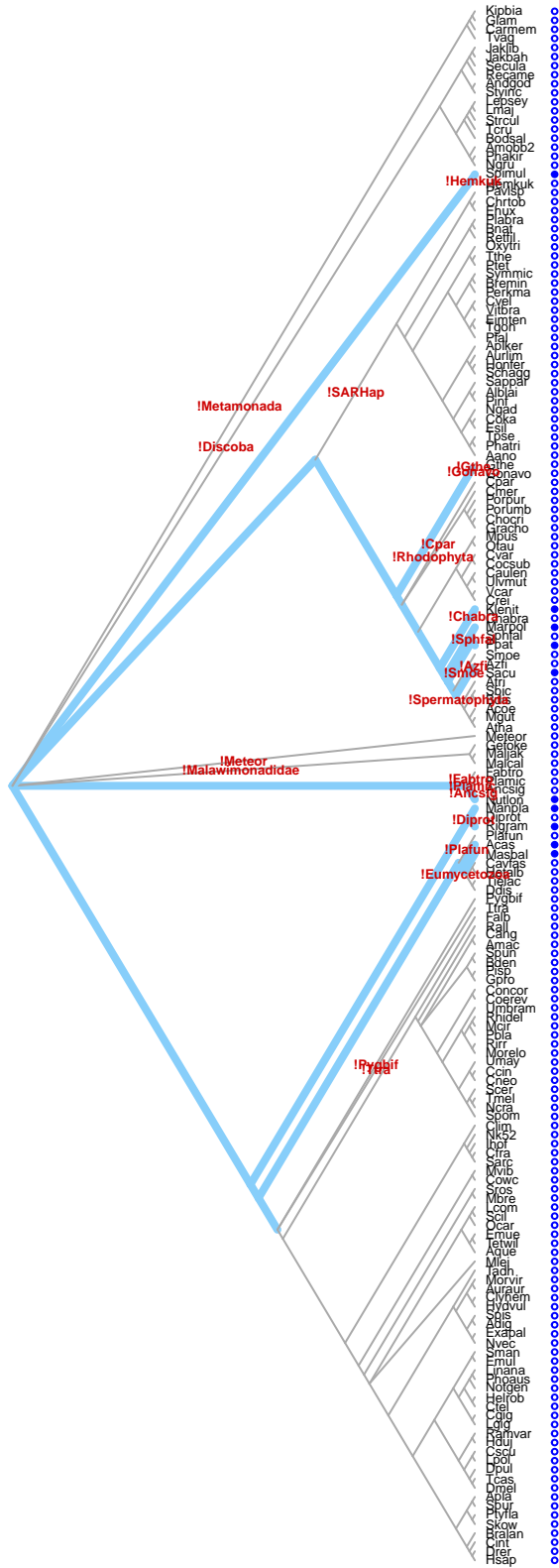
Gain: NA
Presence Eukaryota prob = 0.98
Present: 6
Losses: NA



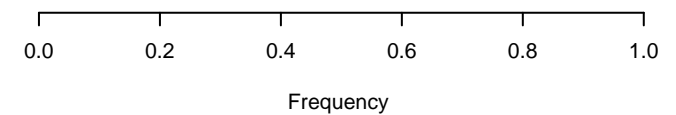
Gain: NA
Presence Eukaryota prob = 0.95
Present: 29
Losses: NA



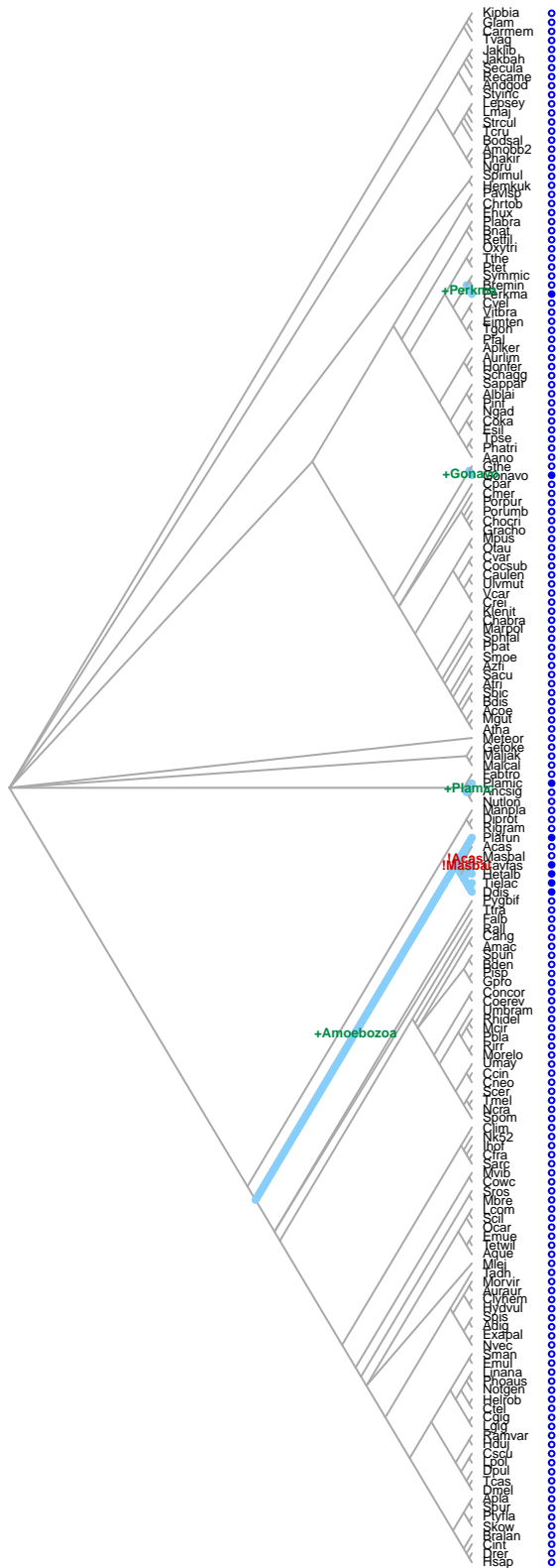
zf-CCHH.HG1.4
like:CHFR



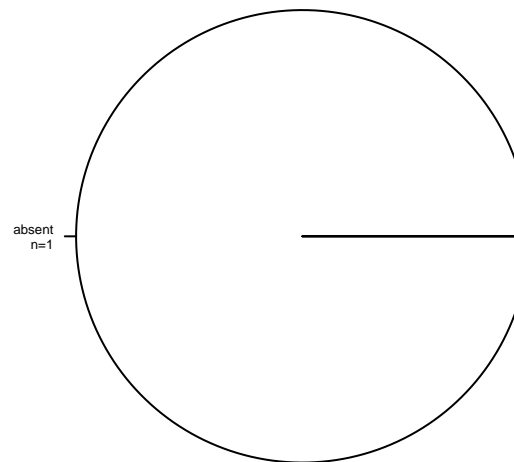
absent absent | n=1



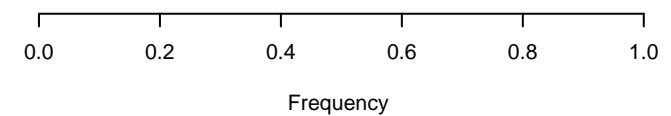
zf-CCHH.HG1.5
like:CHFR



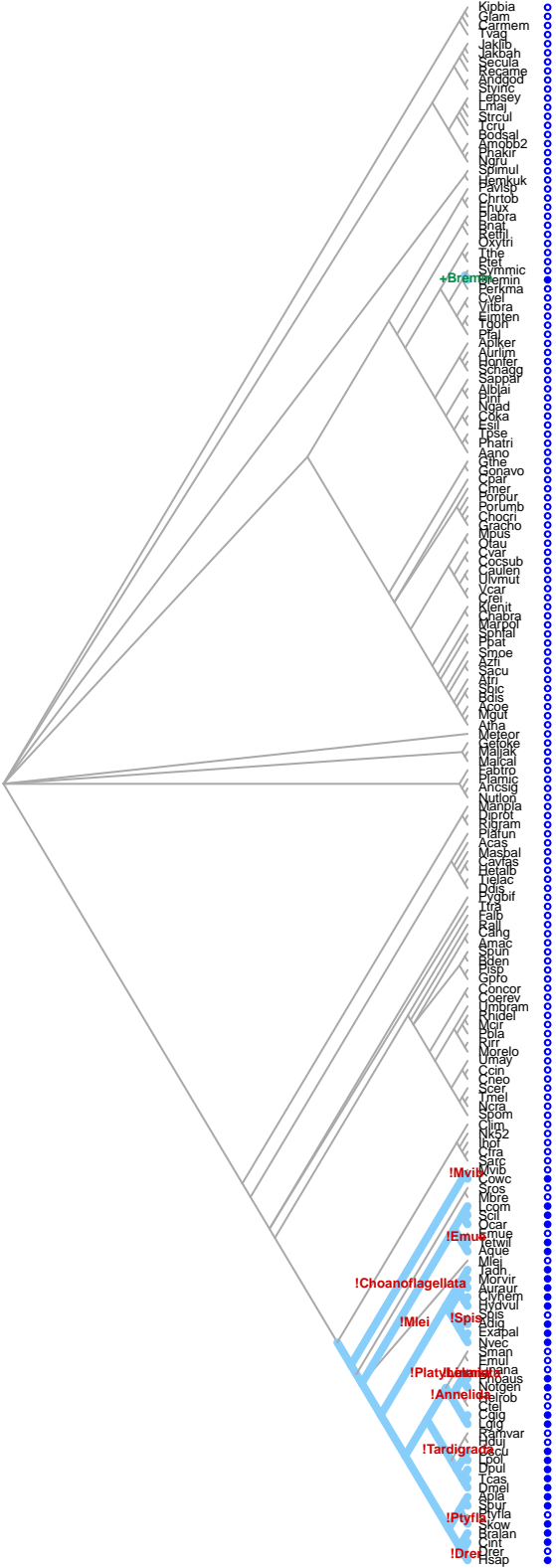
Gain: Amoebozoa,Plamic,Gonavo,Perkma,NA
 Presence Eukaryota prob = 0.13
 Present: 8
 Losses: NA



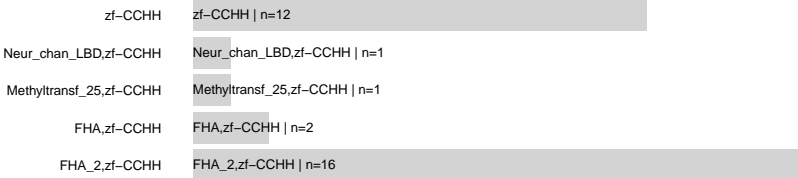
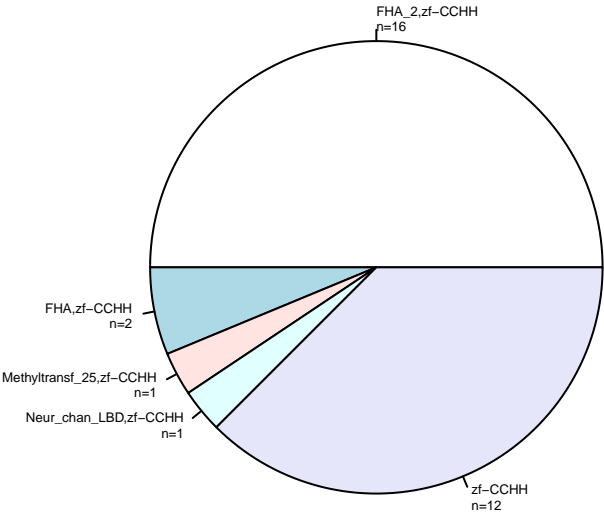
absent absent | n=1



zf-CCHH.HG1.6
APLF

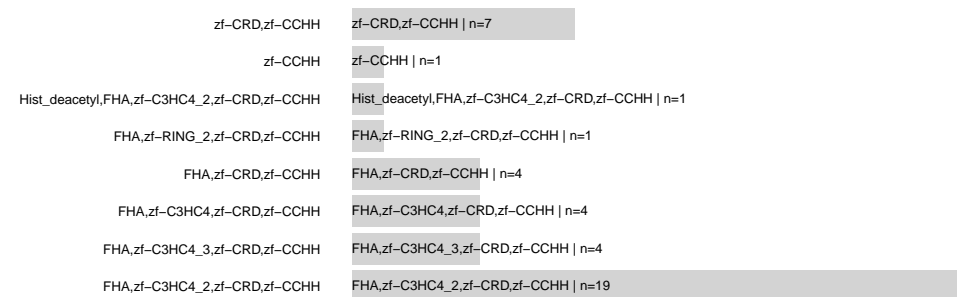
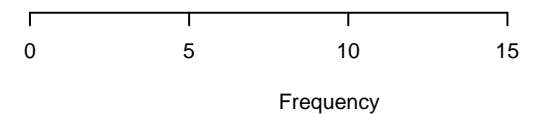


Gain: Bremin,NA
Presence Eukaryota prob = 0.03
Present: 30
Losses: NA

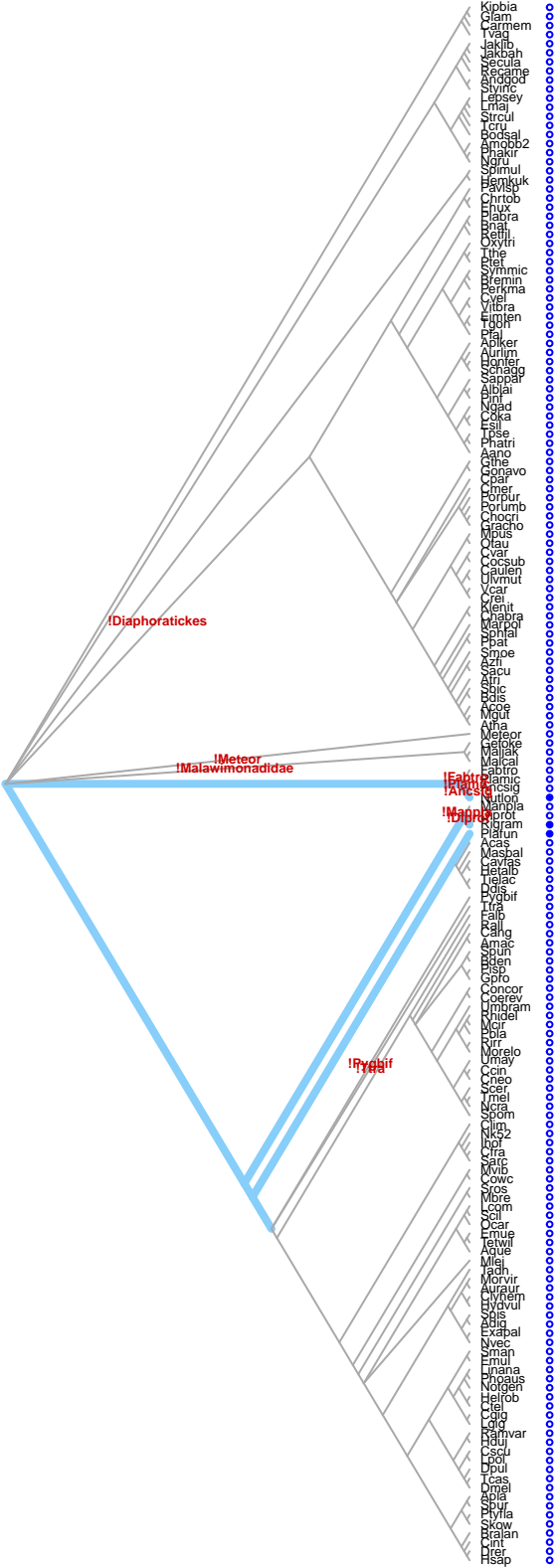


0 5 10 15
Frequency

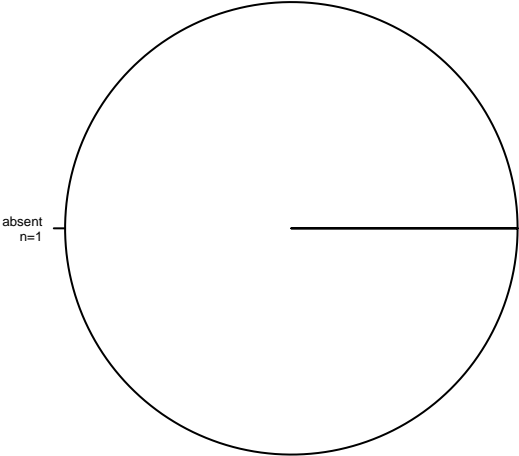
Gain: NA
Presence Eukaryota prob = 1.00
Present: 40
Losses: NA



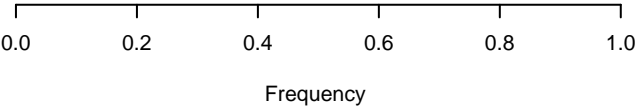
zf-CCHH.HG1.8
like:APLF



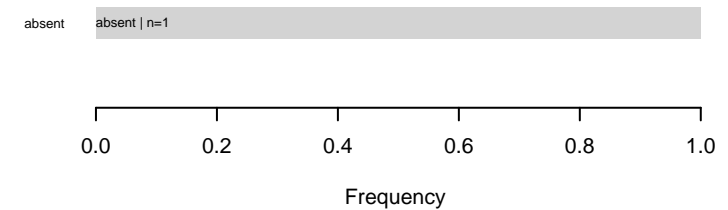
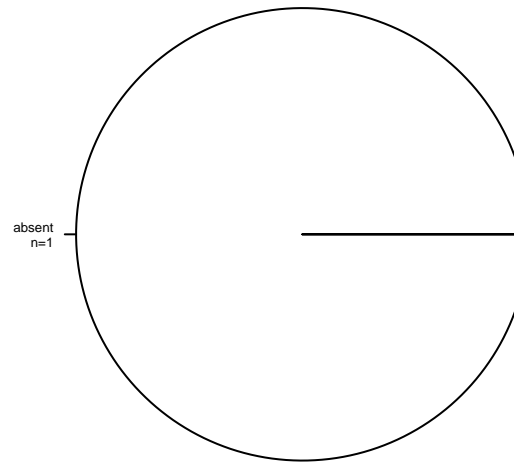
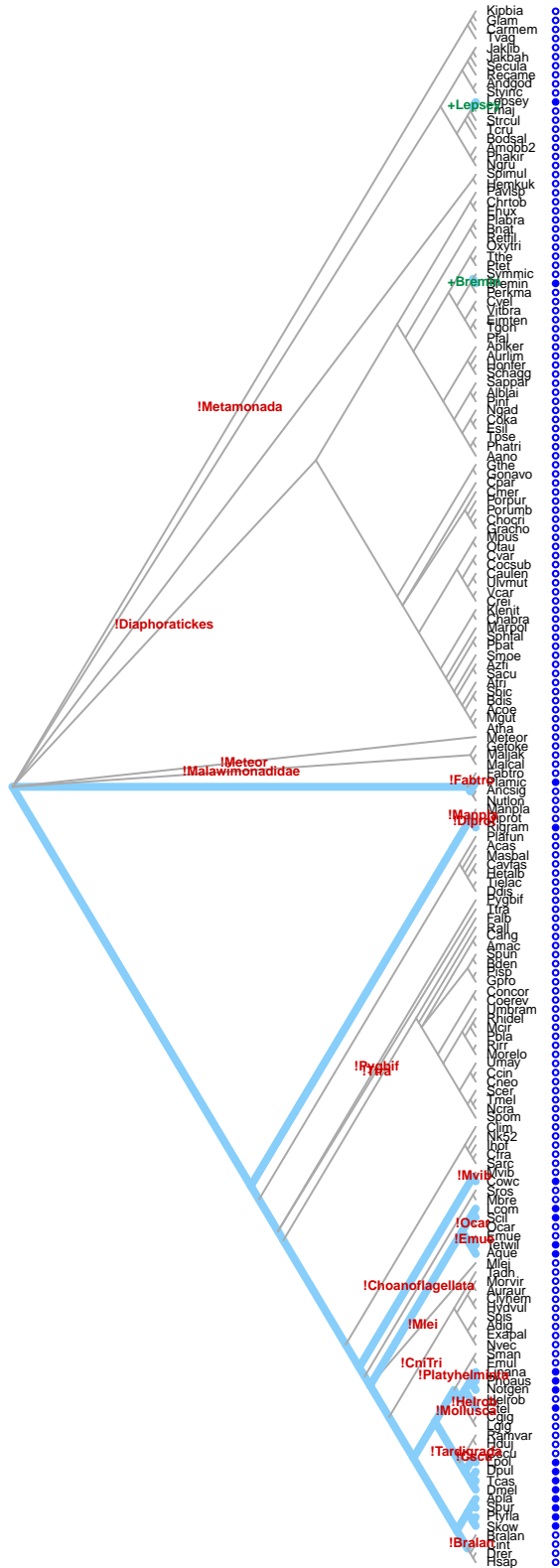
Gain: NA
Presence Eukaryota prob = 0.61
Present: 3
Losses: NA



absent absent | n=1

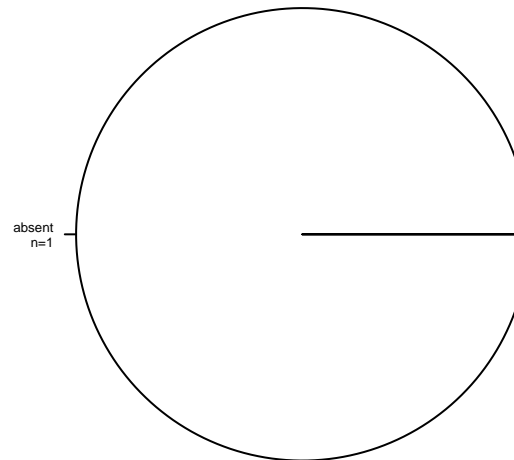
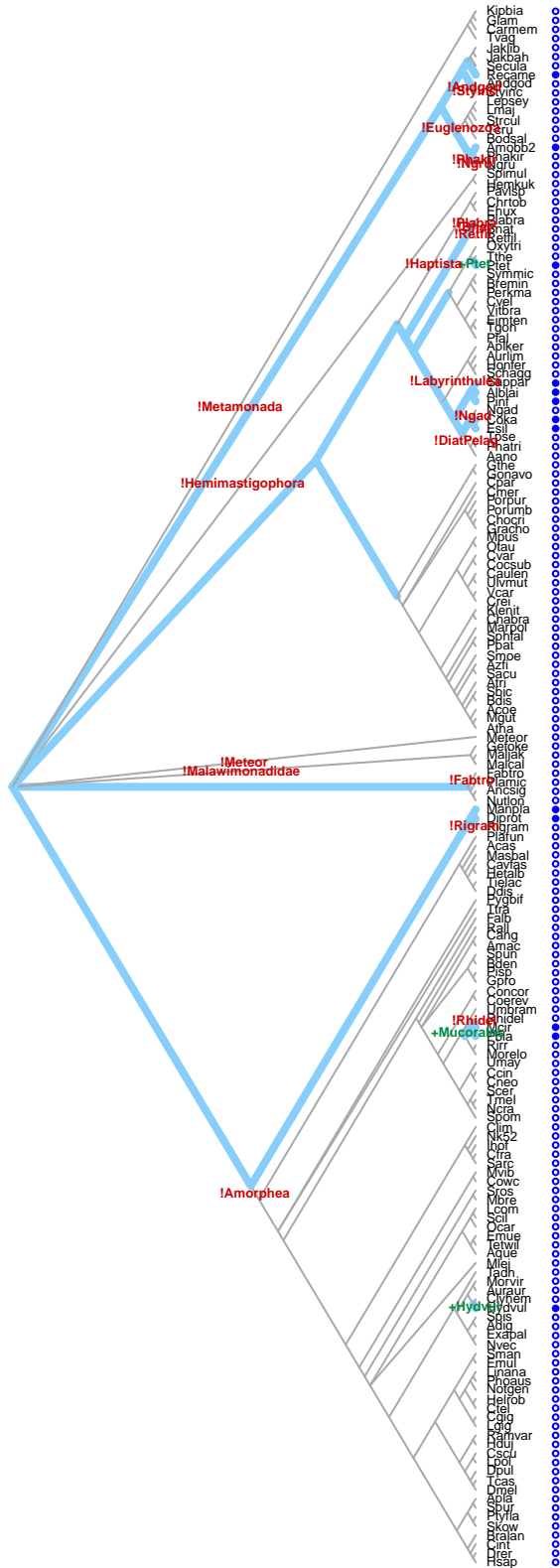


zf-CCHH.HG1.9
like:APLF

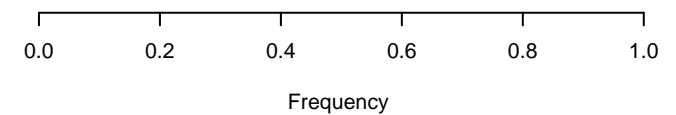


Gain: Bremin,Lepsey,NA
Presence Eukaryota prob = 0.74
Present: 21
Losses: NA

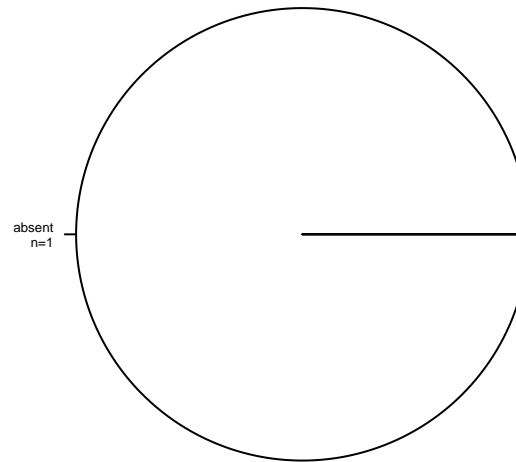
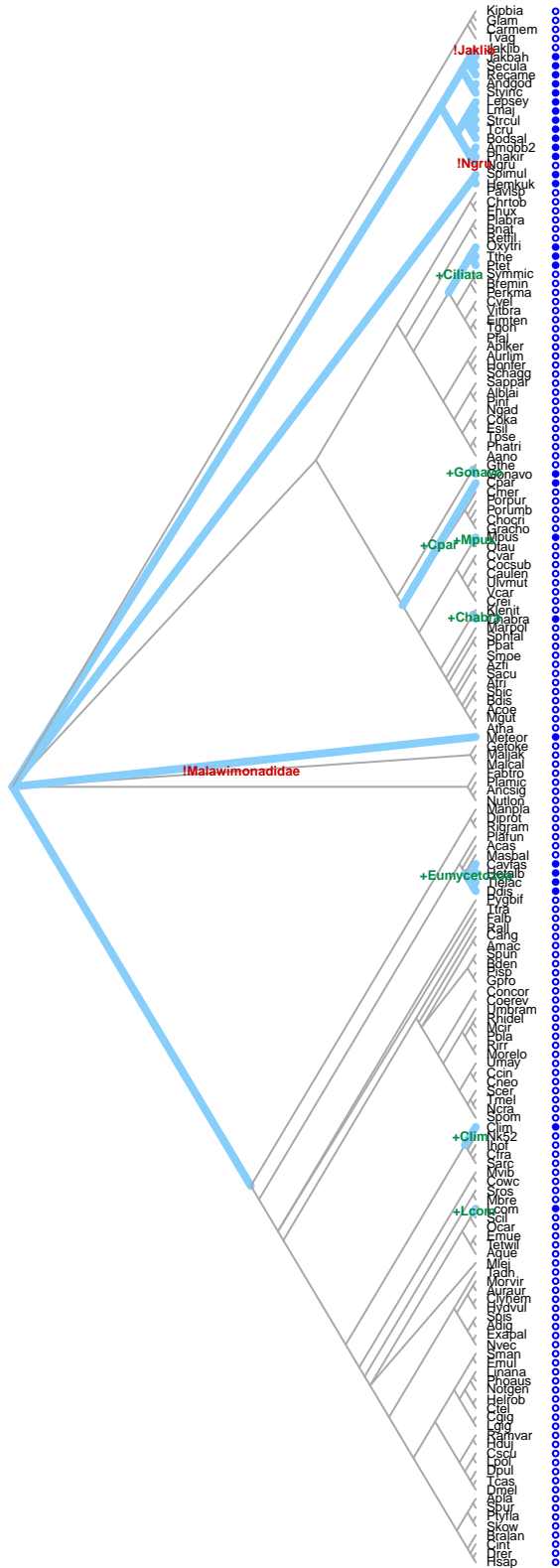
**Nucleoplasmin.HG1.0
like:NPM1/NPM2/NPM3**



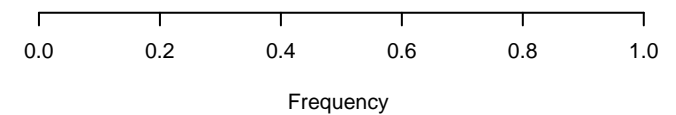
absent absent | n=1



**Nucleoplasmin.HG1.1
like:NPM1/NPM2/NPM3**

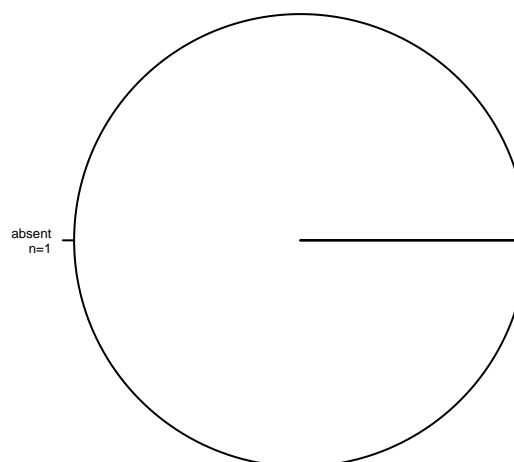
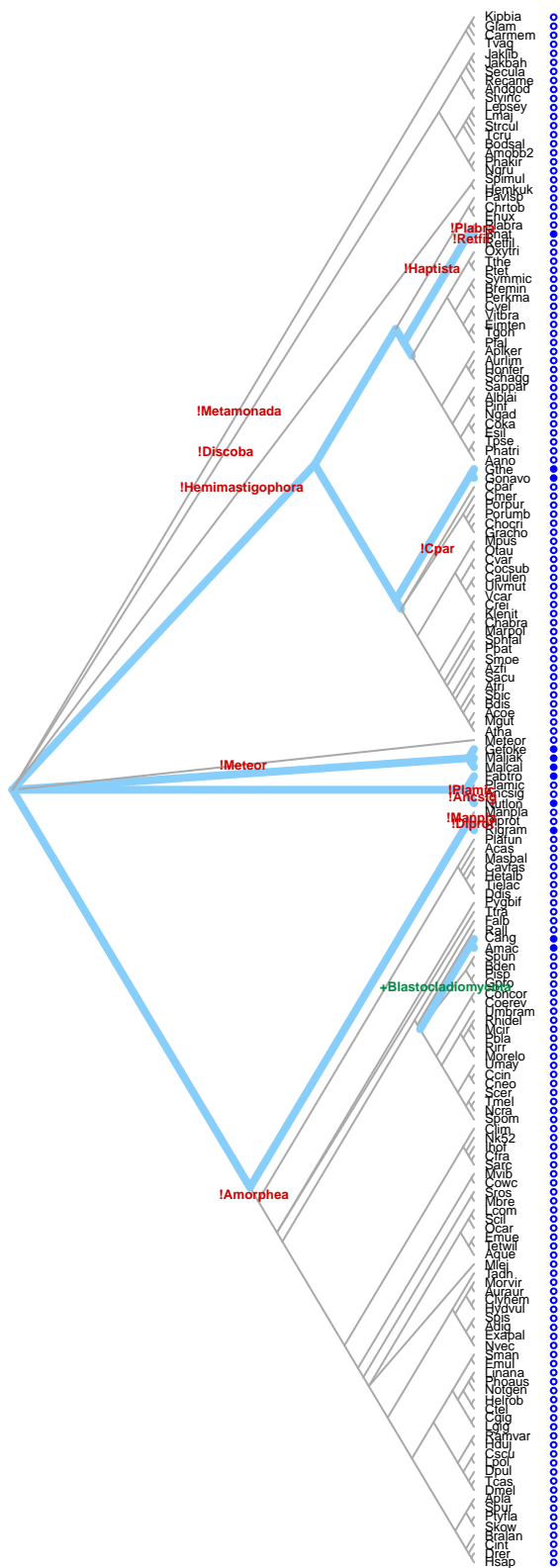


absent absent | n=1

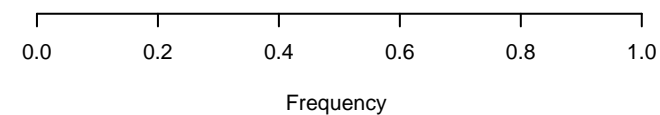


Gain: Lcom, Clim, Eumycetozoa, Chabra, Mpus, Cpar, Gonavo, Ciliata, NA
 Presence Eukaryota prob = 0.53
 Present: 28
 Losses: NA

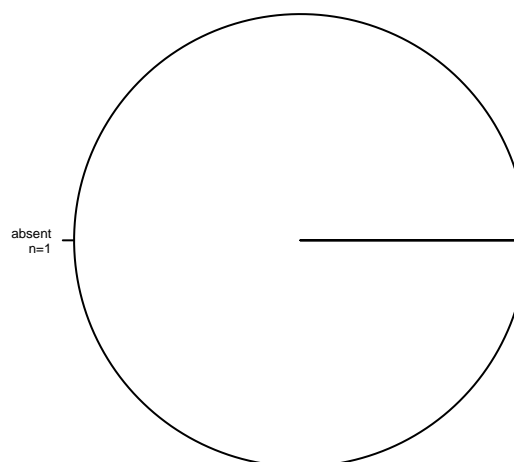
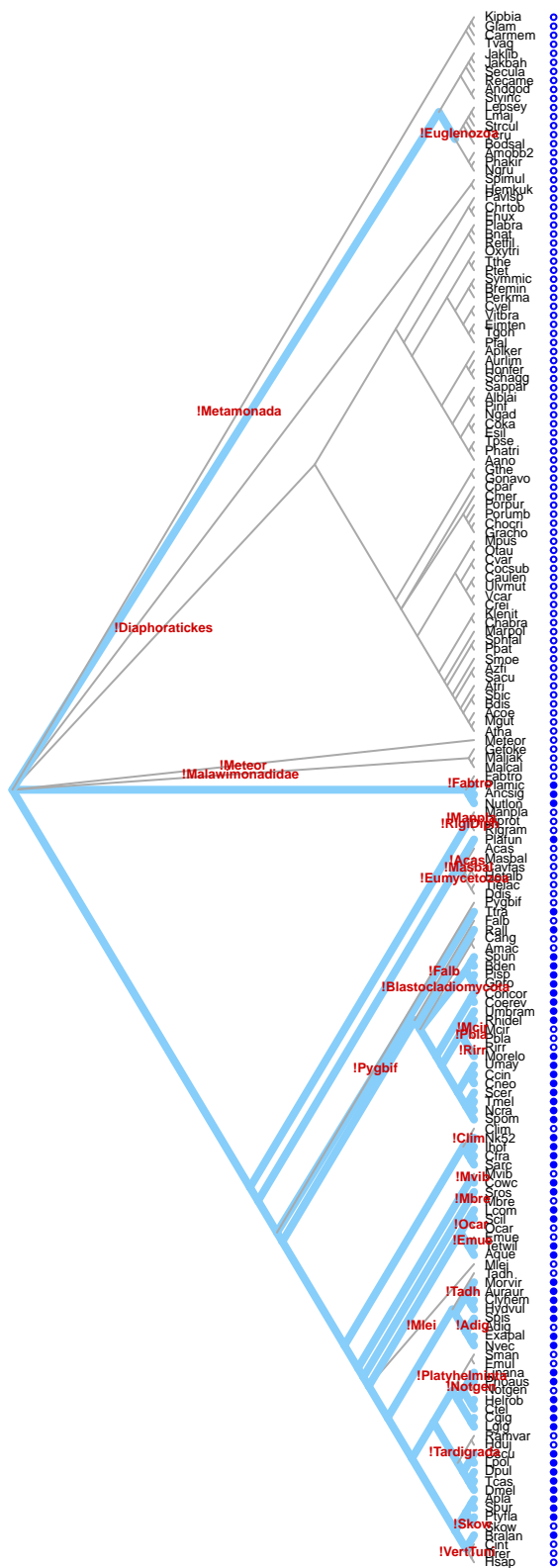
**Nucleoplasmin.HG1.2
like:NPM1/NPM2/NPM3**



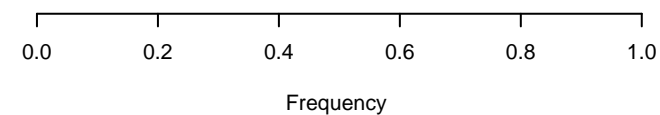
absent absent | n=1

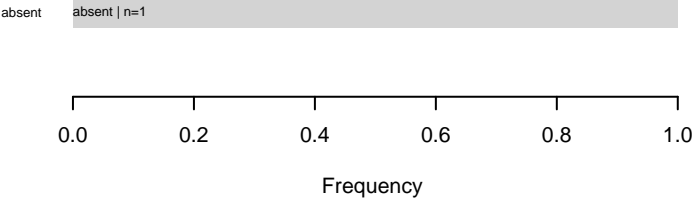
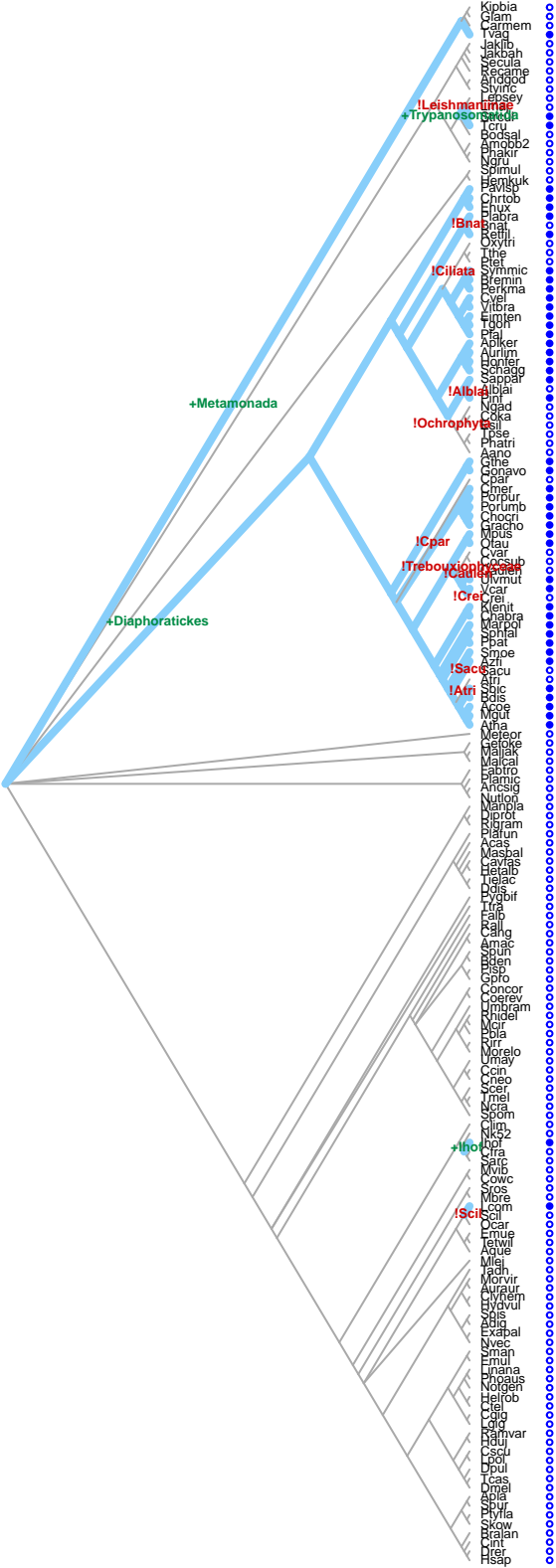


**Nucleoplasmin.HG1.3
like:NPM1/NPM2/NPM3**

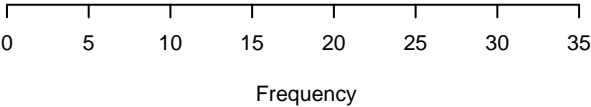
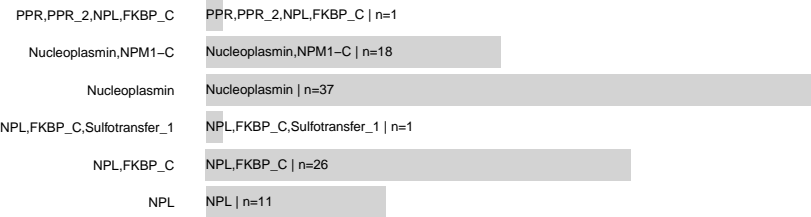
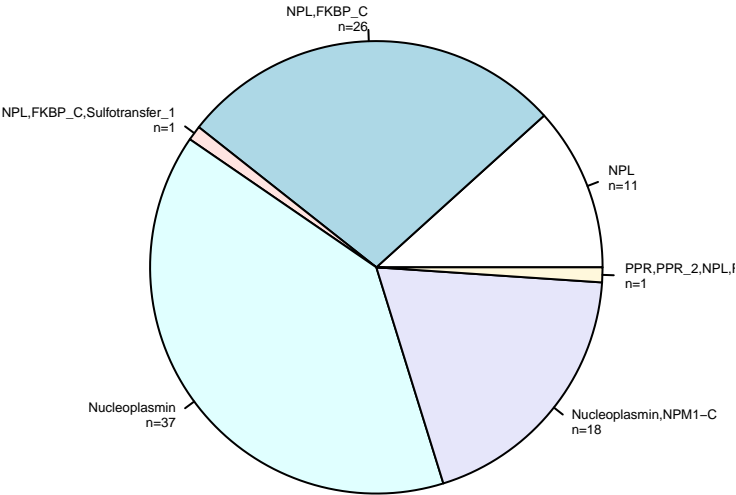
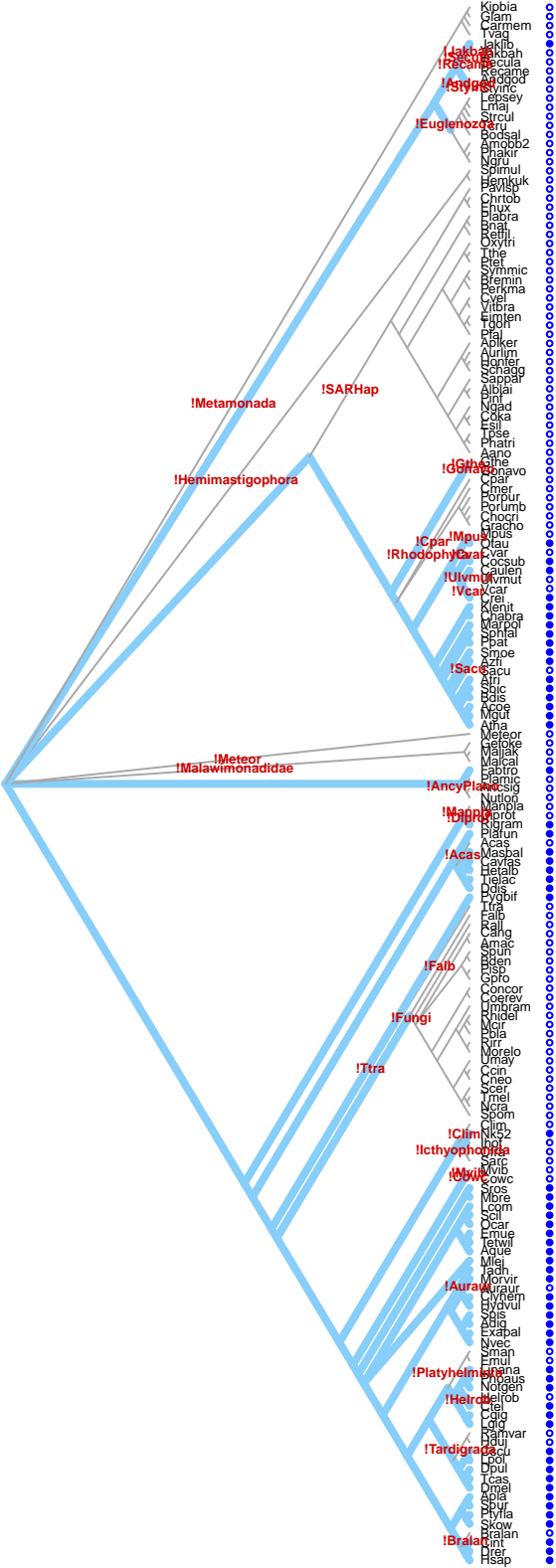


absent absent | n=1





Nucleoplasmin.HG1.5
NPM1/NPM2/NPM3



Gain: NA
Presence Eukaryota prob = 1.00
Present: 63
Losses: NA