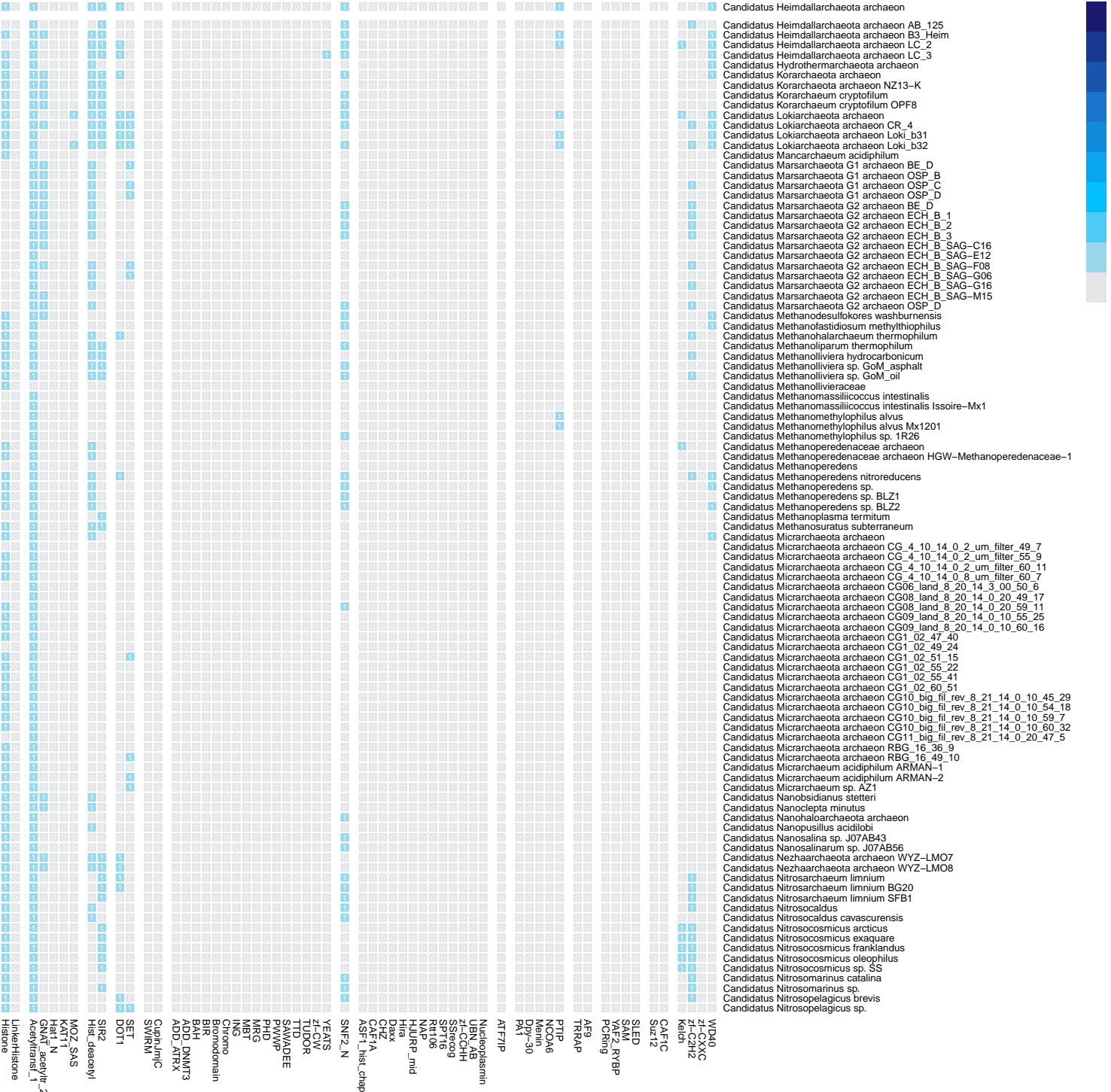


Gene presence per taxon in Archaea (superkingdom) (1/20)

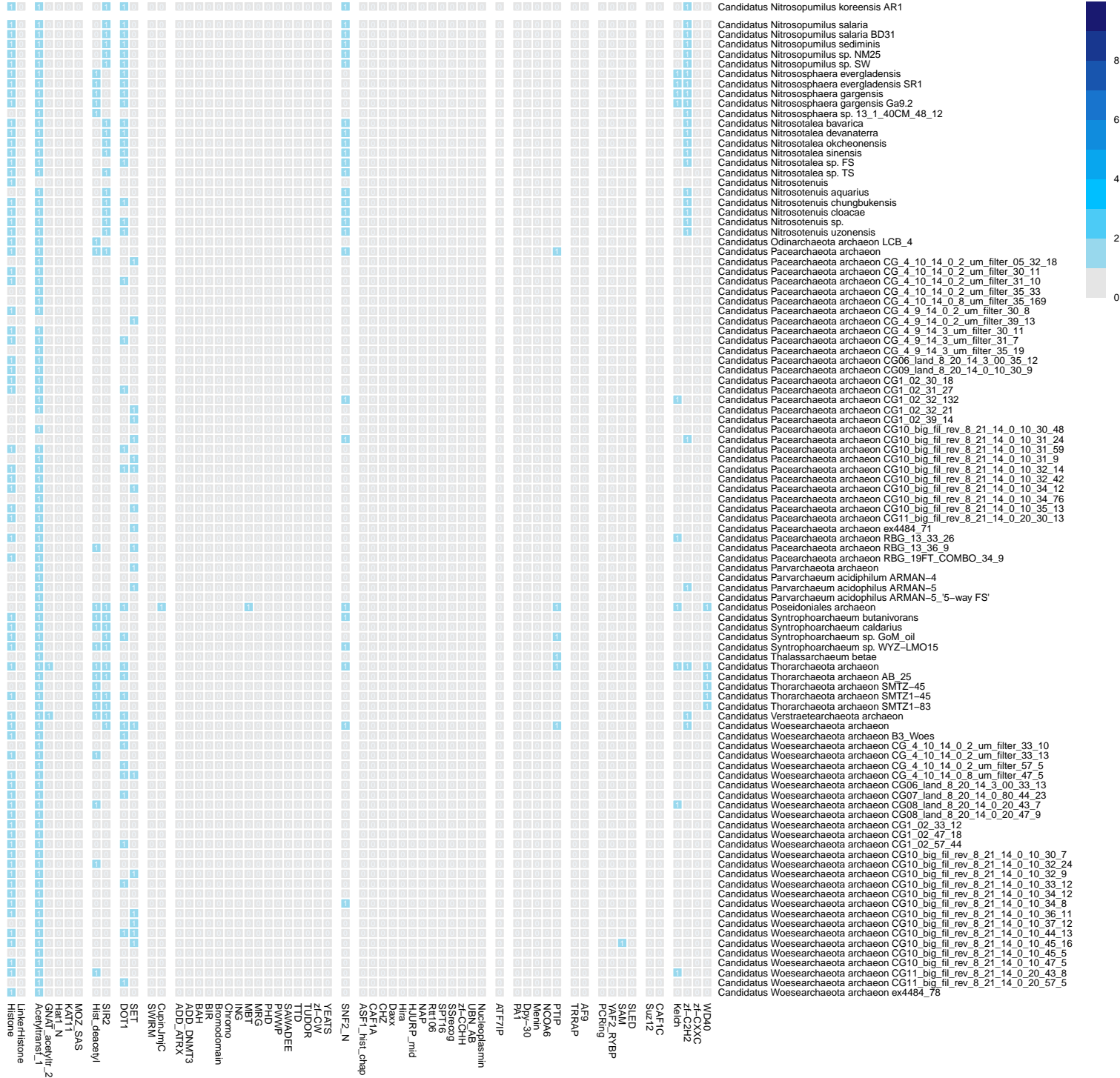


[illegible]

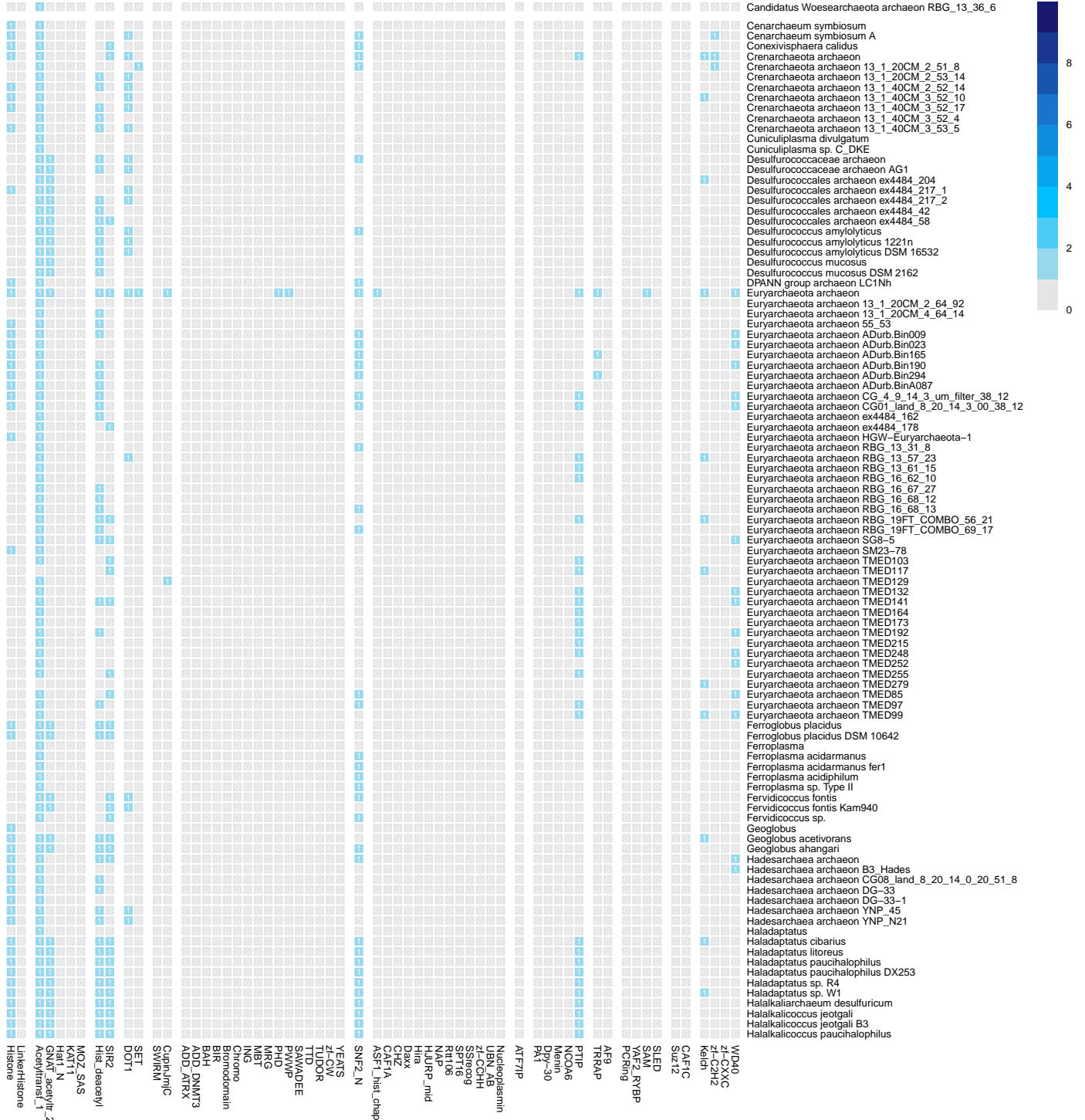
Gene presence per taxon in Archaea (superkingdom) (3/20)



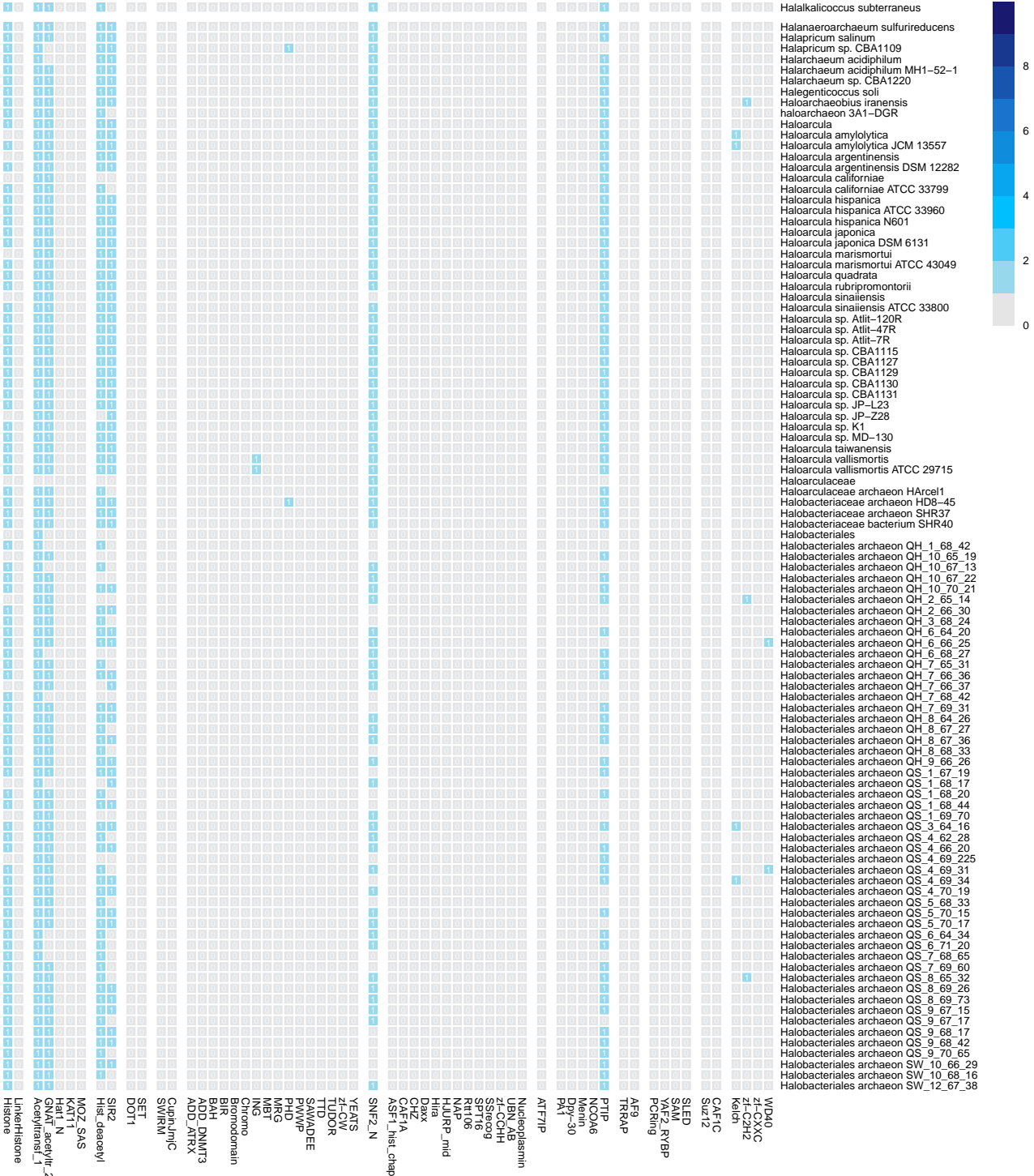
Gene presence per taxon in Archaea (superkingdom) (4/20)



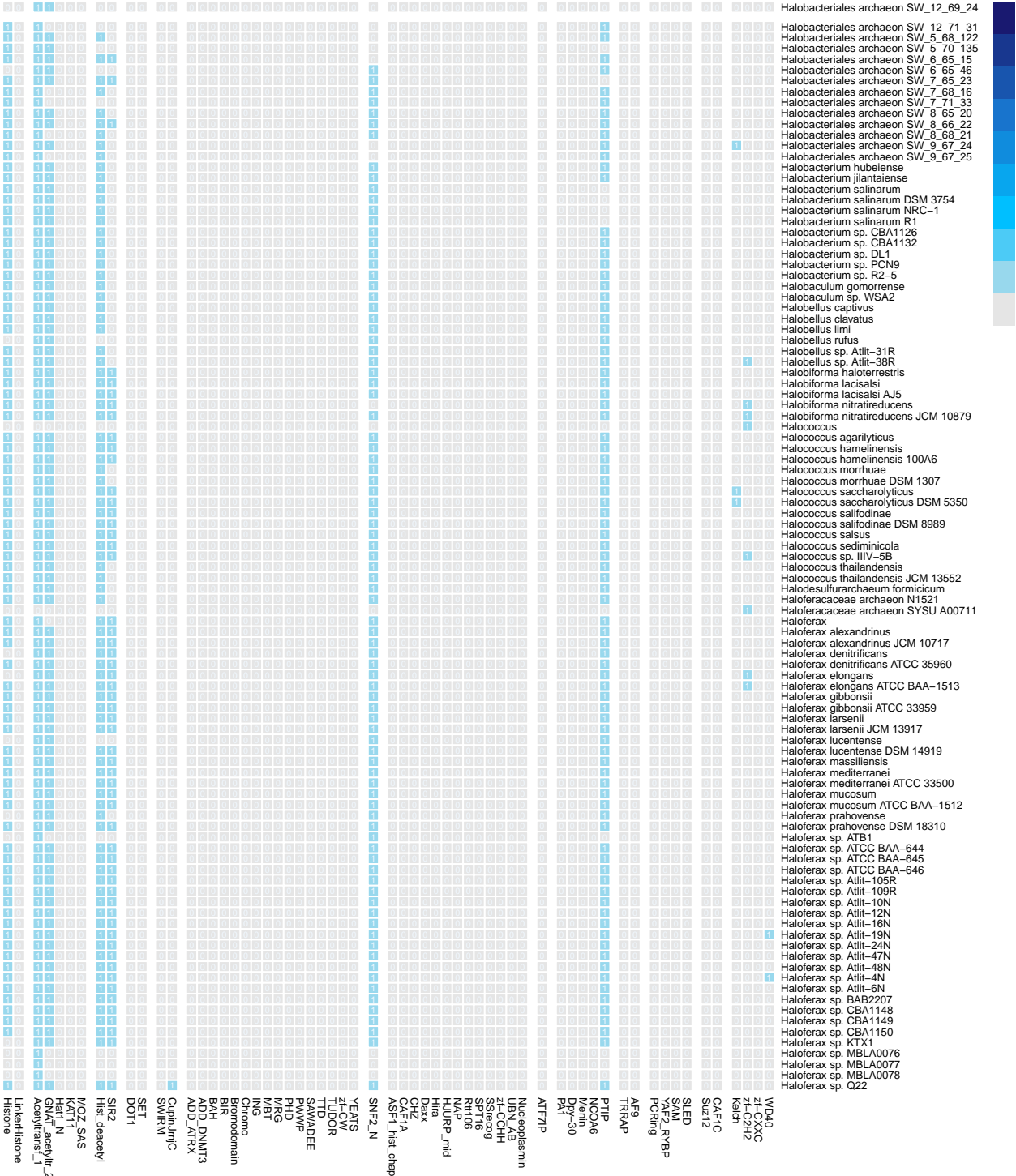
Gene presence per taxon in Archaea (superkingdom) (5/20)



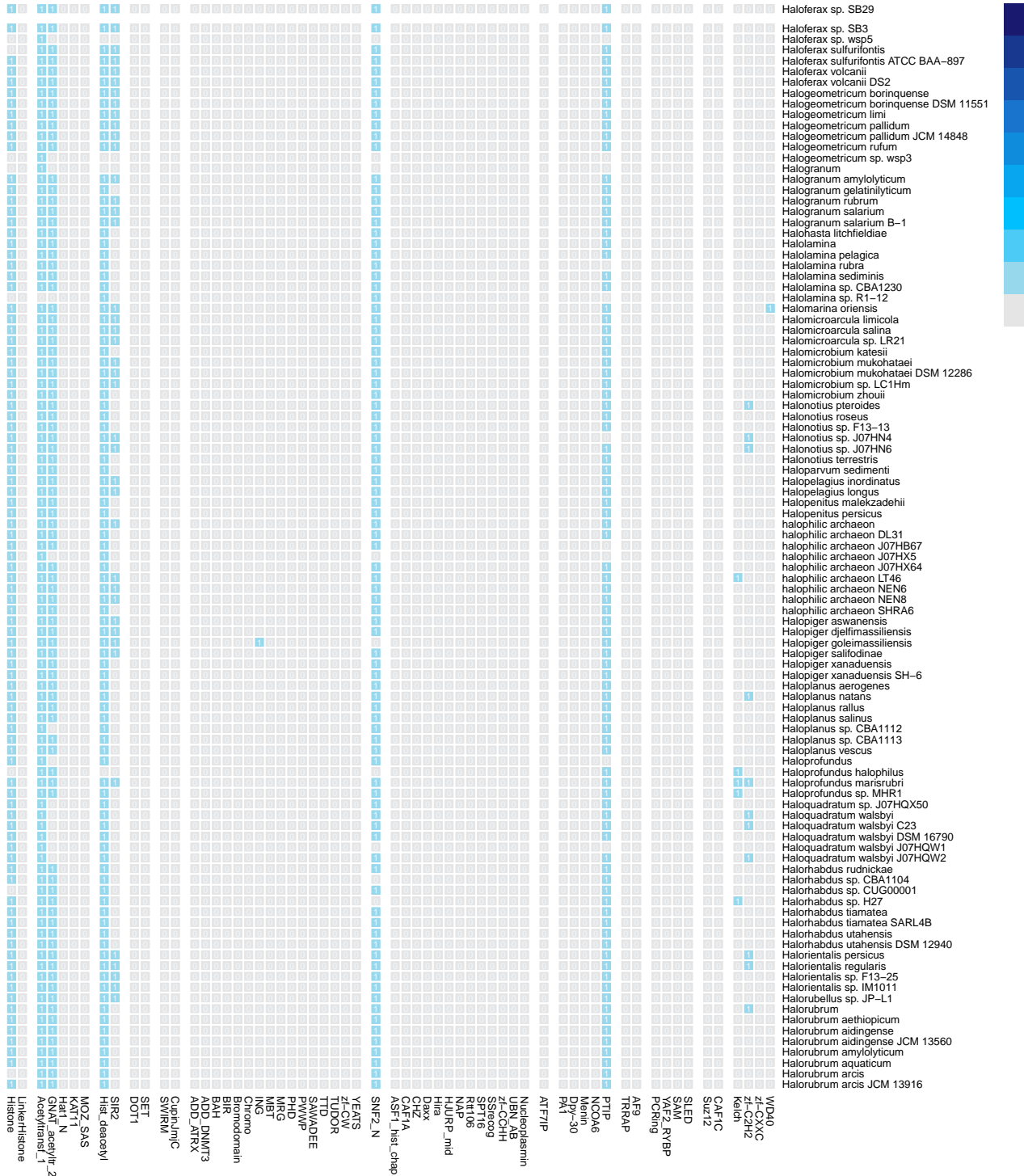
Gene presence per taxon in Archaea (superkingdom) (6/20)



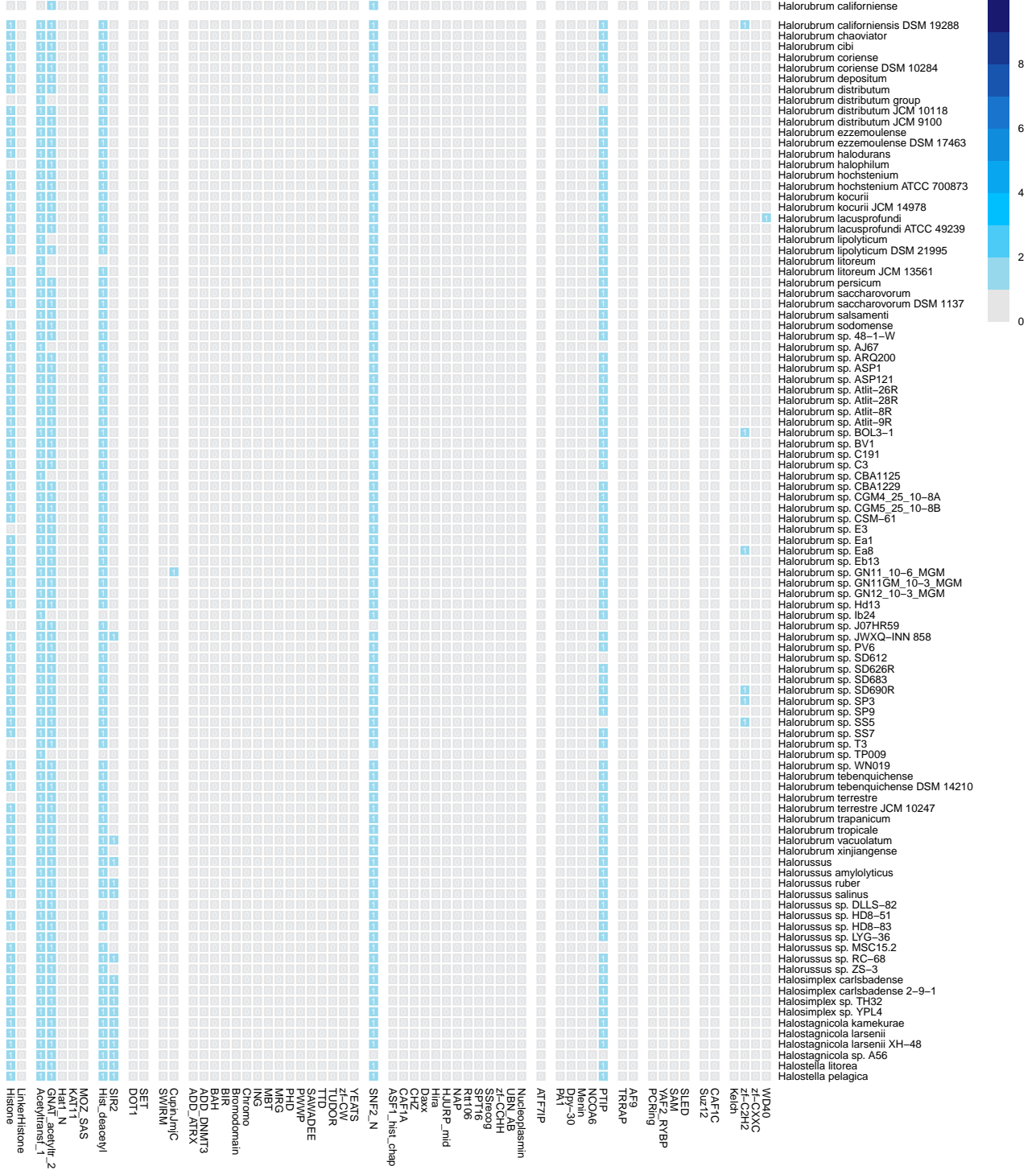
Gene presence per taxon in Archaea (superkingdom) (7/20)



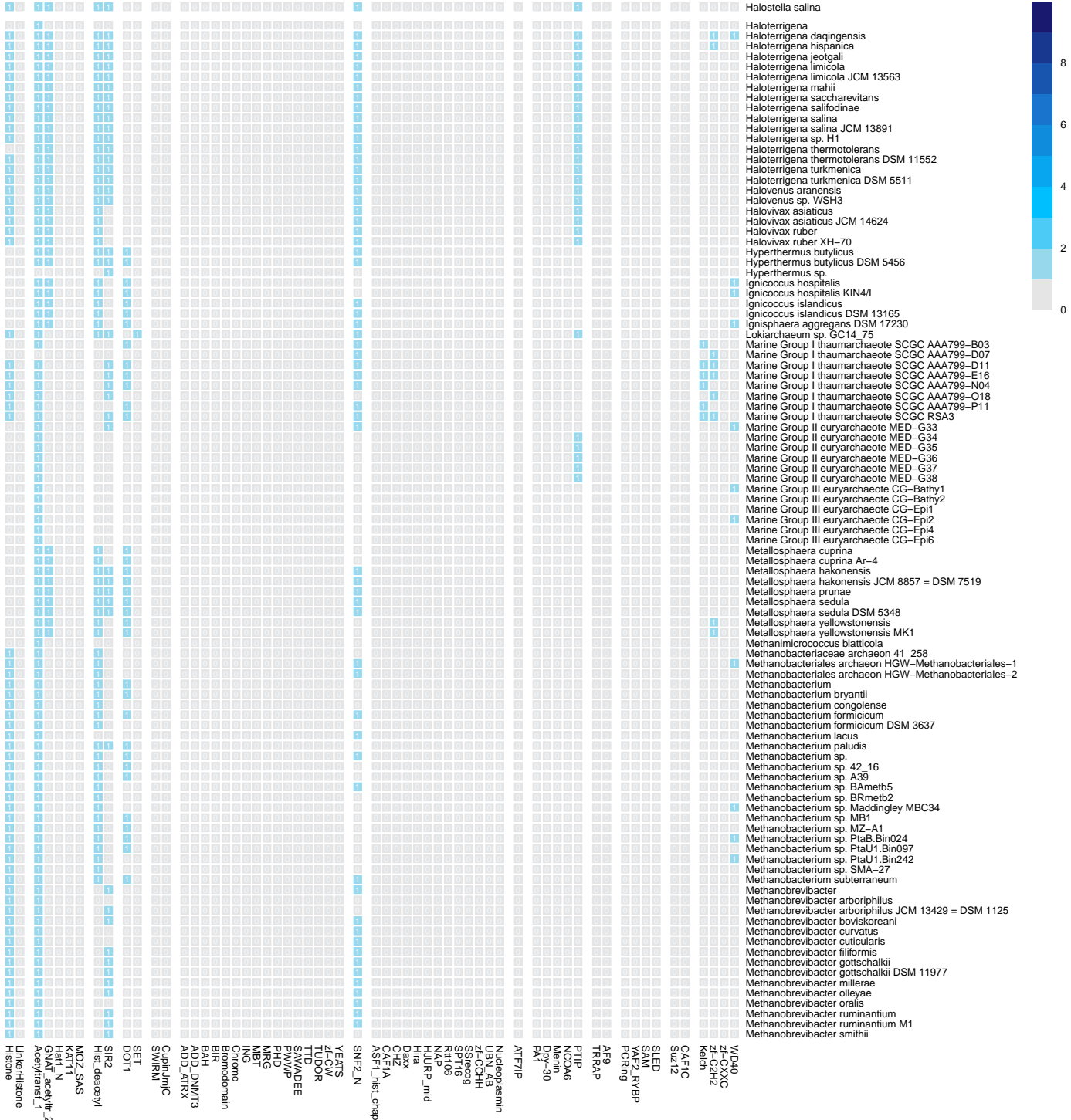
Gene presence per taxon in Archaea (superkingdom) (8/20)



Gene presence per taxon in Archaea (superkingdom) (9/20)



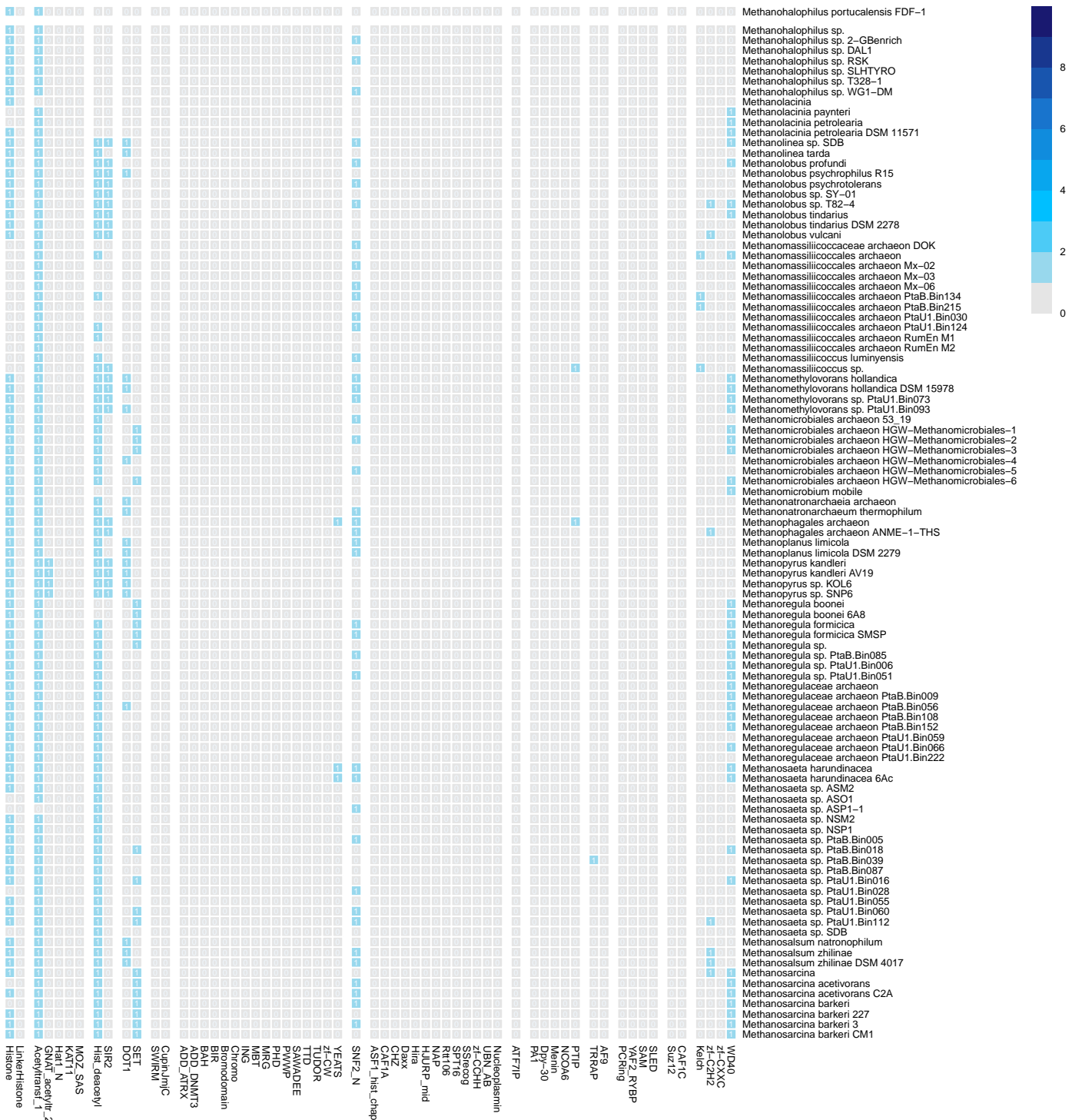
Halostella salina



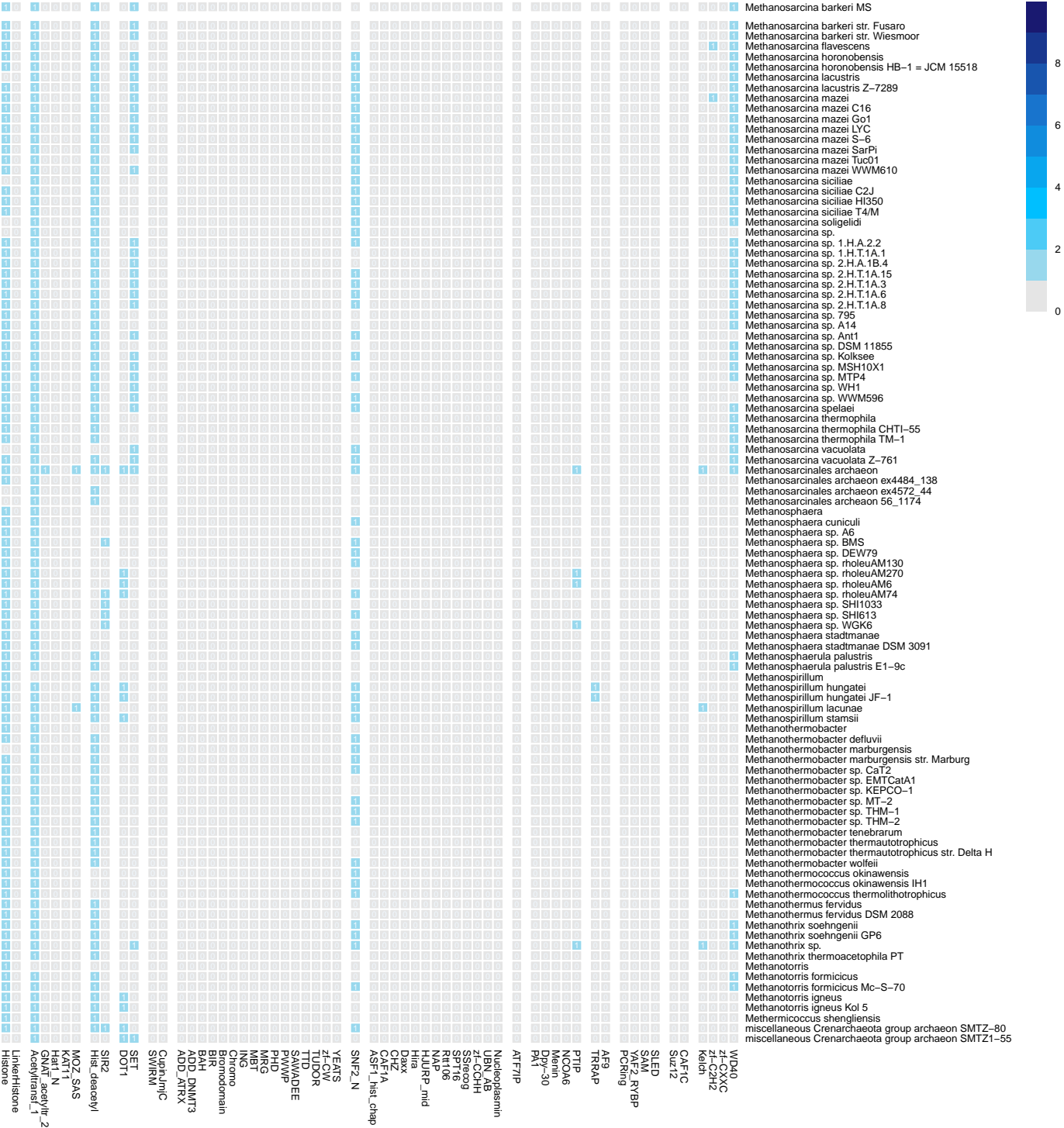
Gene presence per taxon in Archaea (superkingdom) (11/20)



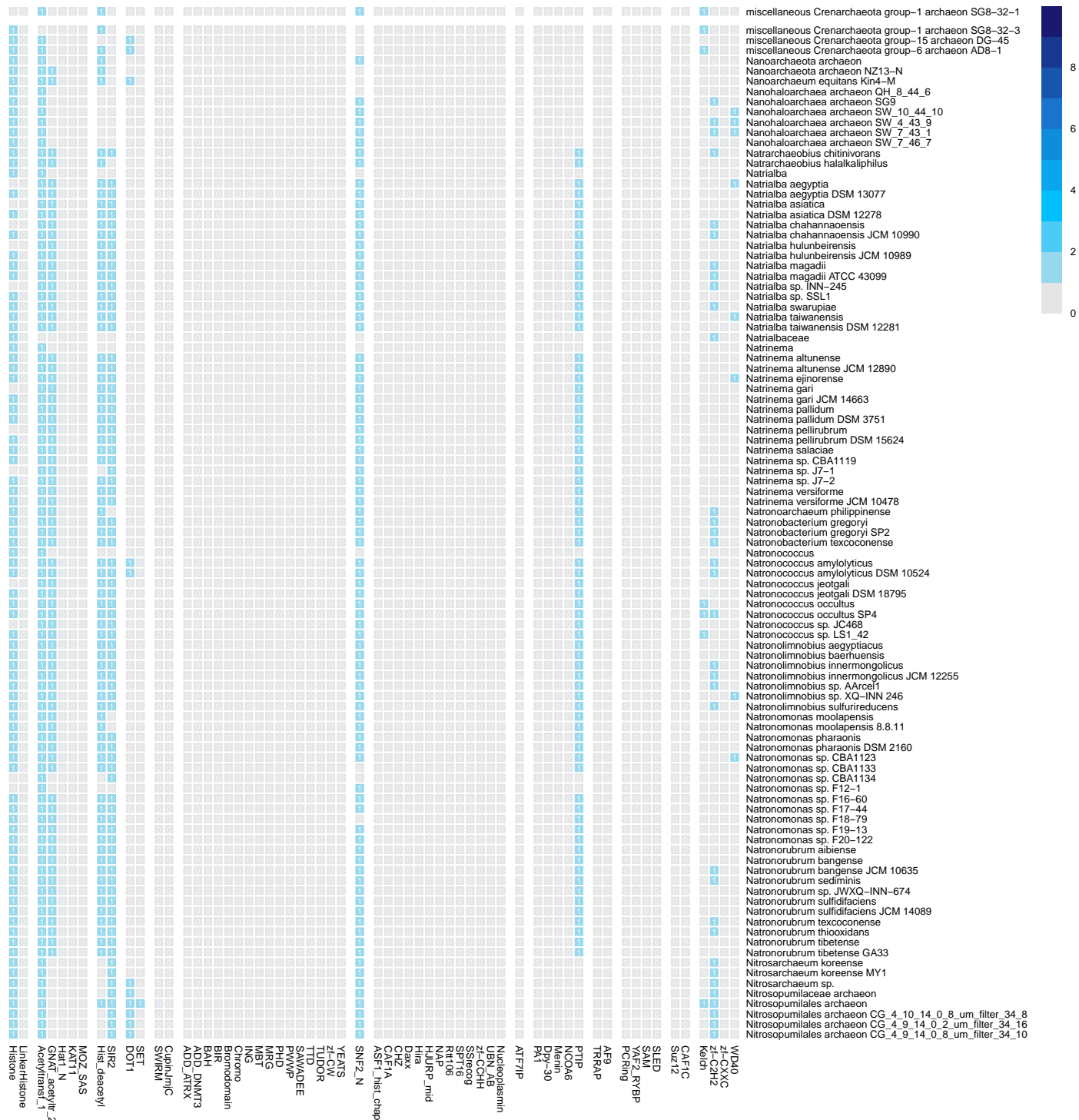
Gene presence per taxon in Archaea (superkingdom) (12/20)



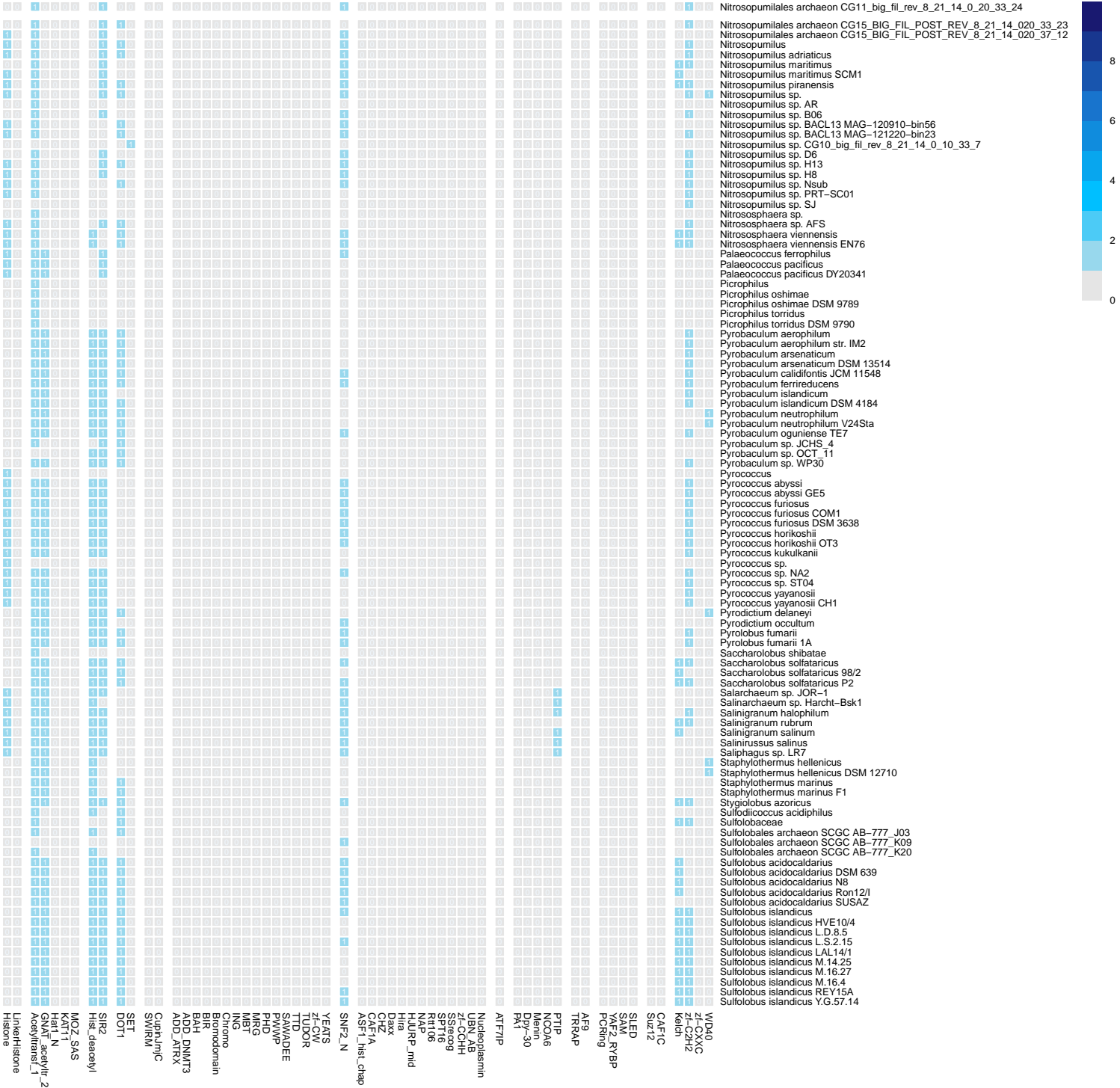
Gene presence per taxon in Archaea (superkingdom) (13/20)



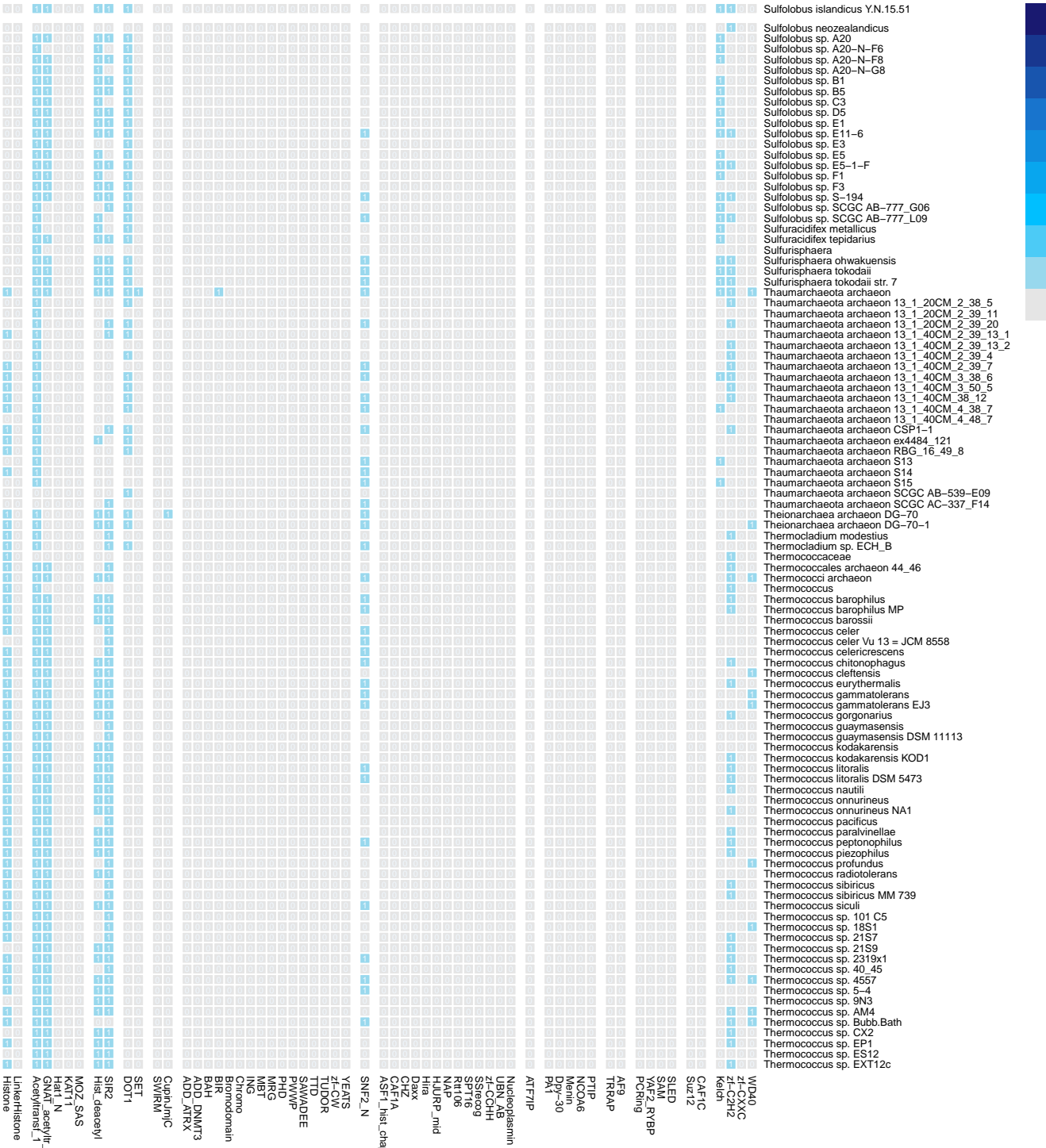
Gene presence per taxon in Archaea (superkingdom) (14/20)



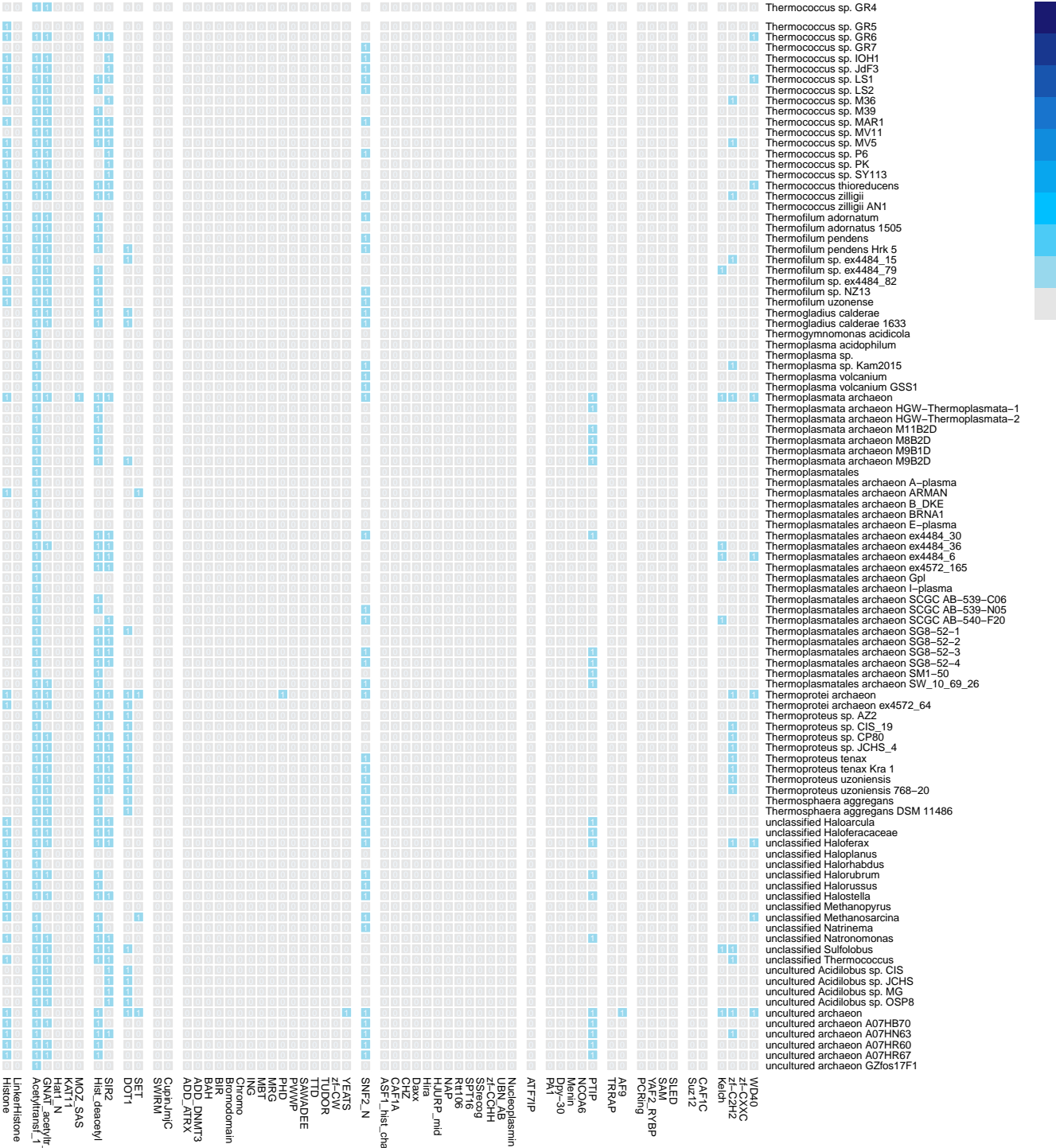
Gene presence per taxon in Archaea (superkingdom) (15/20)



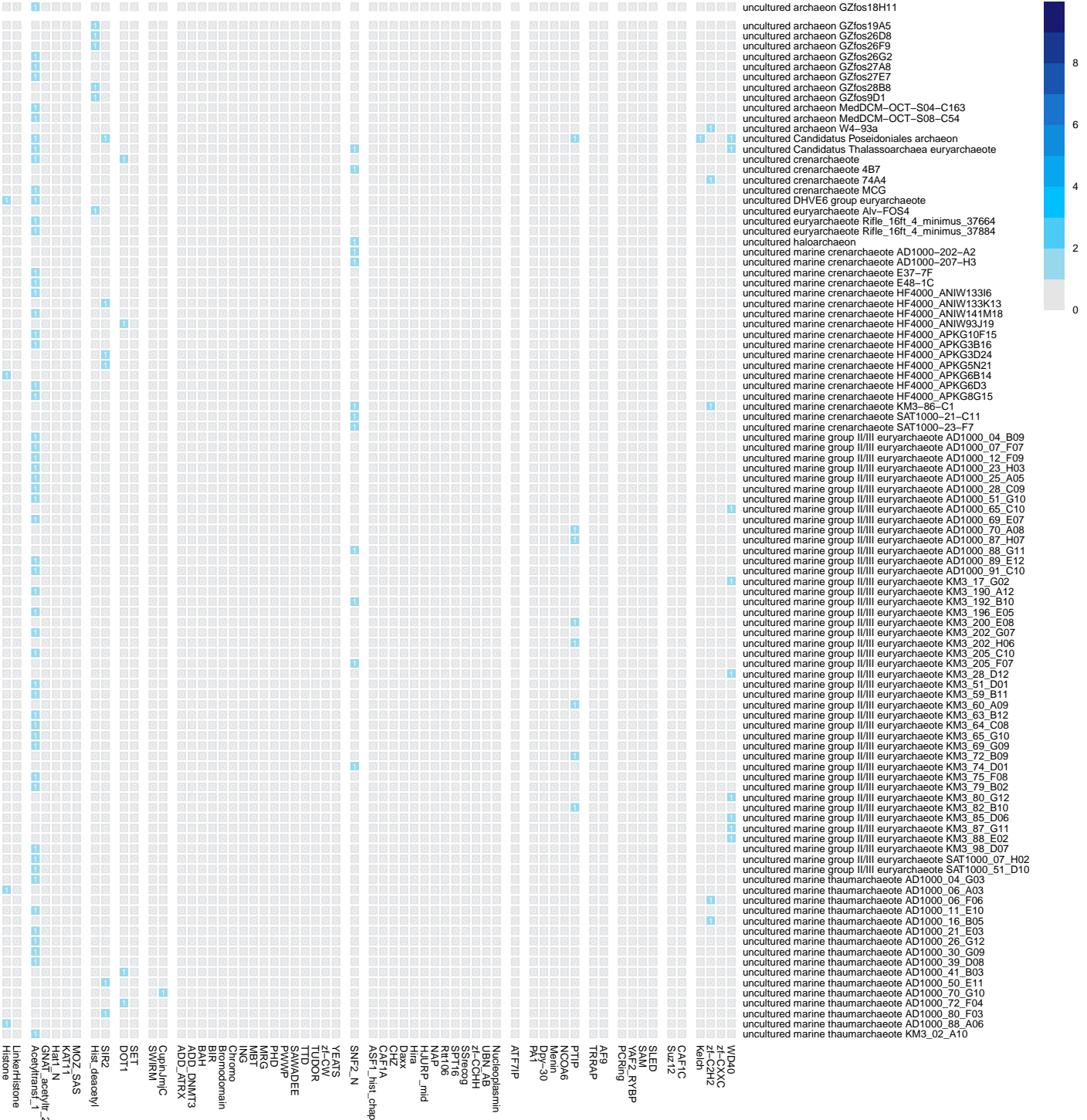
Gene presence per taxon in Archaea (superkingdom) (16/20)



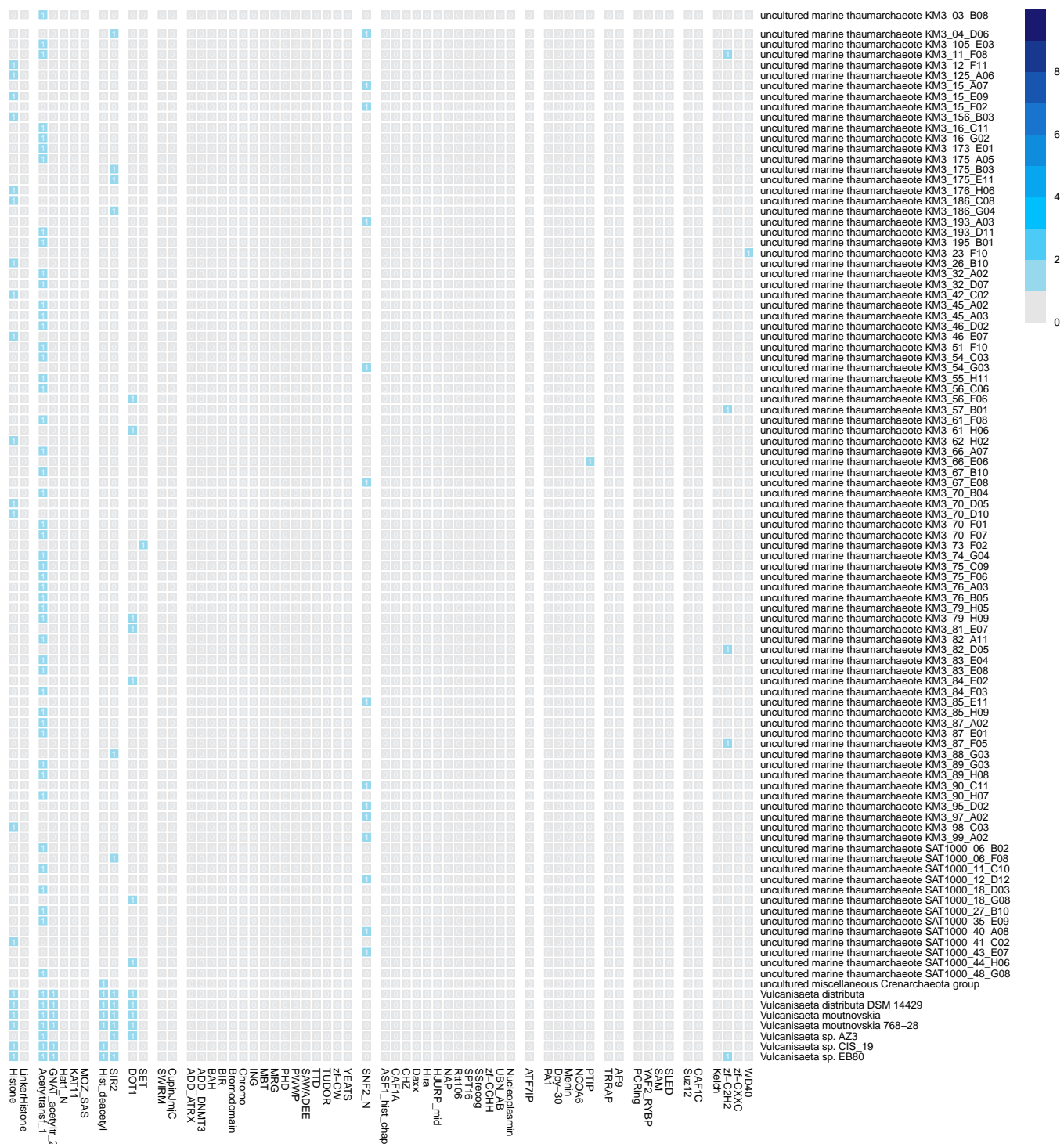
Gene presence per taxon in Archaea (superkingdom) (17/20)

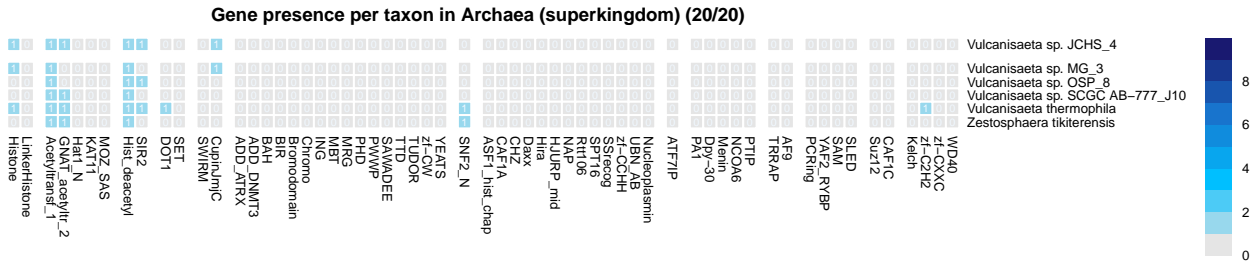


Gene presence per taxon in Archaea (superkingdom) (18/20)



Gene presence per taxon in Archaea (superkingdom) (19/20)





Protein Domains (Y-axis):

- Acidianus
- Acidianus ambivalens
- Acidianus brierleyi
- Acidianus hospitalis
- Acidianus hospitalis W1
- Acidianus infernus
- Acidianus manzaensis
- Acidianus sulfidivorans
- Acidianus sulfidivorans JP7
- Acidianus tengchongensis
- Acidilobus saccharovorans
- Acidilobus saccharovorans 345-15
- Acidilobus sp. 7A
- Acidilobus sp. SCGC AC-742_E15
- Acidilobus sp. SCGC AC-742_M05
- Acidiplasma
- Acidiplasma aeolicum
- Acidiplasma cupricumulans
- Acidiplasma sp. MBA-1
- Aciduliprofundum boonei
- Aciduliprofundum boonei T469
- Aciduliprofundum sp.
- Aciduliprofundum sp. MAR08-339
- Aeropyrum camini
- Aeropyrum camini SY1 = JCM 12091
- Aeropyrum pernix
- Aeropyrum pernix K1
- Algarchoeota archaeon NZ13_MG1
- Anaerobic archaeon MK-D1
- ANME-1 cluster archaeon ex4572_4
- ANME-2 cluster archaeon
- ANME-2 cluster archaeon HR1
- Archaeoglobales archaeon
- Archaeoglobales archaeon ex4484_92
- Archaeoglobi archaeon
- Archaeoglobus fulgidus
- Archaeoglobus fulgidus DSM 4304
- Archaeoglobus fulgidus DSM 8774
- Archaeoglobus profundus
- Archaeoglobus profundus DSM 5631
- Archaeoglobus sp.
- Archaeoglobus sulfatocalidus
- Archaeoglobus sulfatocalidus PM70-1
- Archaeoglobus veneficus
- Archaeoglobus veneficus SNP6
- archaeon
- archaeon 13_1_20CM_2_51_12
- archaeon 13_1_20CM_2_54_9
- archaeon 13_1_20CM_52_20
- archaeon 13_1_40CM_2_52_13
- archaeon 13_1_40CM_2_52_4
- archaeon 13_1_40CM_4_53_4
- archaeon 13_2_20CM_2_52_21
- archaeon 13_2_20CM_2_53_6
- archaeon ADurb_Bin336
- archaeon BMS3Abin16
- archaeon BMS3Abin17
- archaeon BMS3Bbin15
- archaeon BMS3Bbin16
- archaeon CG_4_10_14_0_2_um_filter_Archaea_38_6
- archaeon CG_4_8_14_3_um_filter_38_5
- archaeon CG06_land_8_20_14_3_00_37_11
- archaeon CG07_land_8_20_14_0_80_38_8
- archaeon CG10_big_fil_rev_8_21_14_0_10_43_11
- archaeon CG2_30_31_98
- archaeon D22
- archaeon GW2011_AR1
- archaeon GW2011_AR10
- archaeon GW2011_AR11
- archaeon GW2011_AR13
- archaeon GW2011_AR15
- archaeon GW2011_AR16
- archaeon GW2011_AR17
- archaeon GW2011_AR18
- archaeon GW2011_AR19
- archaeon GW2011_AR20
- archaeon GW2011_AR21
- archaeon GW2011_AR3
- archaeon GW2011_AR4
- archaeon GW2011_AR5
- archaeon GW2011_AR6
- archaeon GW2011_AR9
- archaeon HR01
- archaeon HR02
- archaeon HR03
- archaeon HR04
- archaeon HR05
- archaeon HR06
- archaeon MnTg01
- archaeon RBG_16_50_20
- archaeon SCG-AAA382B04
- Caldisphaera lagunensis
- Caldisphaera lagunensis DSM 15908
- Caldisphaera sp.
- Caldivirga maquiltingensis
- Caldivirga maquiltingensis IC-167
- Caldivirga sp. CIS_19
- Caldivirga sp. JCHS_4
- Caldivirga sp. MG_3

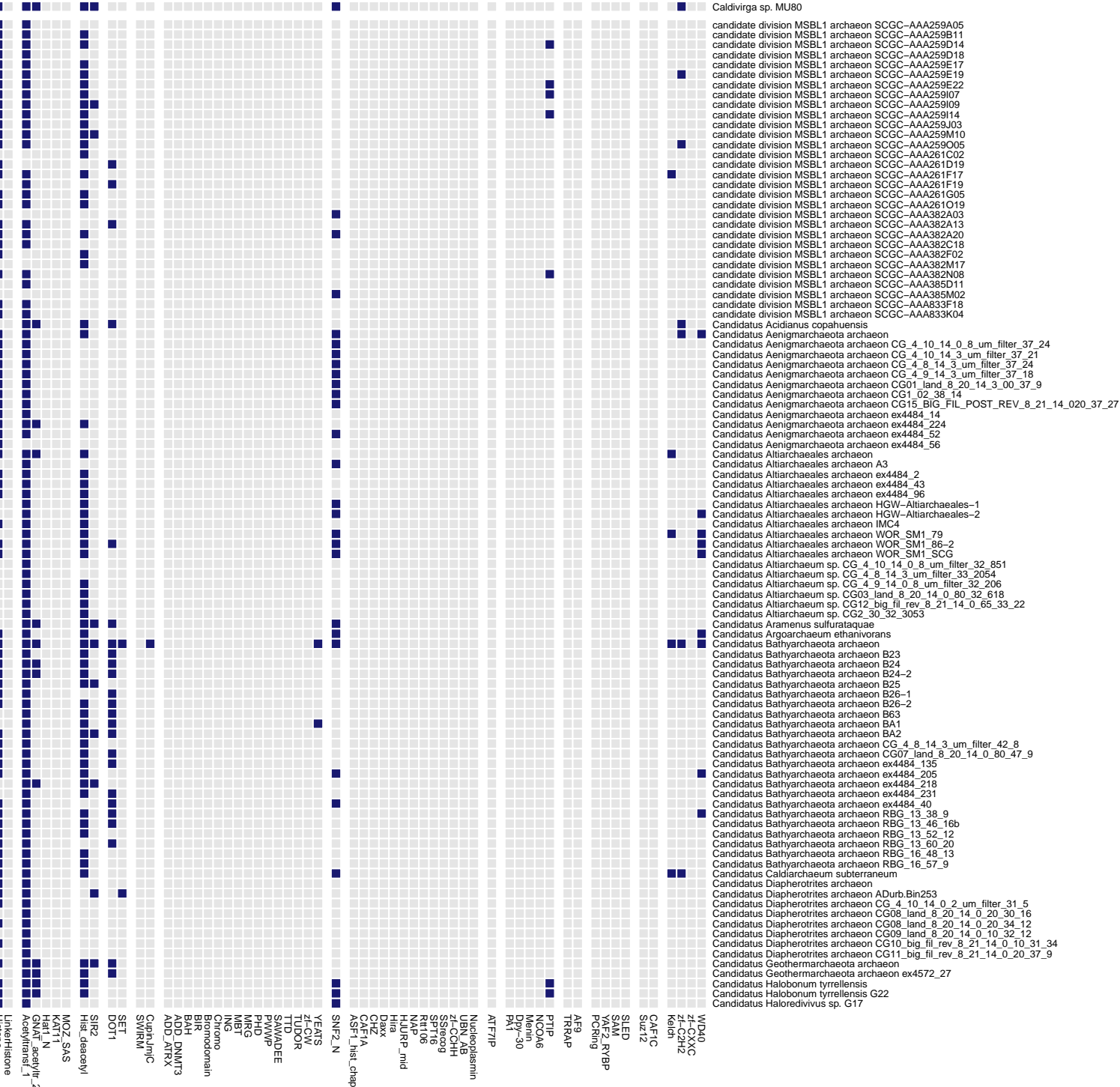
Species (X-axis):

- Nucleoplasmin
- Blebbistatin
- ZnOx
- Steroid
- SPR16
- RIT06
- NAP
- HJURP_mid
- Daxx
- Chaz
- CAF1A
- ASF1_hist_chap
- SNF2_N
- YEATS
- Zf-CNV
- TTROR
- SAWADDEE
- PWWP
- PHD
- MIRG
- MIRG
- MIRG
- Chromo
- Bromodomain
- BIR
- ADD_DNM73
- ADD_ARX
- Cytoplasmic SWIRM
- SET1
- DOT1
- SRF2
- Hist_deacetyl
- MOZ SAS
- KAT1 N
- KAT1 N
- GNAI_acetyl_2
- Acetylrns_1
- LineH1stone
- Histone

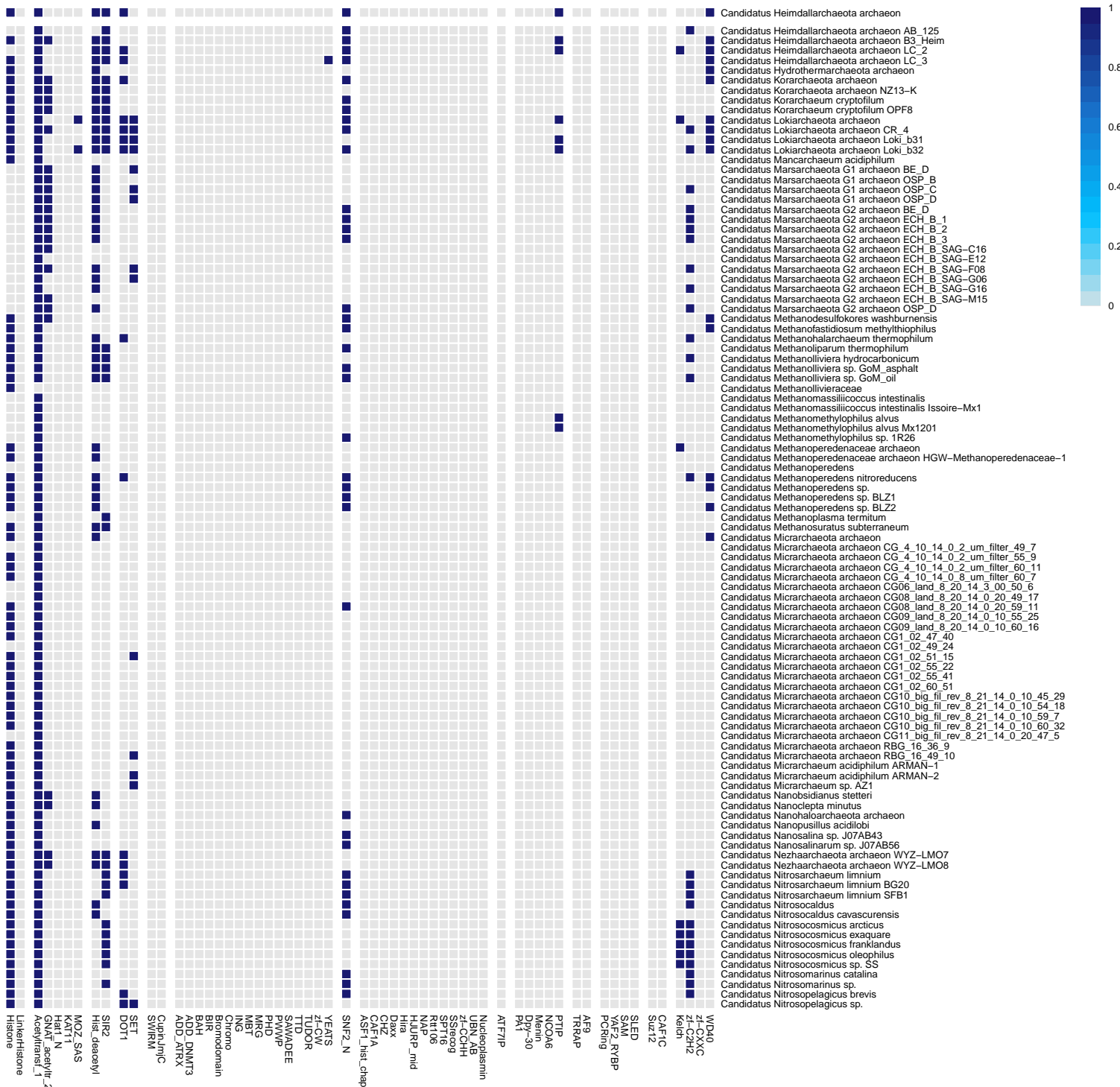
Legend:

- 0 (Light Blue)
- 0.2
- 0.4
- 0.6
- 0.8
- 1 (Dark Blue)

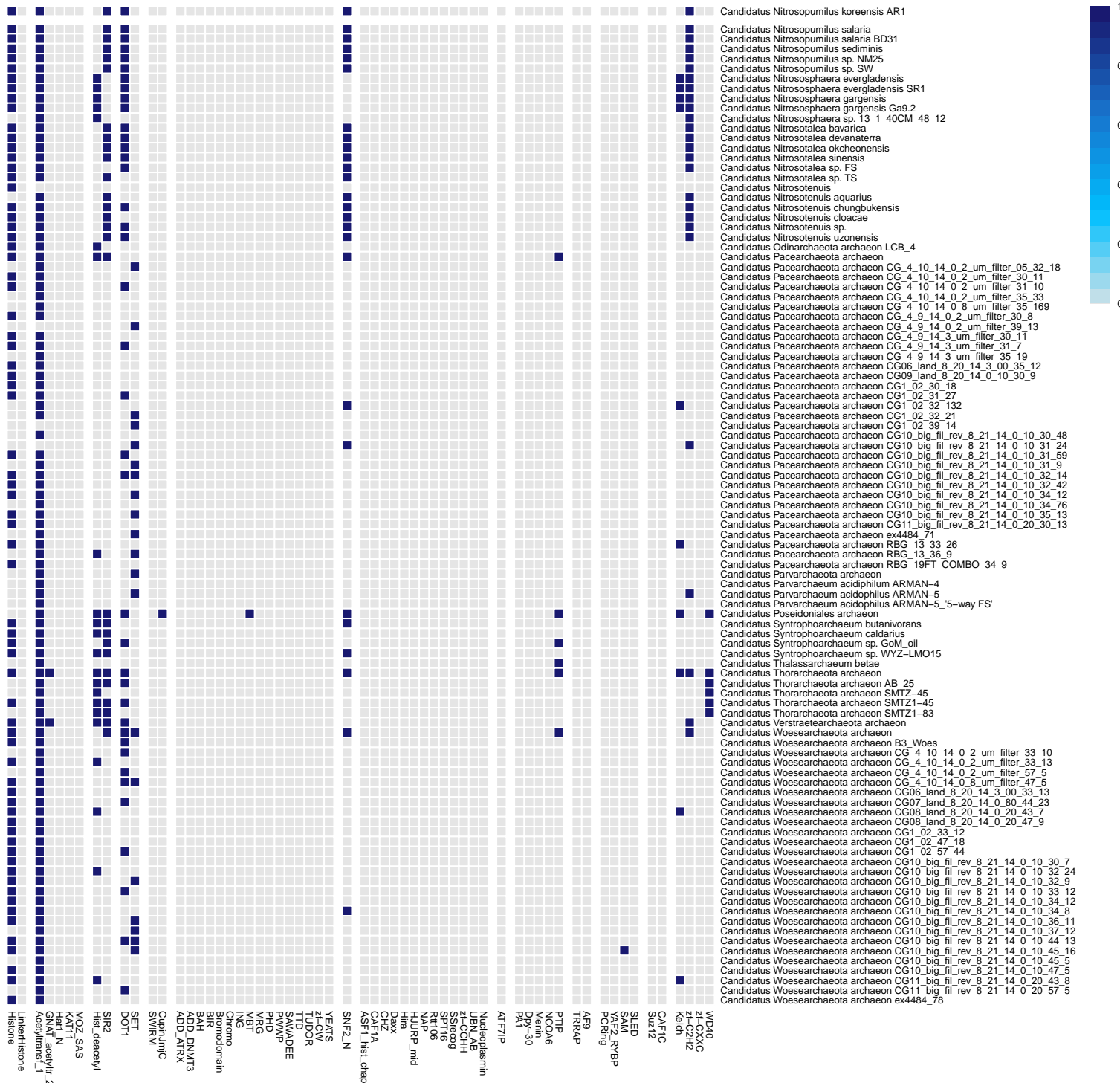
Gene presence per taxon in Archaea (superkingdom) (2/20) (fraction of species)



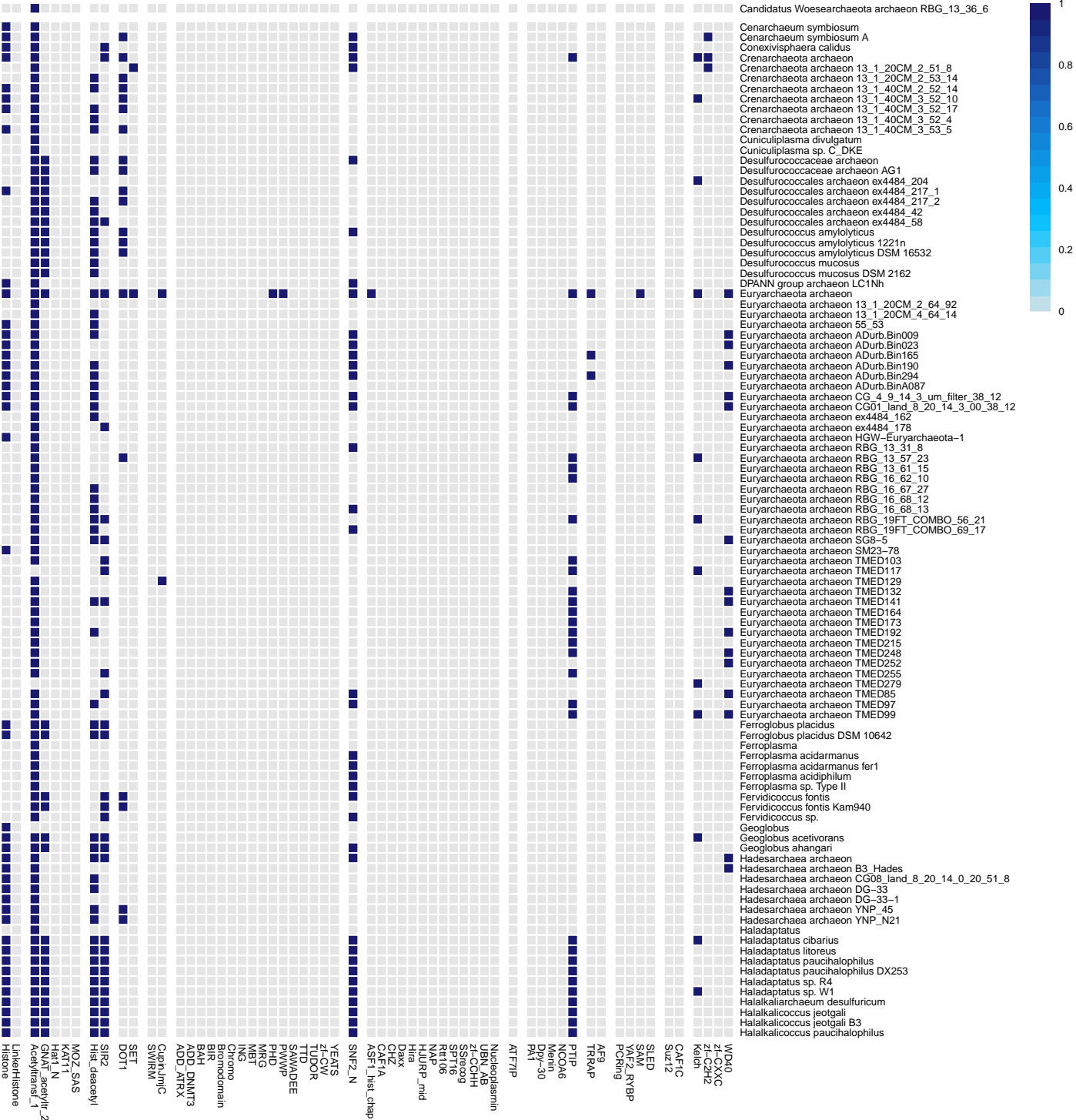
Gene presence per taxon in Archaea (superkingdom) (3/20) (fraction of species)

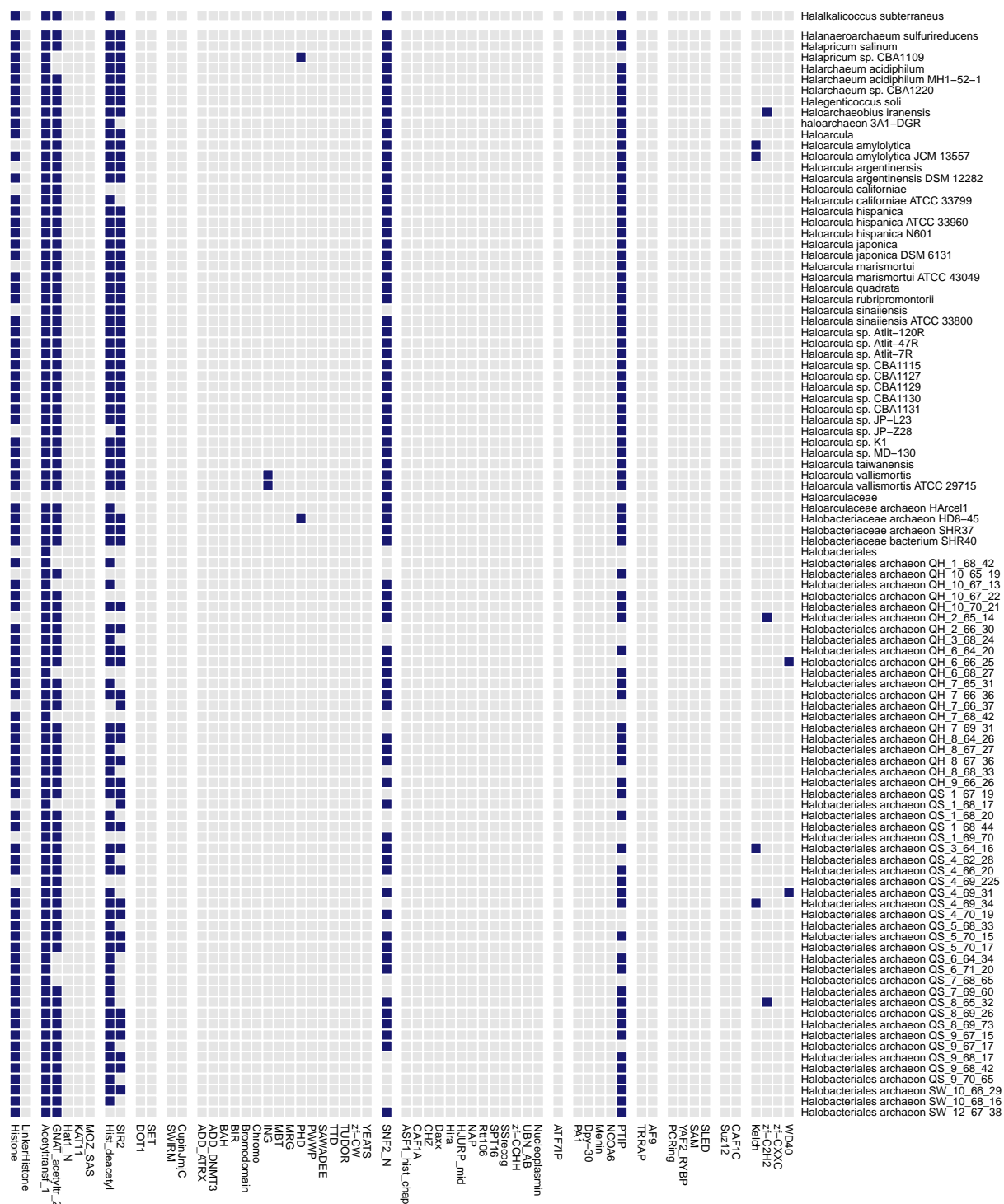


Gene presence per taxon in Archaea (superkingdom) (4/20) (fraction of species)

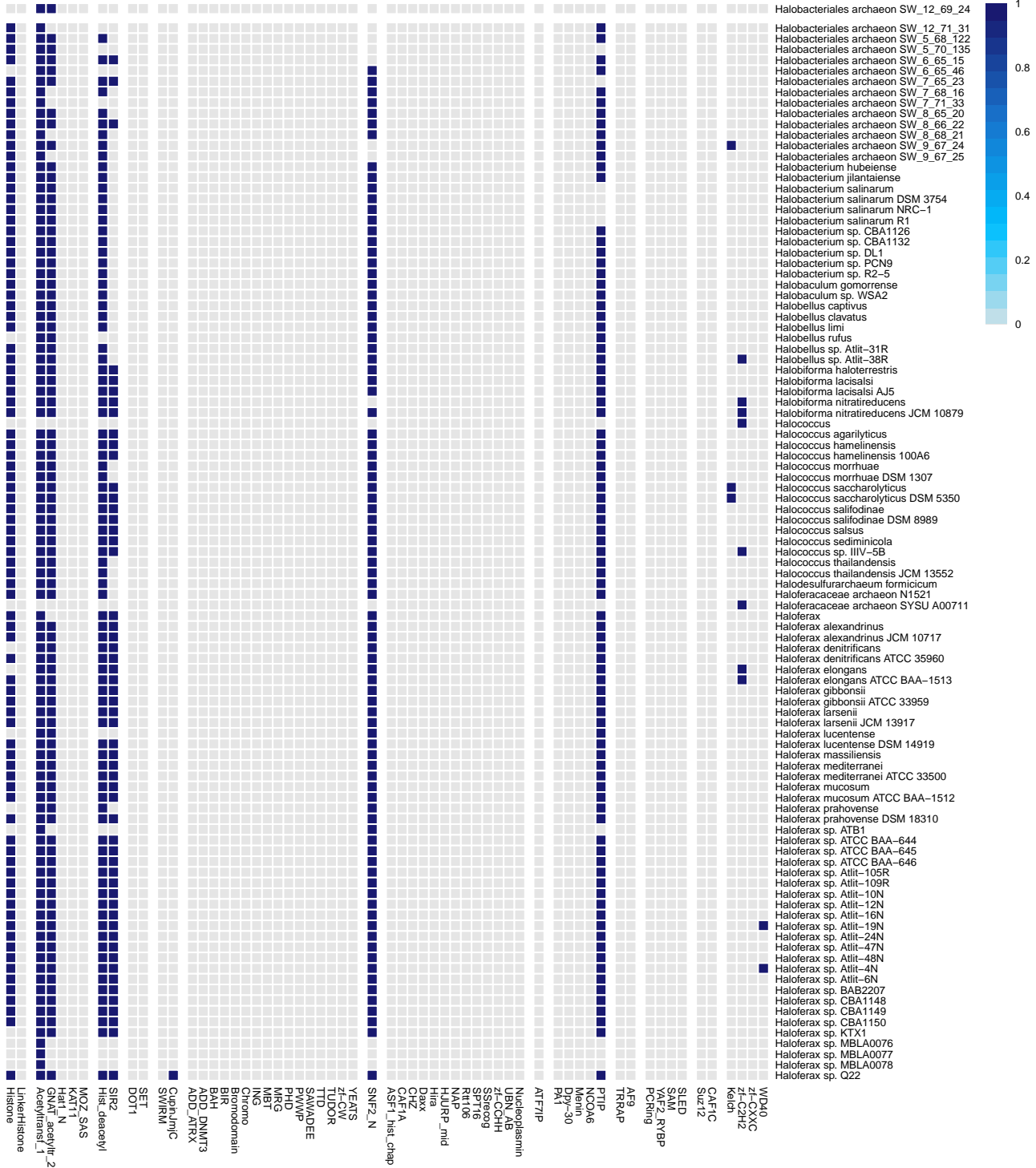


Gene presence per taxon in Archaea (superkingdom) (5/20) (fraction of species)

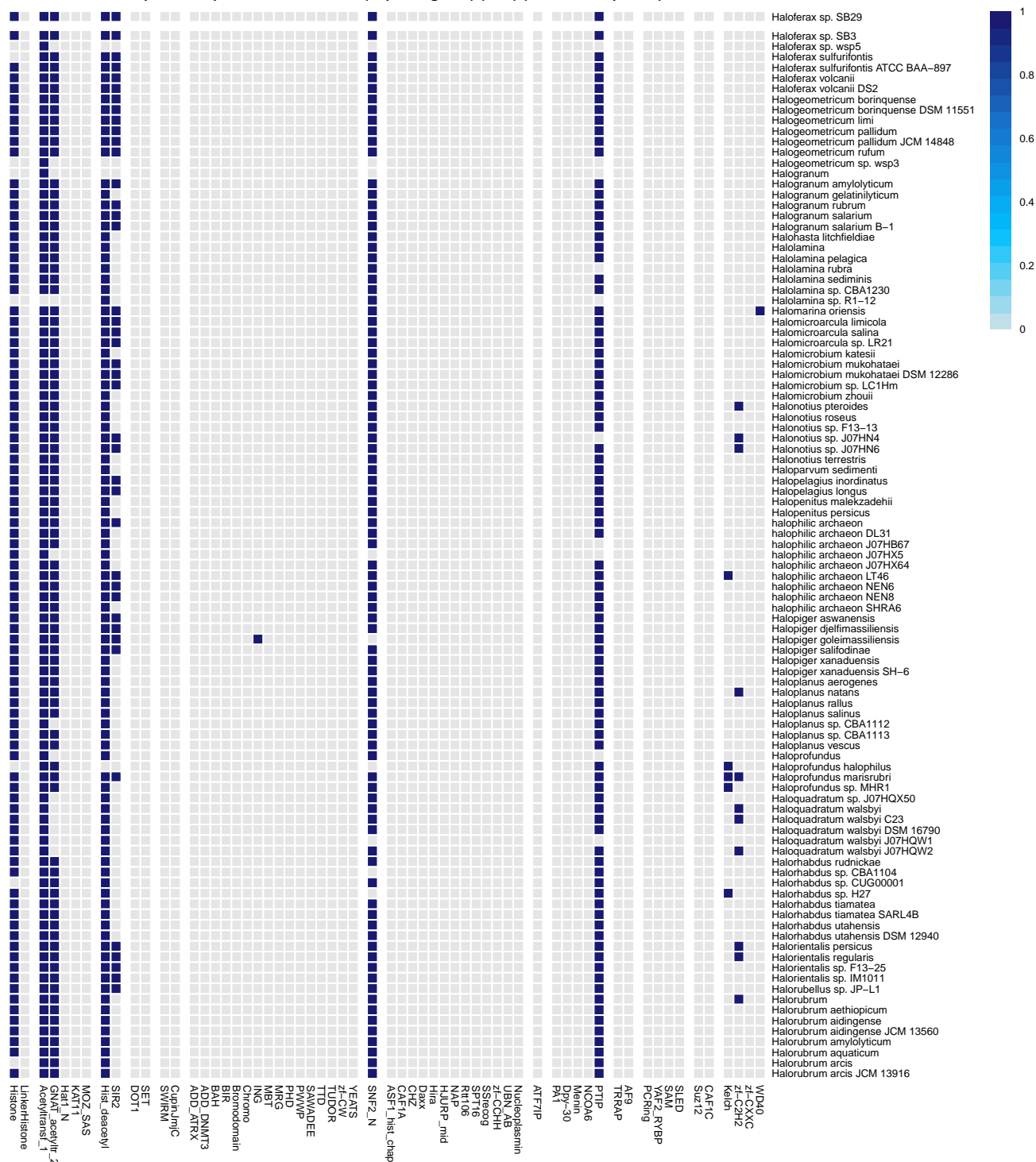




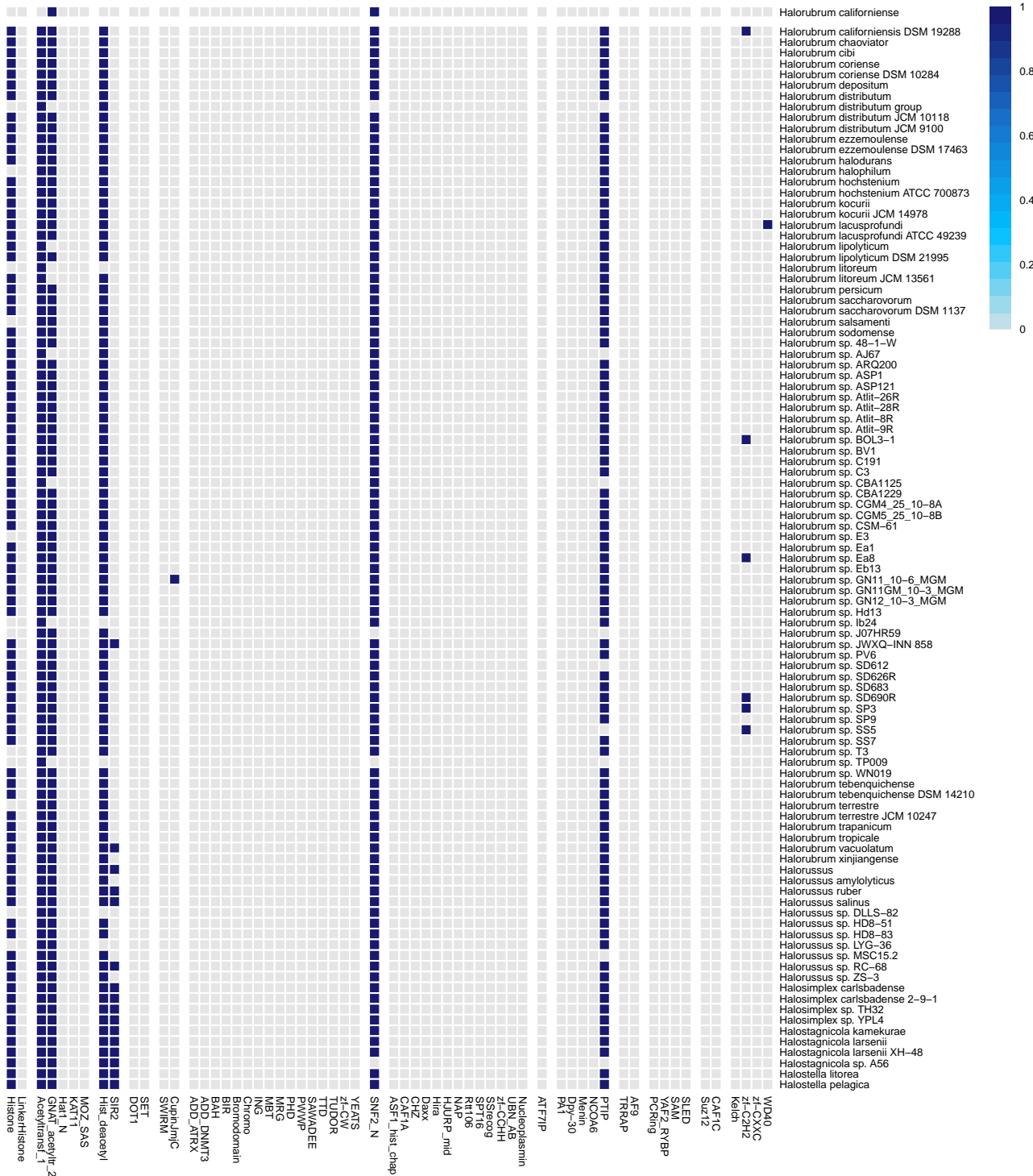
Gene presence per taxon in Archaea (superkingdom) (7/20) (fraction of species)



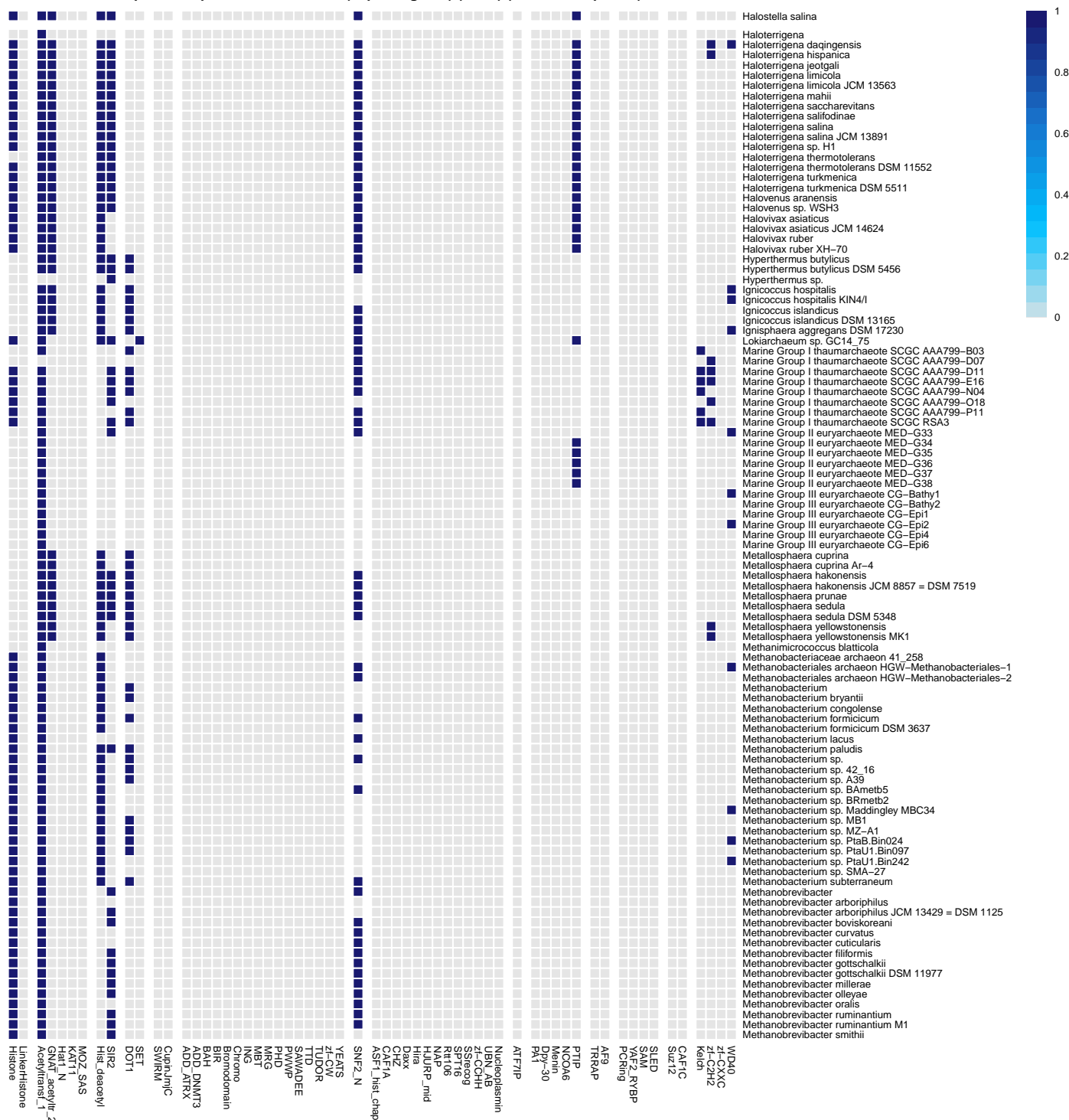
Gene presence per taxon in Archaea (superkingdom) (8/20) (fraction of species)



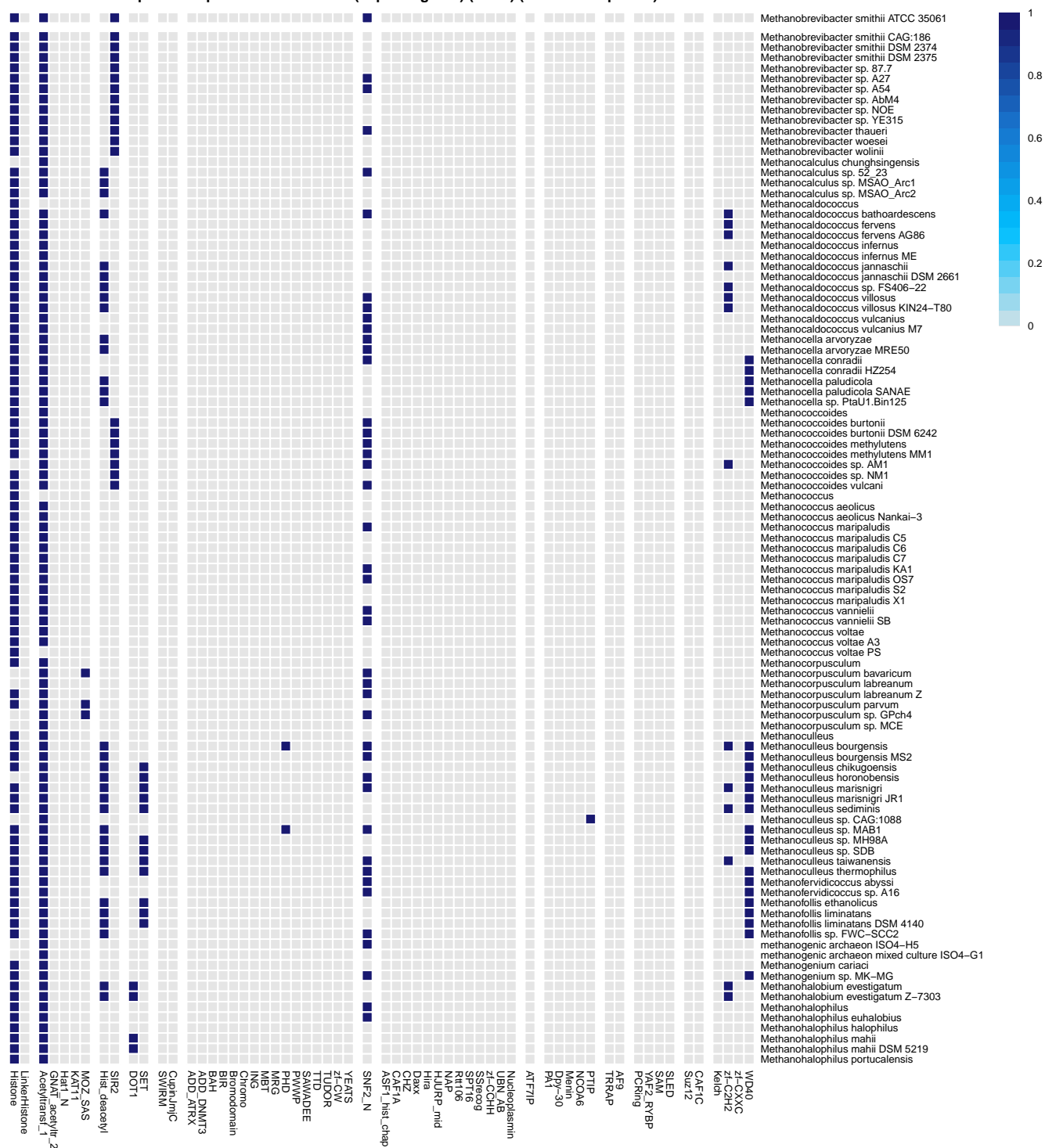
Gene presence per taxon in Archaea (superkingdom) (9/20) (fraction of species)



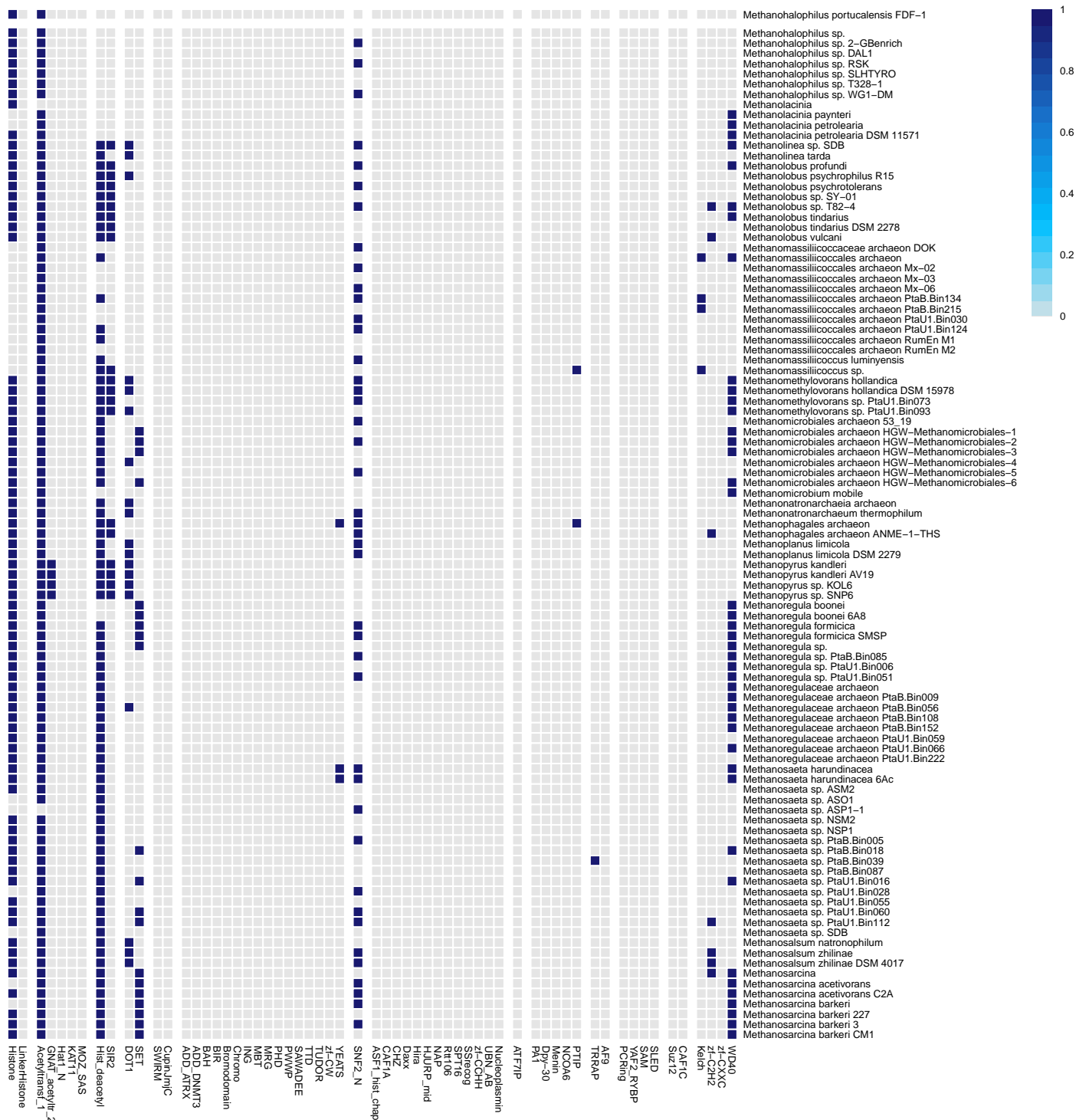
Gene presence per taxon in Archaea (superkingdom) (10/20) (fraction of species)



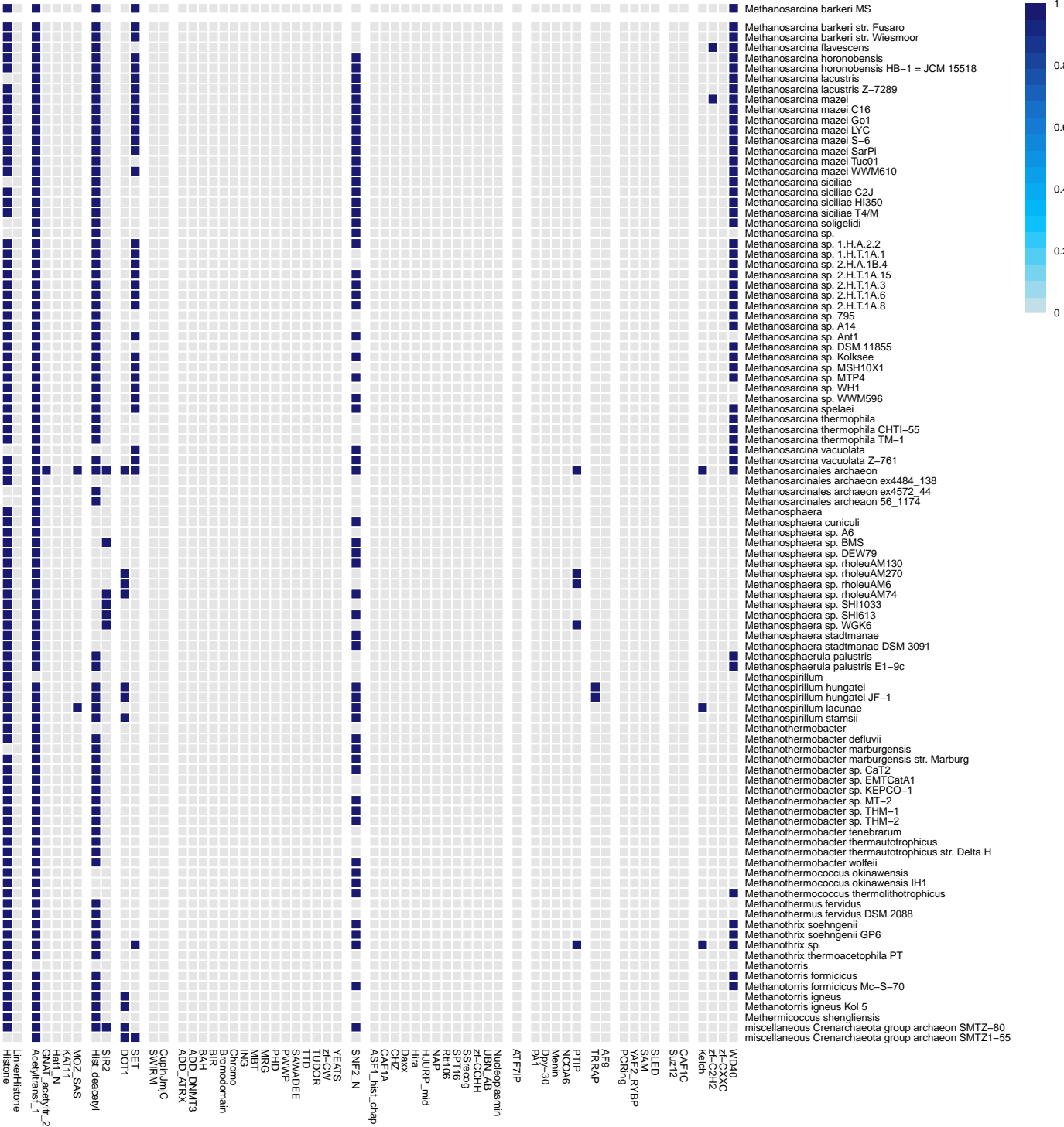
Gene presence per taxon in Archaea (superkingdom) (11/20) (fraction of species)



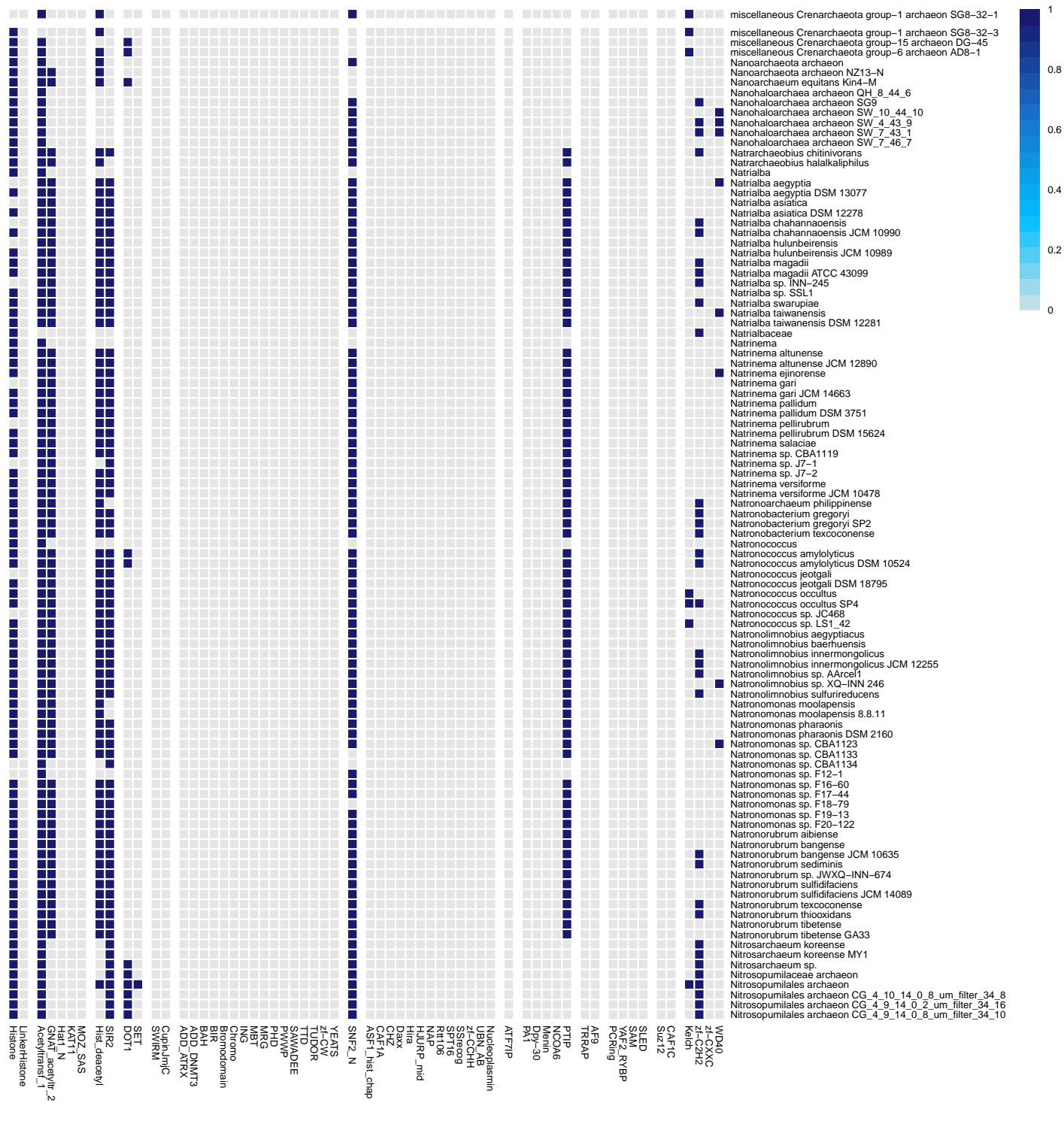
Gene presence per taxon in Archaea (superkingdom) (12/20) (fraction of species)



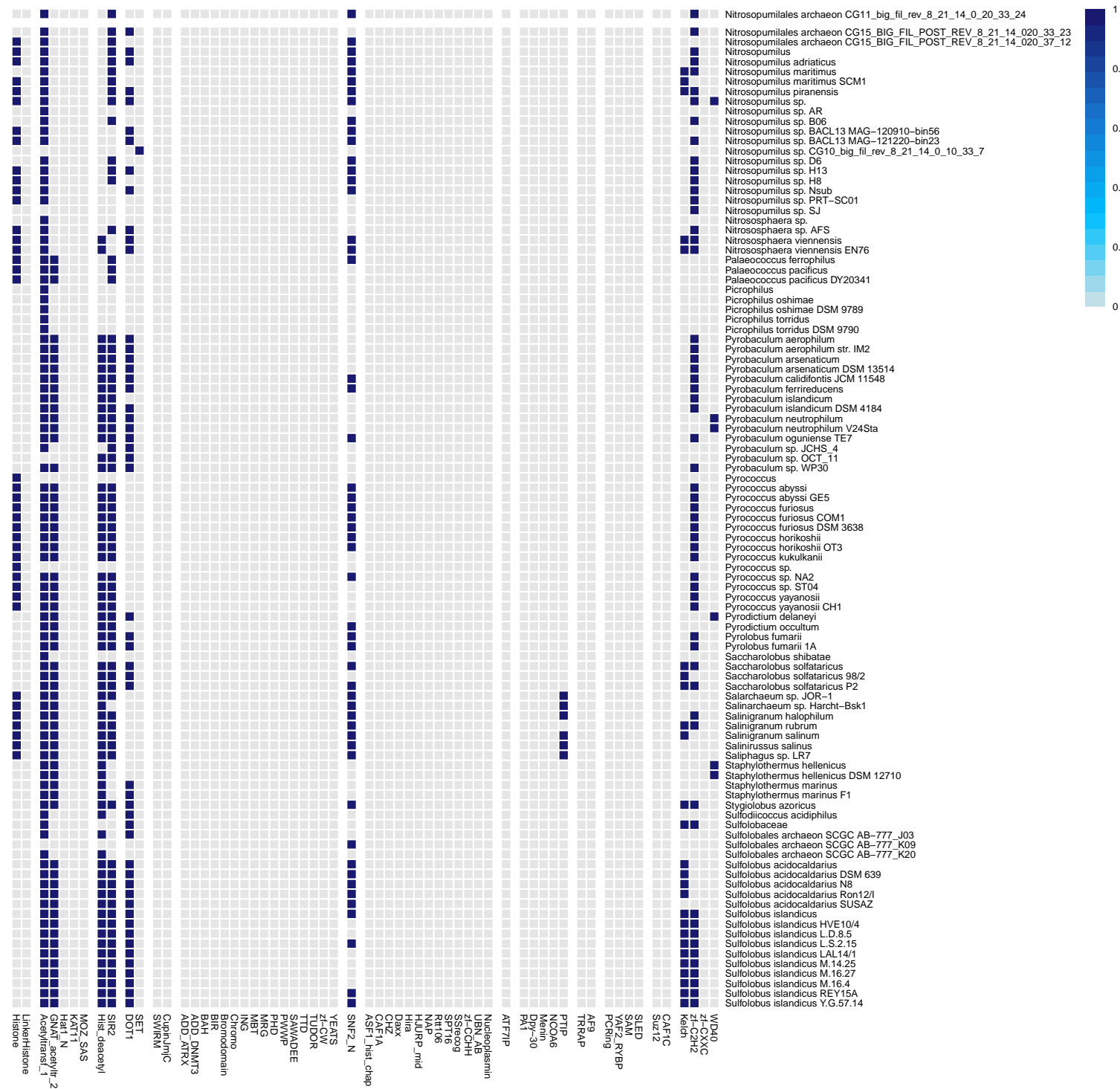
Gene presence per taxon in Archaea (superkingdom) (13/20) (fraction of species)



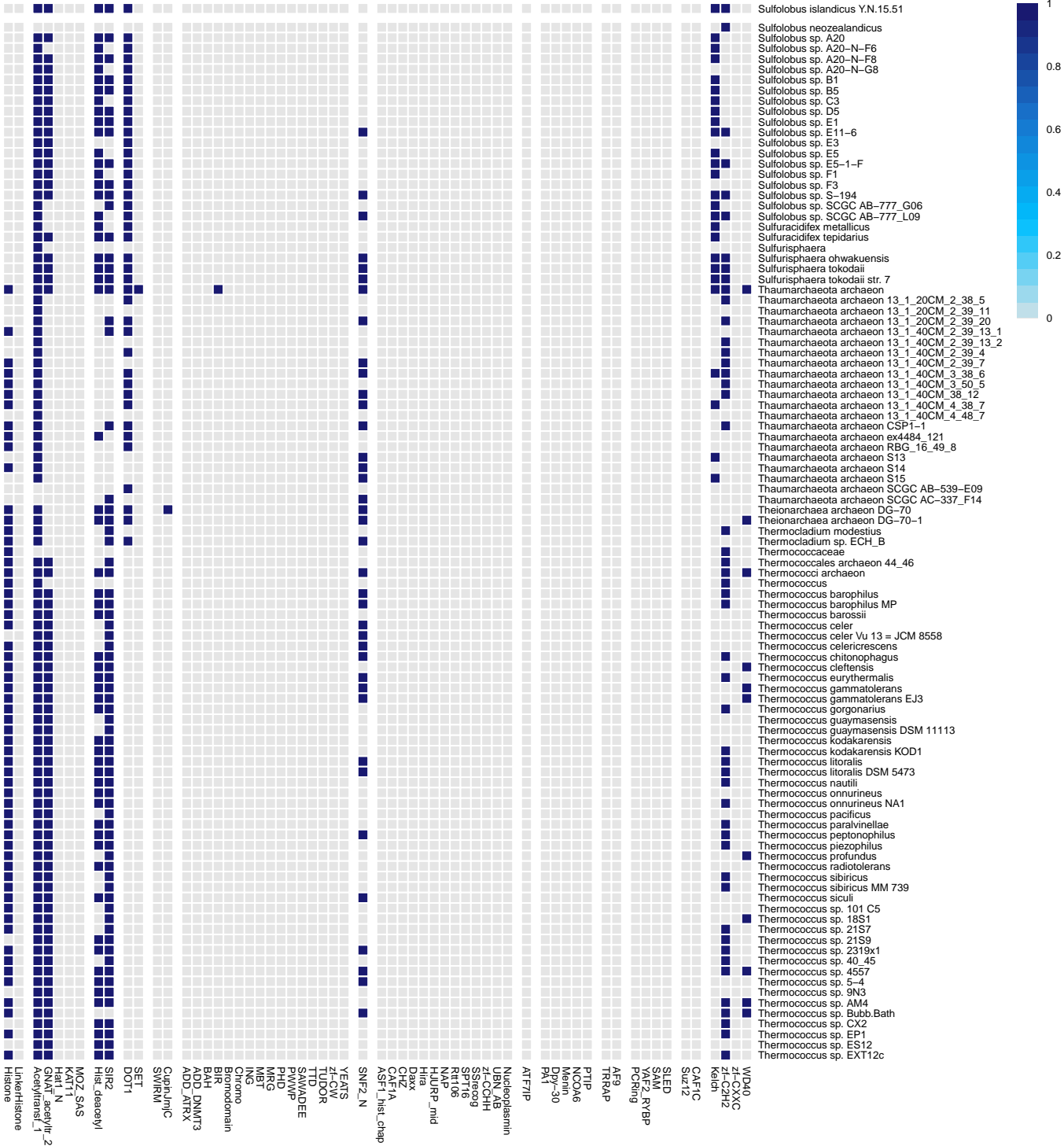
Gene presence per taxon in Archaea (superkingdom) (14/20) (fraction of species)



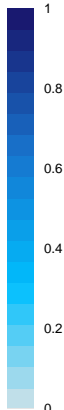
Gene presence per taxon in Archaea (superkingdom) (15/20) (fraction of species)



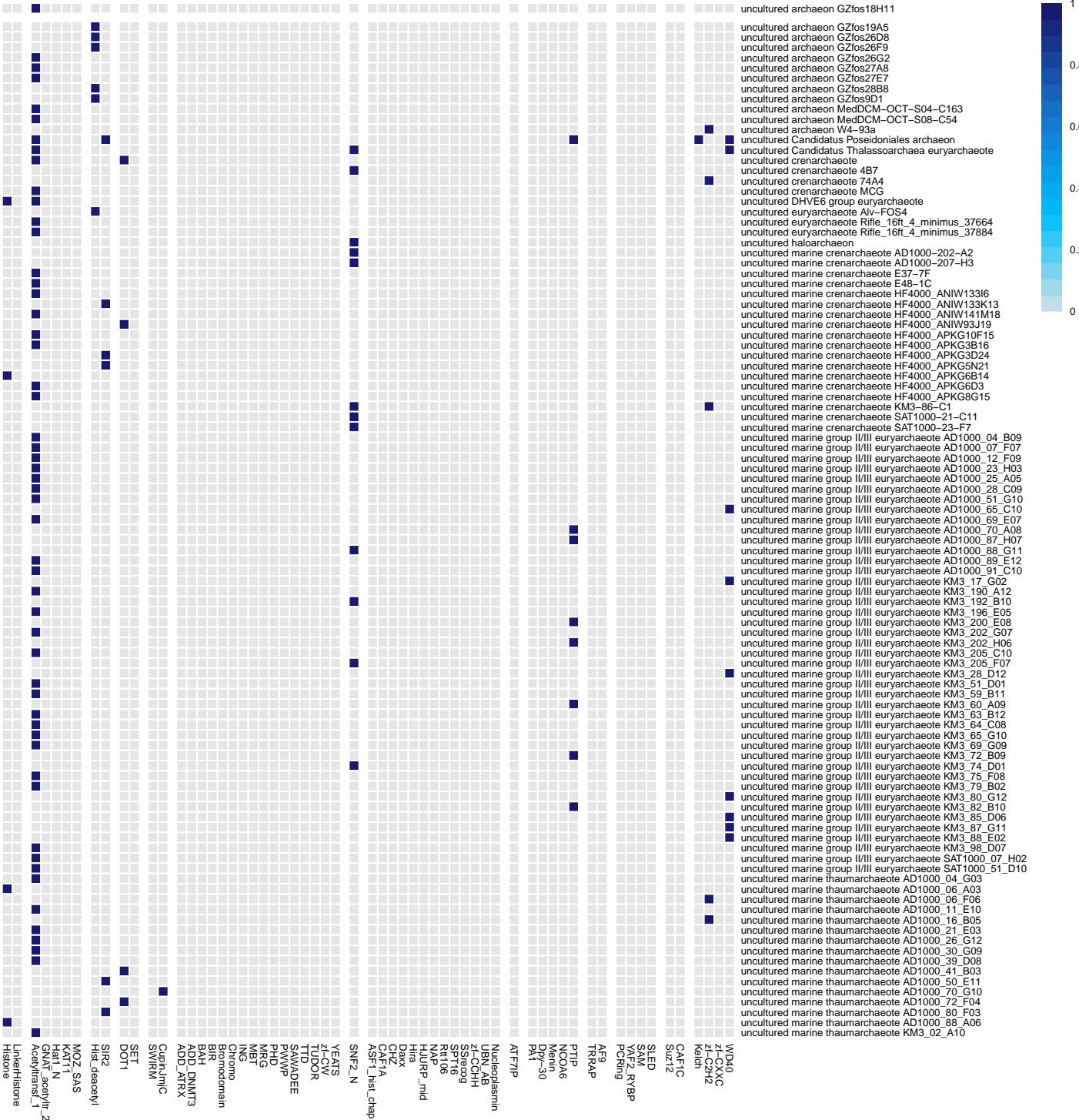
Gene presence per taxon in Archaea (superkingdom) (16/20) (fraction of species)



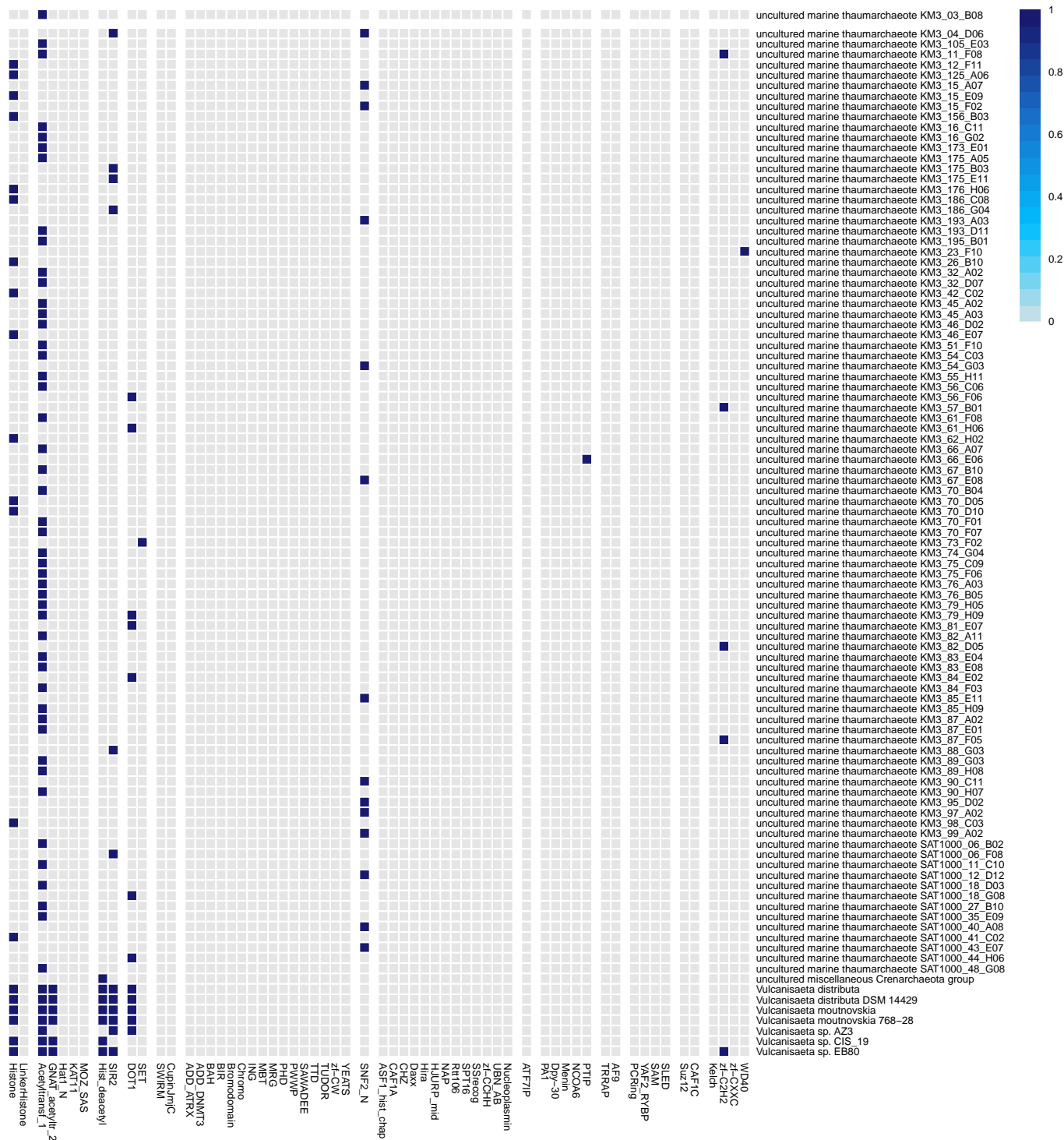
Thermococcus sp. GR4

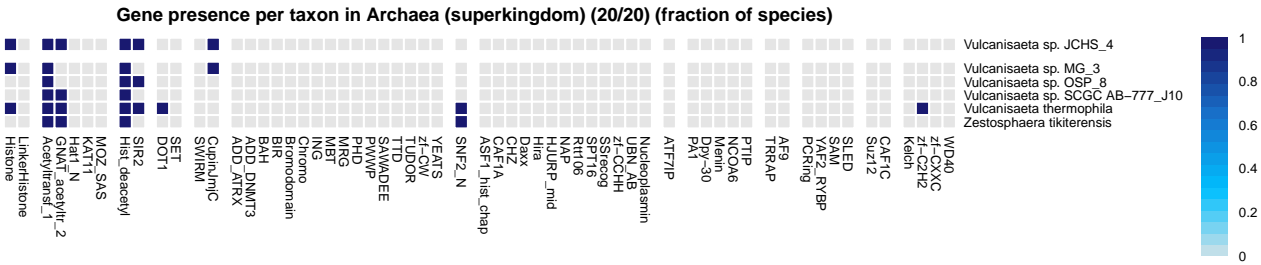


Gene presence per taxon in Archaea (superkingdom) (18/20) (fraction of species)



uncultured marine thaumarchaeote KM3 03 B08





A vertical number line with tick marks at 0, 1000, 2000, 3000, and 4000.

