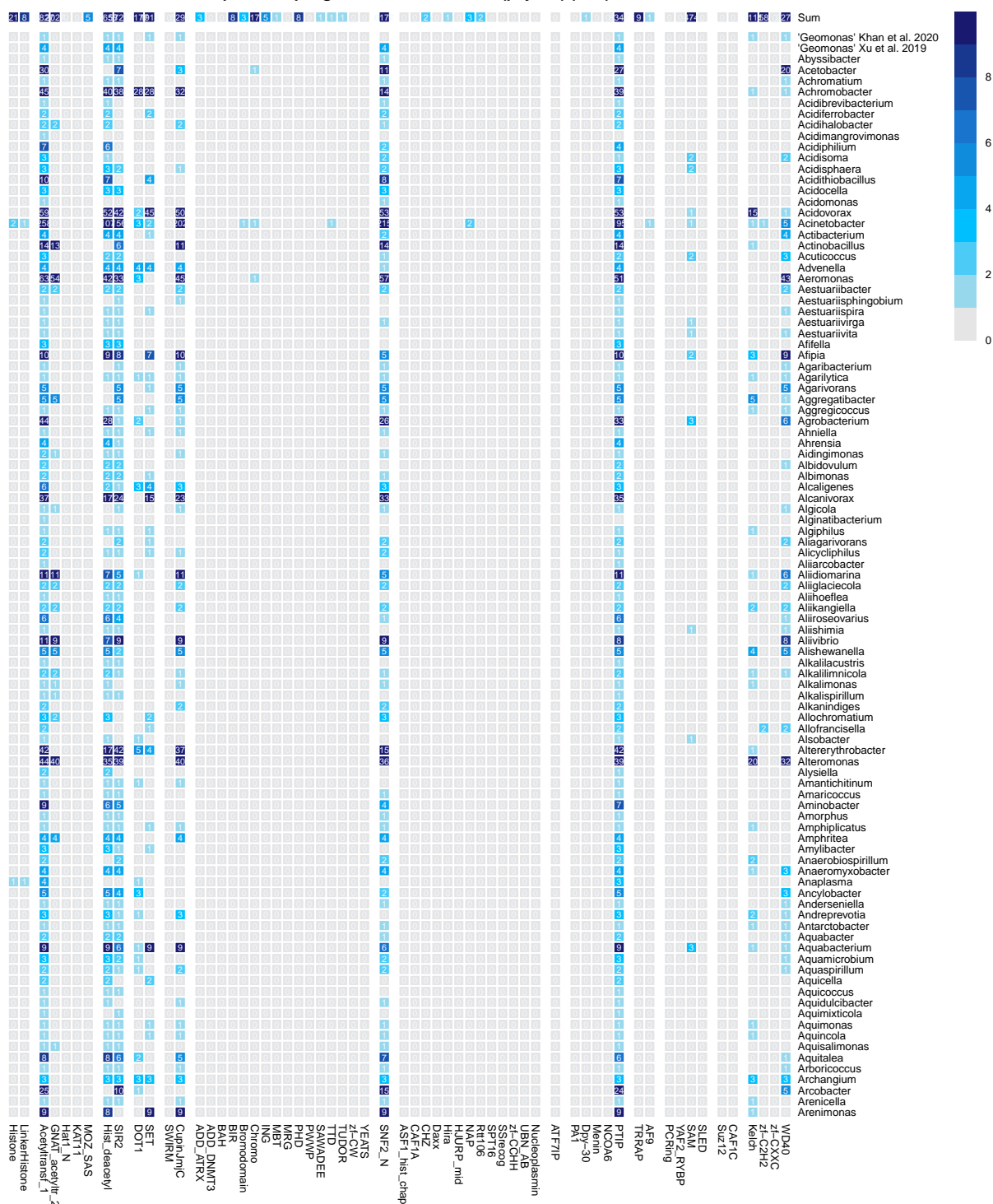
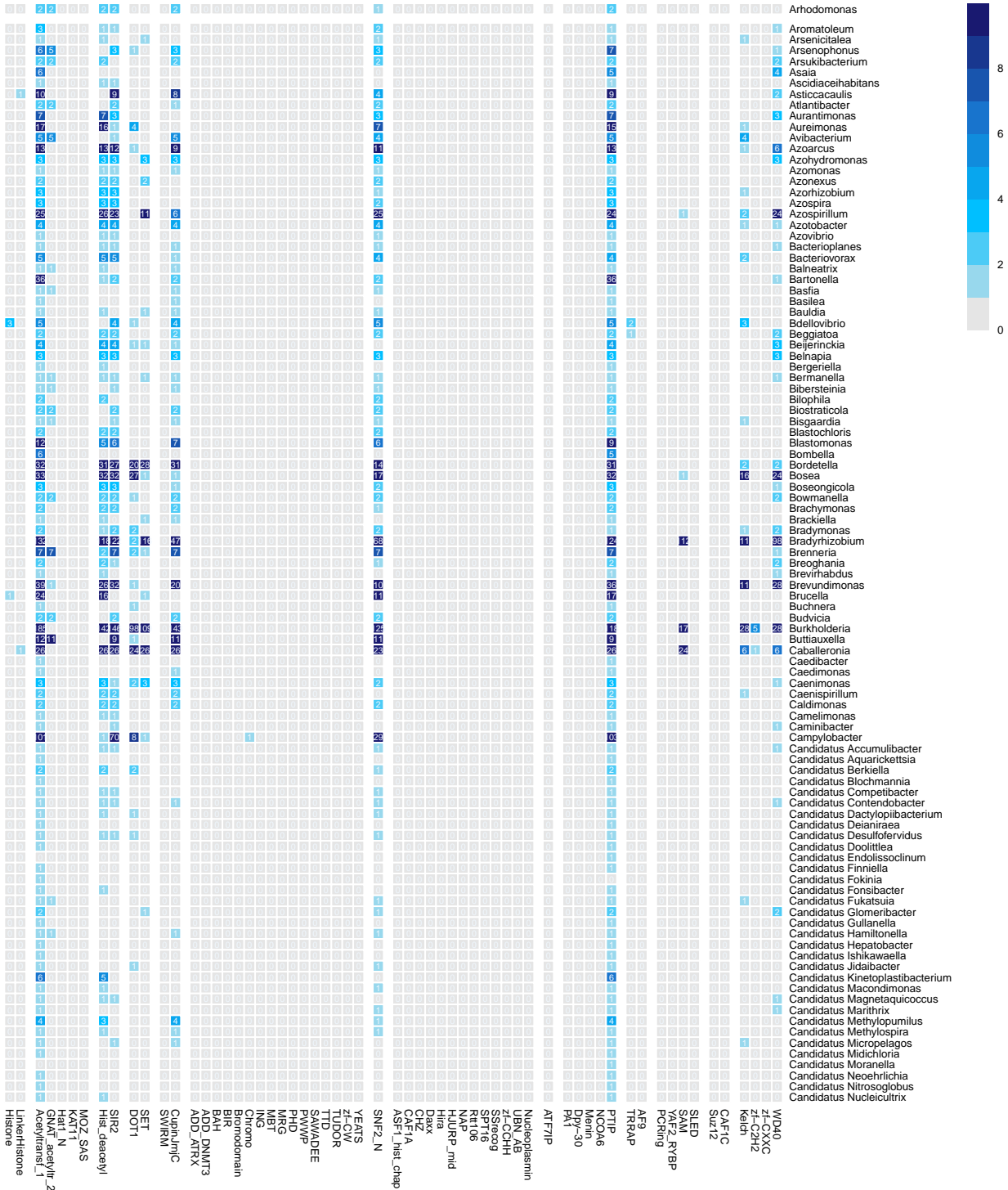


Gene presence per genus in Proteobacteria (phylum) (1/12)



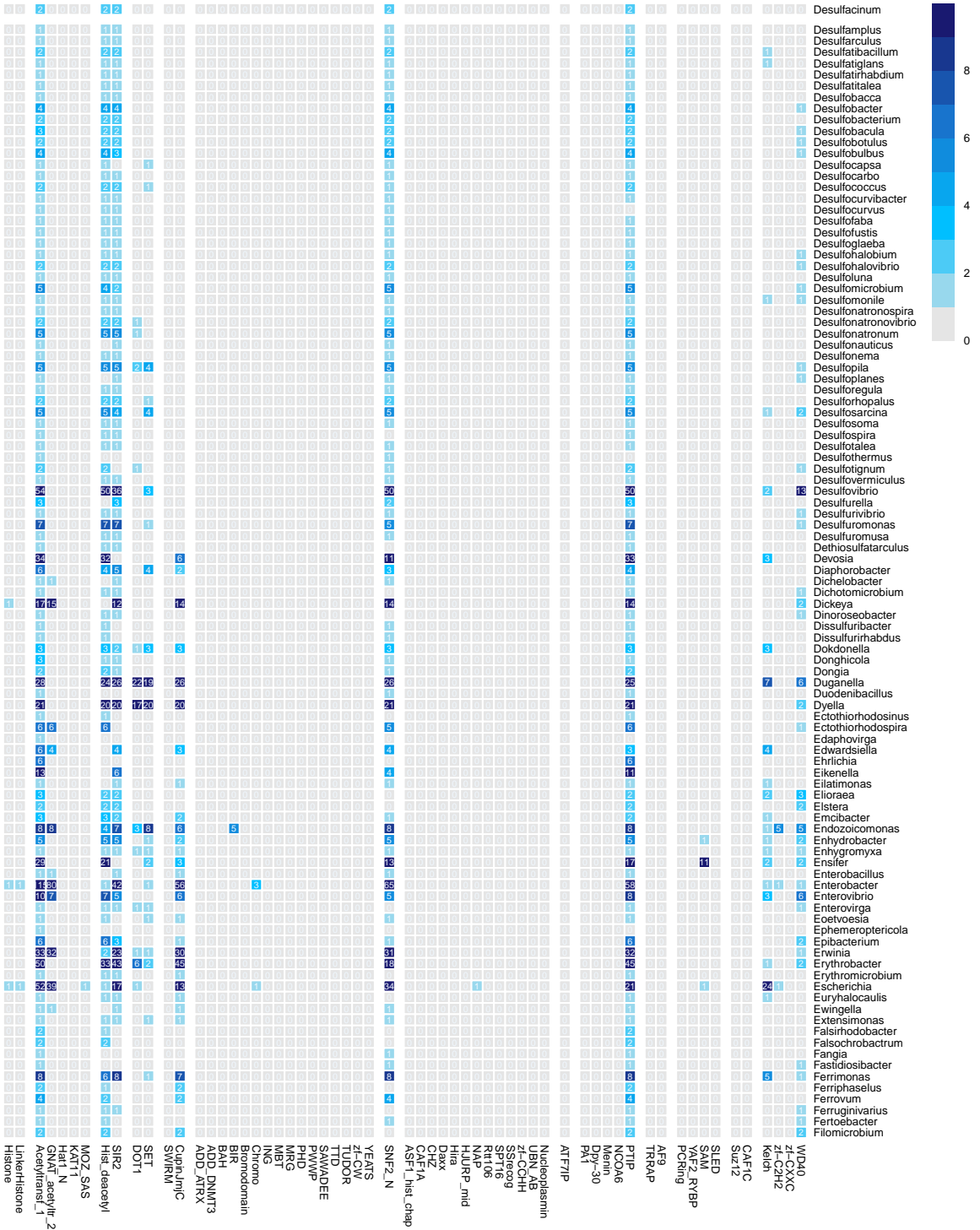
Gene presence per genus in Proteobacteria (phylum) (2/12)



Gene presence per genus in Proteobacteria (phylum) (3/12)



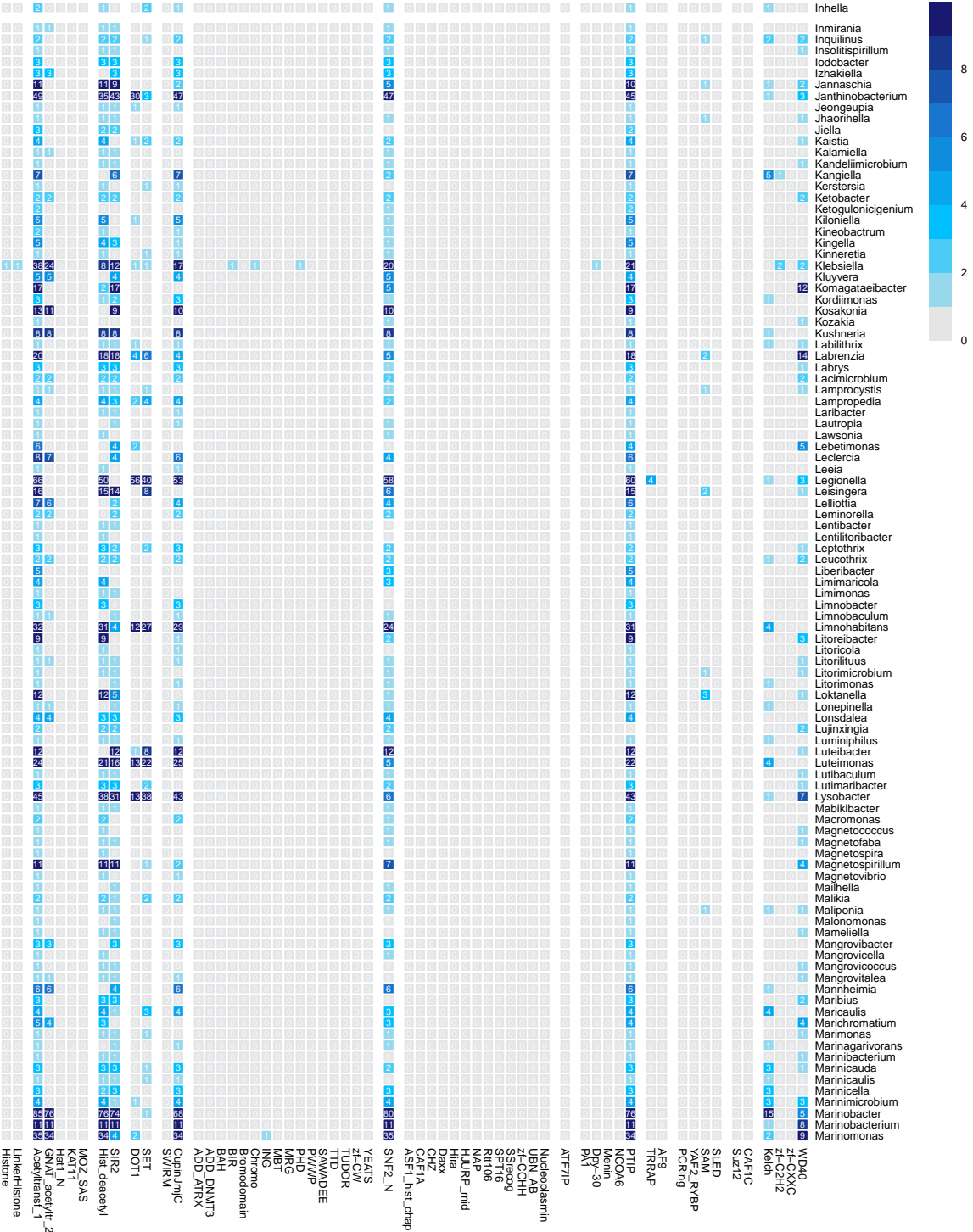
Gene presence per genus in Proteobacteria (phylum) (4/12)



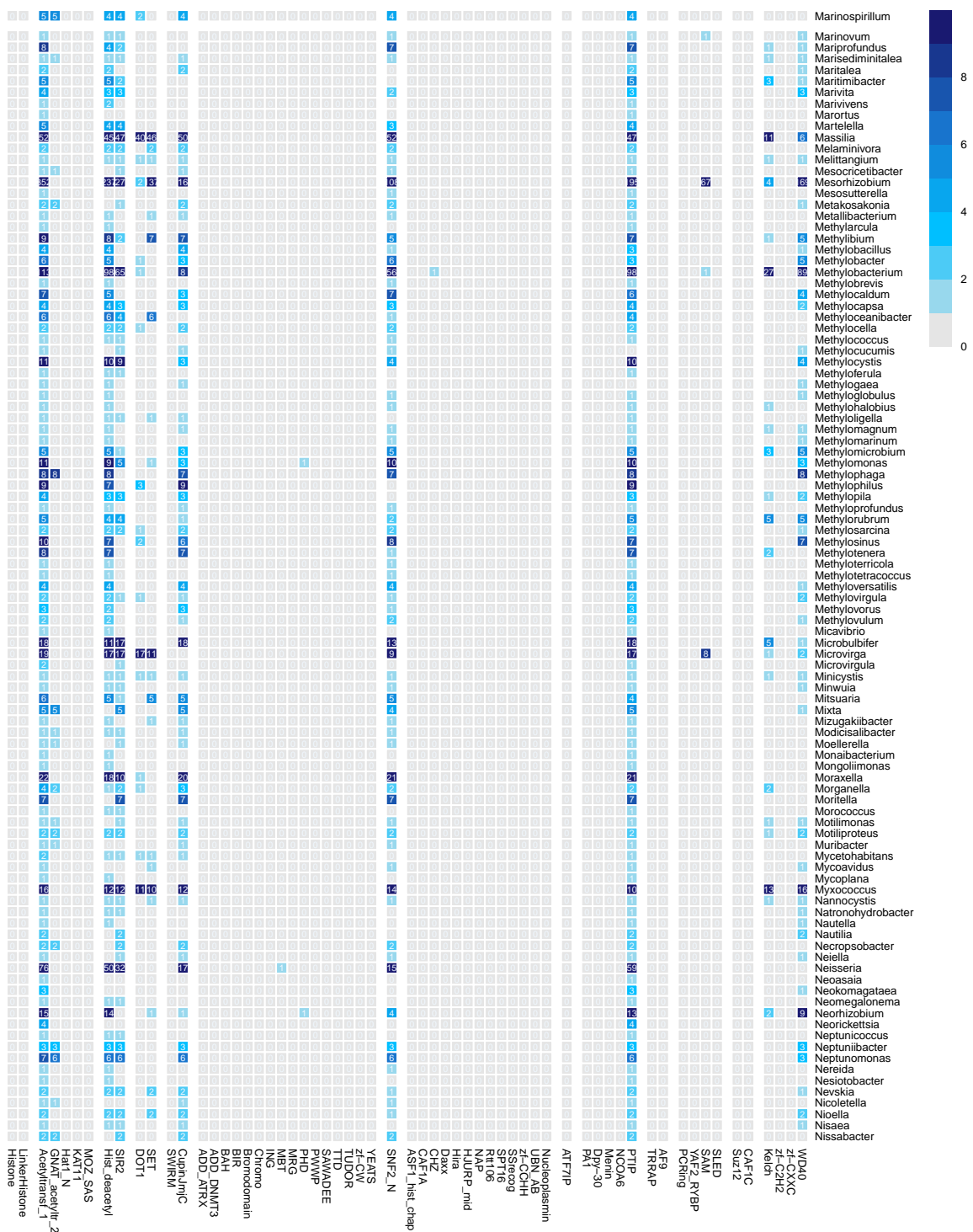
Gene presence per genus in Proteobacteria (phylum) (5/12)



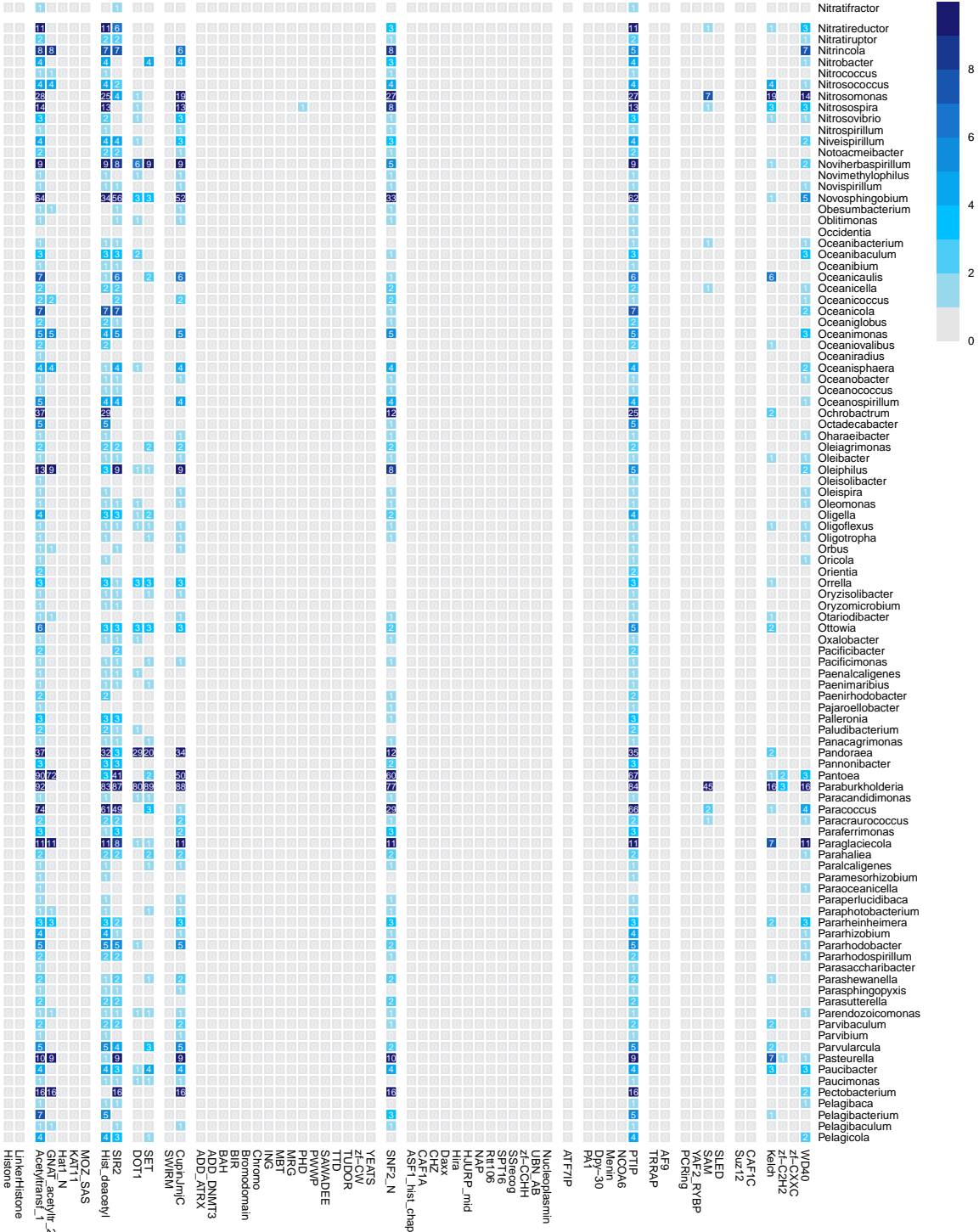
Gene presence per genus in Proteobacteria (phylum) (6/12)



**Gene presence per genus in Proteobacteria (phylum) (7/12)**

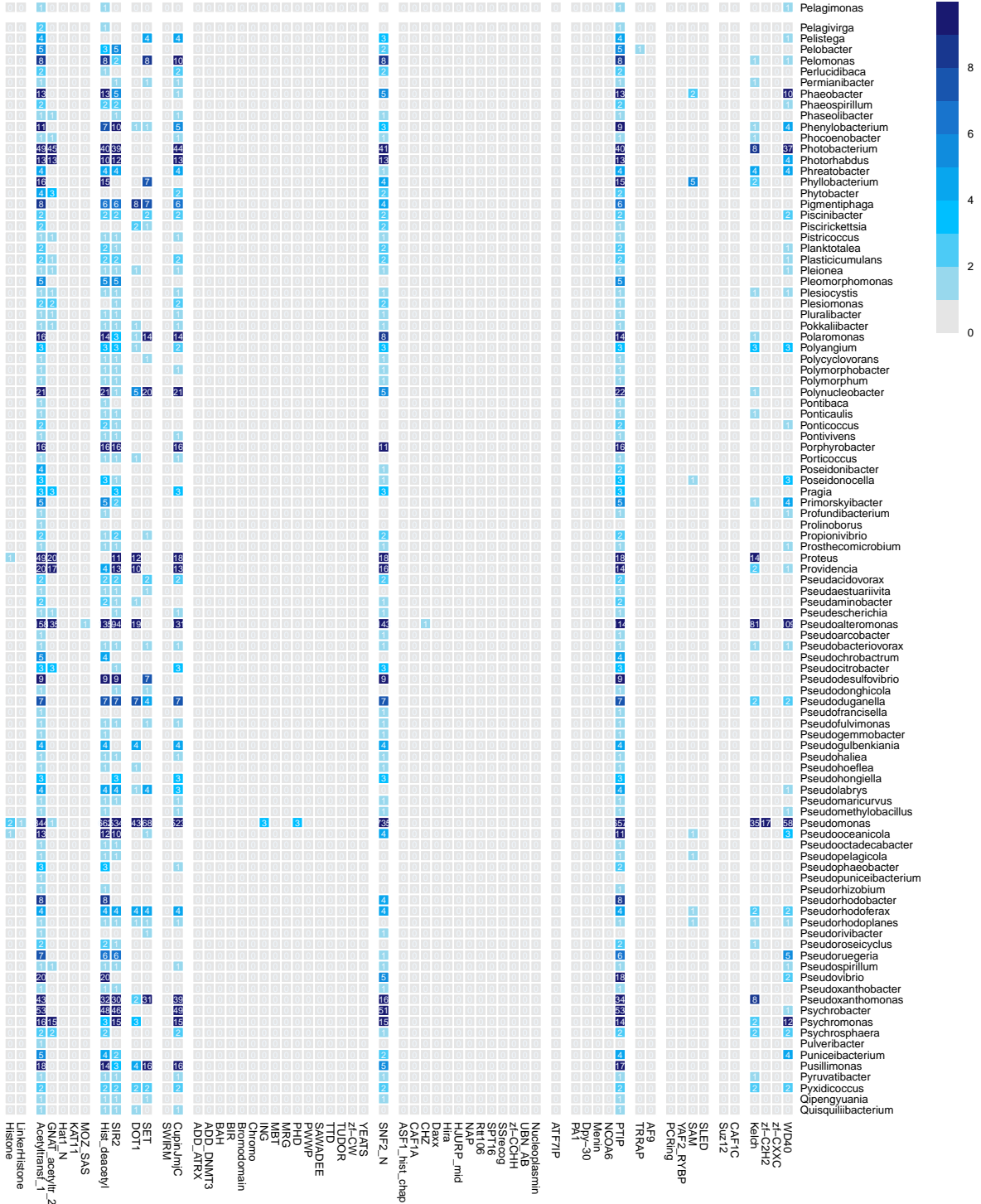


Gene presence per genus in Proteobacteria (phylum) (8/12)





Gene presence per genus in Proteobacteria (phylum) (9/12)



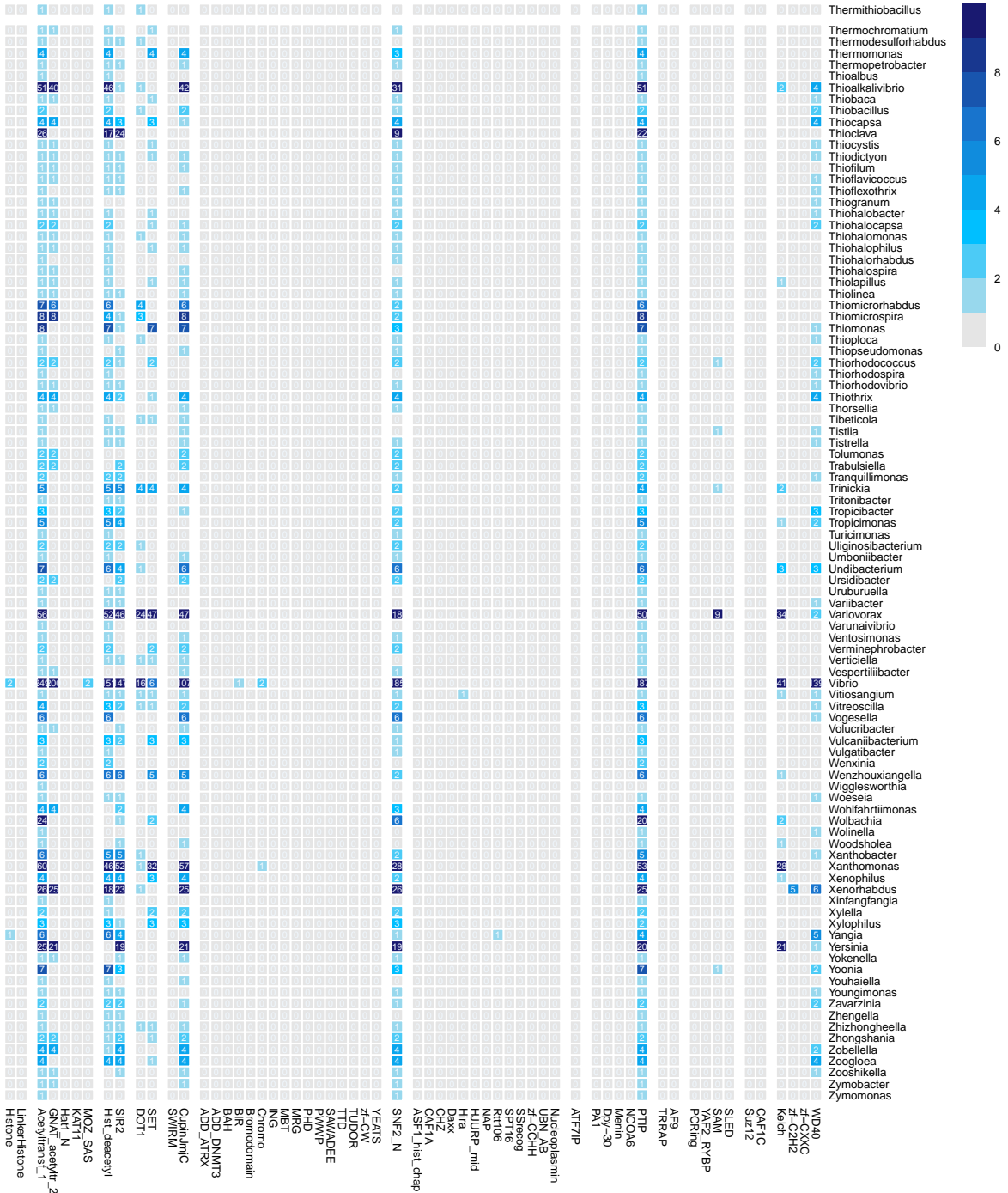
Gene presence per genus in Proteobacteria (phylum) (10/12)



Gene presence per genus in Proteobacteria (phylum) (11/12)



Gene presence per genus in Proteobacteria (phylum) (12/12)



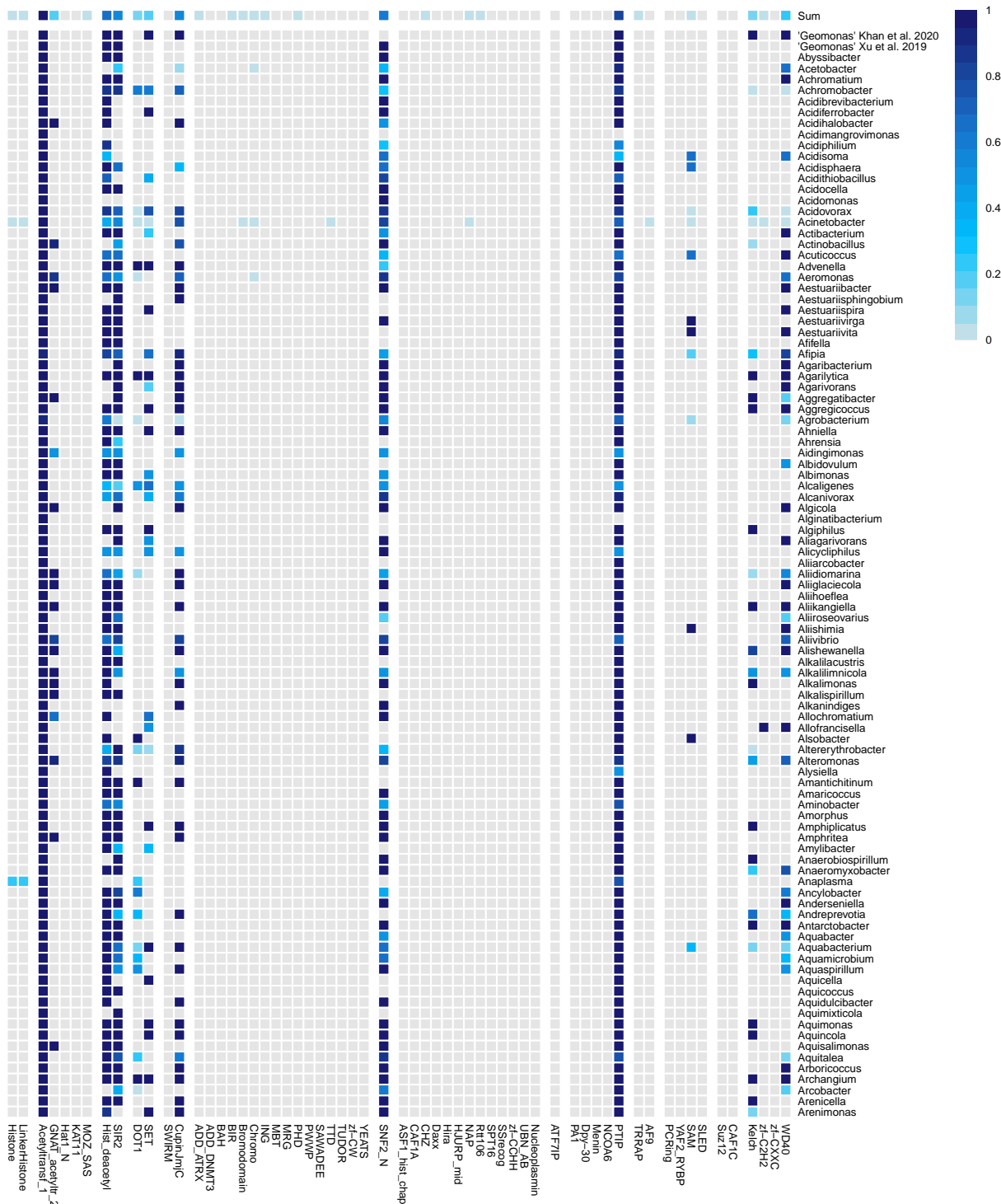
Heatmap showing the relative abundance of 100 bacterial taxa across 100 samples. The taxa are listed on the right, and the samples are listed on the bottom. A color scale on the right indicates relative abundance from 0 (light blue) to 1 (dark blue).

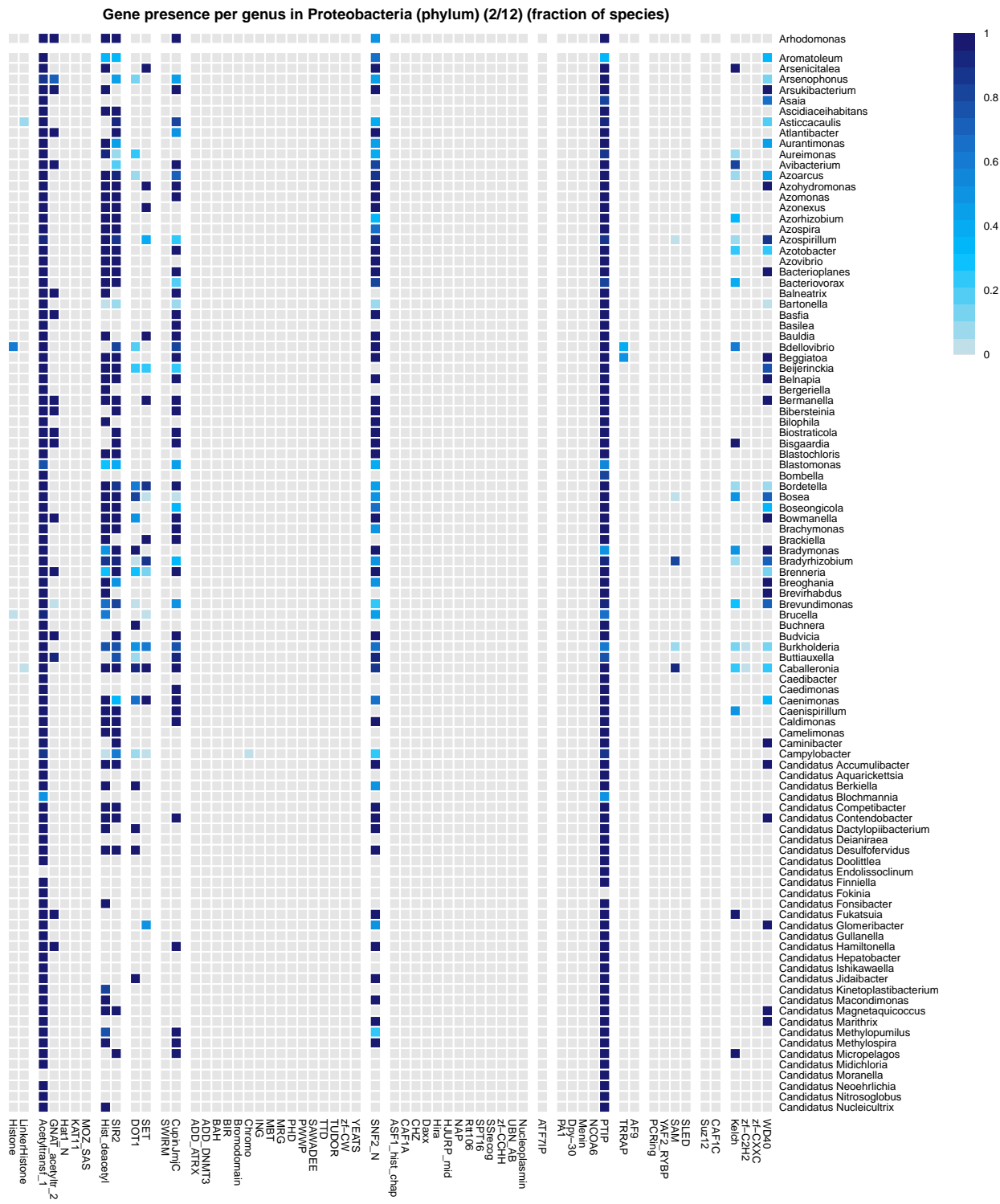
**Taxa (Right):**

- 'Geomonas' Khan et al. 2020
- 'Geomonas' Xu et al. 2019
- Abyssibacter
- Acetobacter
- Achromatium
- Achromobacter
- Acidibrevibacterium
- Acidiferrrobacter
- Acidihalobacter
- Acidimangroviimonas
- Acidiphilium
- Acidisoma
- Acidispheara
- Acidithiobacillus
- Acidocella
- Acidomonas
- Acidovorax
- Acinetobacter
- Actinobacterium
- Actinobacillus
- Acuticoccus
- Advenella
- Aeromonas
- Aestuariibacter
- Aestuariasphingobium
- Aestuariuspira
- Aestuariusvira
- Aestuariusvita
- Aifella
- Alfia
- Agaribacterium
- Agarolytica
- Agarivorans
- Aggregatibacter
- Aggregicoccus
- Agrobacterium
- Ahniella
- Ahrensia
- Aidingimonas
- Albidovulum
- Albimonas
- Alcaligenes
- Alcanivorax
- Algicola
- Alginatibacterium
- Algiphilus
- Aliagarivorans
- Allicyclophilus
- Aliarobacter
- Aliidiomarina
- Aliiglaciecola
- Aliihoeflea
- Aliikangiella
- Aliiroseovarius
- Aliishimia
- Aliivibrio
- Alishewanella
- Alkalilacustris
- Alkalilimnicola
- Alkalimonas
- Alkalispirillum
- Alkanindiges
- Allochromatium
- Allofrancisella
- Alsobacter
- Altererythrobacter
- Alteromonas
- Alysia
- Amantichitinum
- Amaricoccus
- Aminobacter
- Amorphus
- Amphiplicatus
- Amphritea
- Amylibacter
- Anaerobiospirillum
- Anaeromyxobacter
- Anaplasma
- Ancyllobacter
- Andersenella
- Andreprevotia
- Antarctobacter
- Aquabacter
- Aquabacterium
- Aquamicrobium
- Aquaspirillum
- Aquicella
- Aquicoccus
- Aquidulcibacter
- Aquimixticola
- Aquimonas
- Aquincola
- Aquisalimonas
- Aquitalea
- Arboricoccus
- Archangium
- Arcobacter
- Arenicella
- Arenimonas

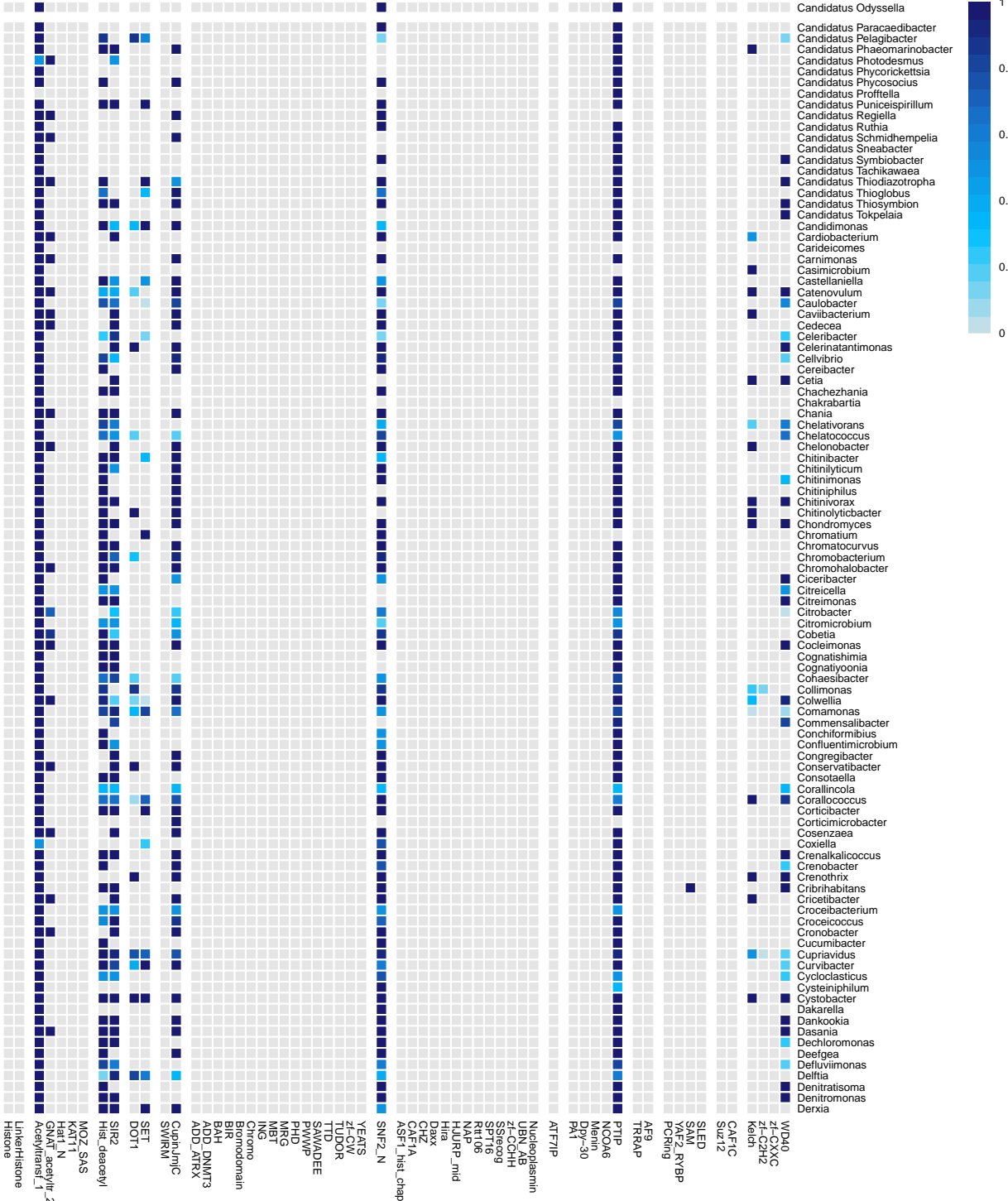
**Samples (Bottom):**

- WD40
- 2F-QXG
- 2F-CH2
- Keoh
- CA-FIC
- Su-12
- SLED
- SAM
- VA-F2-RVP
- PcKing
- AF9
- TRRAP
- PTP
- NCOA6
- Mentm
- Dpy-30
- PA1
- AT7TP
- Nucleoplasm
- UBN\_AB
- UBN\_COH
- SEBCH
- SPT16
- SPT19
- RH106
- NAP
- HJURP\_mid
- HJURP
- CHZ
- CHZ
- CA-F1
- AS-F1\_Inst\_chap
- SNF2\_N
- YEATS
- ZI-QW
- TTT-DOR
- TTT
- SAWADEE
- PVWP
- PHD
- MKG
- MKG
- MKG
- MKG
- Chromo
- Bromodomain
- BIR
- BAH
- ADNMT3
- ADN\_ATRX
- ADN\_ATRX
- CupN\_JnC
- SWRM
- MOZ\_SAS
- KAT1
- KAT1
- Chromatin
- Acetyrans\_1
- Acetyrans\_2
- Acetyrans\_3
- Acetyrans\_4
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- Acetyrans\_98
- Acetyrans\_99
- Acetyrans\_100





Gene presence per genus in Proteobacteria (phylum) (3/12) (fraction of species)



Heatmap showing the relative abundance of 100 bacterial taxa across 100 samples. The taxa are listed on the right, and the samples are listed on the bottom. The color scale ranges from 0 (light blue) to 1 (dark blue).

**Relative Abundance Scale:** 0 (light blue) to 1 (dark blue).

**Top 10 Bacterial Taxa (by relative abundance in sample 100):**

- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum

**Bottom 10 Bacterial Taxa (by relative abundance in sample 100):**

- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum
- Desulfatibaculum

**Sample Labels (Bottom):**

100, 99, 98, 97, 96, 95, 94, 93, 92, 91, 90, 89, 88, 87, 86, 85, 84, 83, 82, 81, 80, 79, 78, 77, 76, 75, 74, 73, 72, 71, 70, 69, 68, 67, 66, 65, 64, 63, 62, 61, 60, 59, 58, 57, 56, 55, 54, 53, 52, 51, 50, 49, 48, 47, 46, 45, 44, 43, 42, 41, 40, 39, 38, 37, 36, 35, 34, 33, 32, 31, 30, 29, 28, 27, 26, 25, 24, 23, 22, 21, 20, 19, 18, 17, 16, 15, 14, 13, 12, 11, 10, 9, 8, 7, 6, 5, 4, 3, 2, 1.





Heatmap showing the relative abundance of 100 bacterial taxa across 100 samples. The taxa are listed on the right, and the samples are listed on the left. The color scale ranges from 0 (light blue) to 1 (dark blue).

**Flavimaricola**

**Fluoribacter**

**Fluviibacterium**

**Fluviicoccus**

**Fluviispira**

**Fodinicurvata**

**Fontimonas**

**Formivibrio**

**Francisella**

**Franconibacter**

**Frateuria**

**Frederiksenia**

**Frigidibacter**

**Frischella**

**Fulvimarina**

**Fulvimonas**

**Gallaecimonas**

**Gallibacterium**

**Gallionella**

**Gayadomonas**

**Geminicoccus**

**Gemmobacter**

**Geoalkalibacter**

**Geobacter**

**Geopsychrobacter**

**Georhizobium**

**Geothermobacter**

**Ghiorsea**

**Gibbsiella**

**Giesbergeria**

**Gilliamella**

**Gilvamarinus**

**Glaciicola**

**Glaciimonas**

**Glaesserella**

**Gluconacetobacter**

**Gluconobacter**

**Glycocalyx**

**Granulibacter**

**Granulosicoccus**

**Grimontella**

**Grimontia**

**Gulbenkiania**

**Guyarkeria**

**Gyruella**

**Haematobacter**

**Haematospirillum**

**Haemophilus**

**Haefia**

**Halbella**

**Halangium**

**Haliea**

**Halioglobus**

**Halobacteriovorax**

**Halocynthiibacter**

**Halodesulfobivibrio**

**Halofium**

**Halomonas**

**Halopeptonella**

**Halorhodospira**

**Halospina**

**Halotalea**

**Halothiobacillus**

**Halovibrio**

**Hanschlegelia**

**Hartmannibacter**

**Haslibacter**

**Helicobacter**

**Hellea**

**Henriciella**

**Hephaestia**

**Herbaspirillum**

**Herminiimonas**

**Hipaea**

**Hirschia**

**Histidinibacterium**

**Histophilus**

**Hoeflea**

**Holospira**

**Humitalea**

**Hwanghaella**

**Hwanghaeicola**

**Hyalangium**

**Hydrocarboniclastic**

**Hydrocarboniphaga**

**Hydrogenimonas**

**Hydrogenophaga**

**Hydrogenophilus**

**Hydrogenovibrio**

**Hydromonas**

**Hylemonella**

**Hyphobacterium**

**Hyphomicrobium**

**Hyphomonas**

**Ideonella**

**Idiomarina**

**Ignatzschineria**

**Imhoffiella**

**Immundisolibacter**

**Indioceanicola**

**WDAO**

**ZCXC**

**ZCXC2**

**Keich**

**CAFTC**

**Suz12**

**SLED**

**SAM**

**VAF2\_RBP**

**PcKing**

**AF9**

**TRRAP**

**PTP**

**NCO6**

**Menp**

**Dry-30**

**PA1**

**AT7/P**

**Nucleoplasmin**

**UBN\_AB**

**ZI-CGH**

**Strc29**

**Strc29**

**RT106**

**NAP**

**HJURP\_mid**

**Hra**

**Daxx**

**CAF1A**

**ASF1\_hist\_chap**

**SNF2\_N**

**YEATS**

**ZI-CW**

**TUDOR**

**TTD**

**SAWADDEE**

**PHD**

**MRG**

**MBT**

**ING**

**Chromo**

**Biomodomain**

**BMH**

**ADD\_DMNT3**

**ADD\_ATTRX**

**CapnJnC**

**StrKlm**

**SET**

**DOT1**

**SIR2**

**Hist\_Deacetyl**

**MOZ\_SAS**

**KAT1\_N**

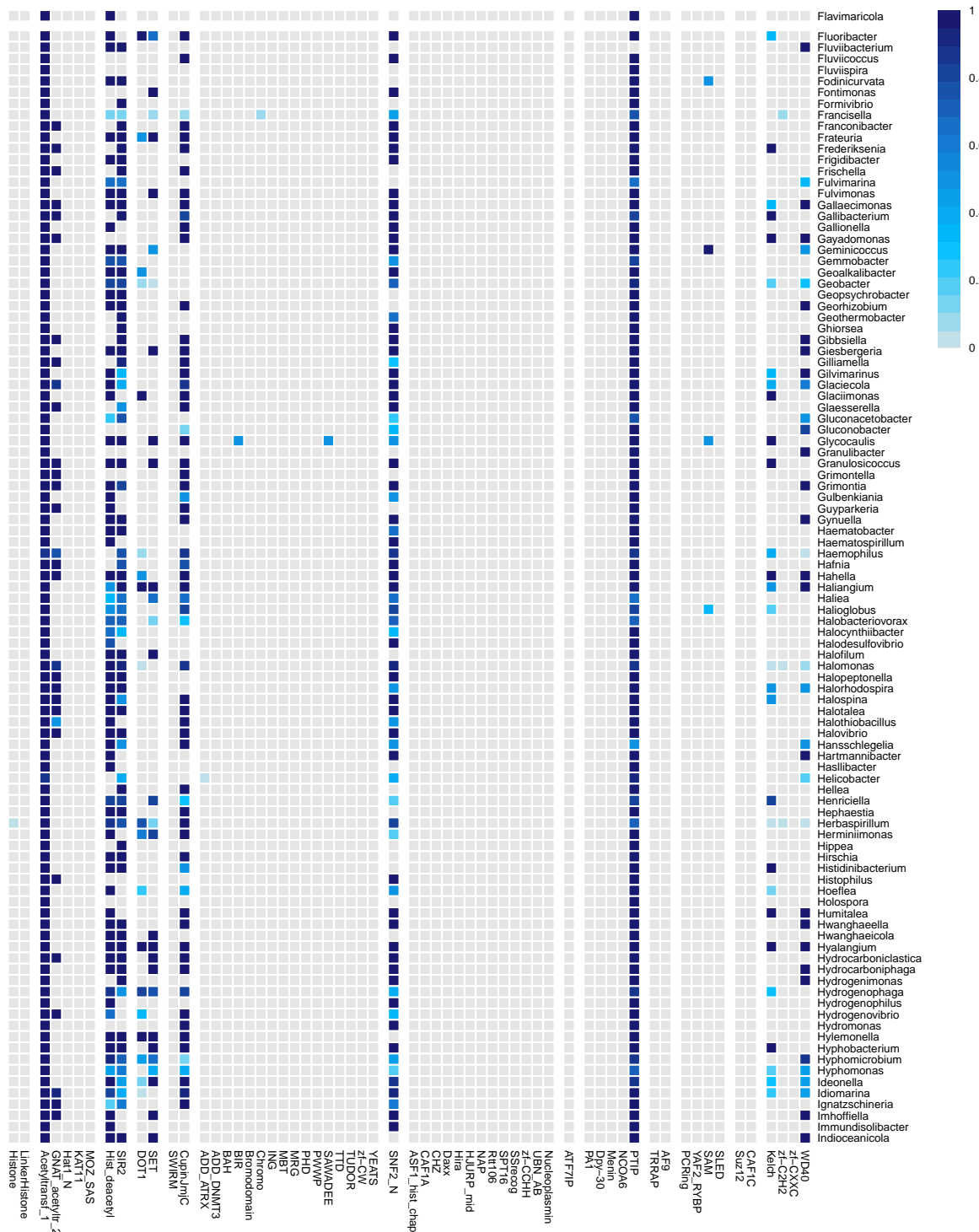
**KAT1\_N**

**GNAT\_acetylT\_2**

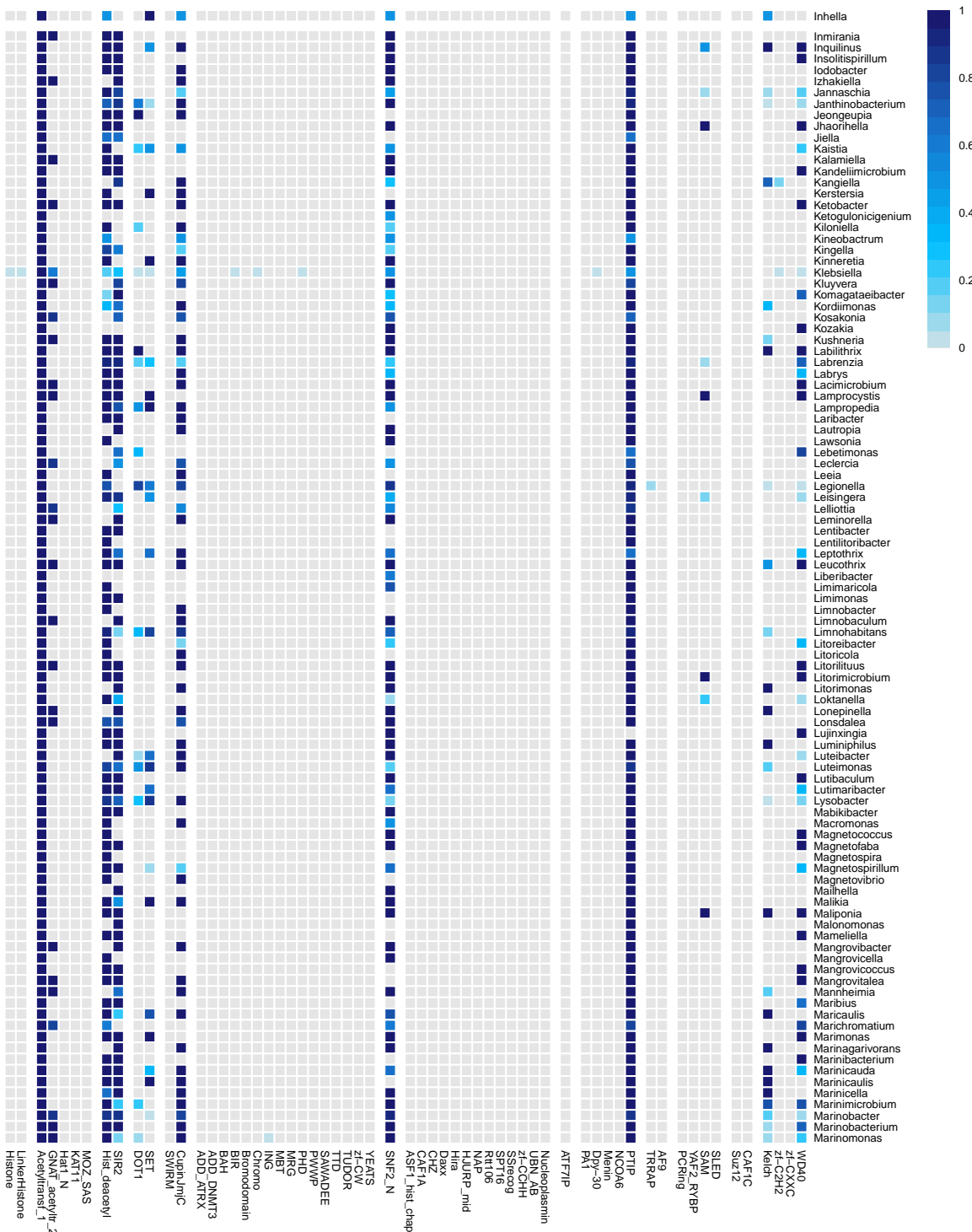
**Acetyltransf\_1**

**LinkerHistone**

**Histone**



Heatmap showing the relative abundance of 100 bacterial taxa across 100 samples. The taxa are listed on the y-axis, and the samples are listed on the x-axis. A color scale on the right indicates relative abundance from 0 (light blue) to 1 (dark blue). The taxa are grouped into several clusters, with some taxa showing high abundance across many samples and others showing more specific patterns.



Heatmap showing the relative abundance of 100 bacterial taxa across 100 samples. The taxa are listed on the right, and the samples are listed on the left. A color scale on the far right indicates relative abundance from 0 (light blue) to 1 (dark blue).

**Taxa (Right):**

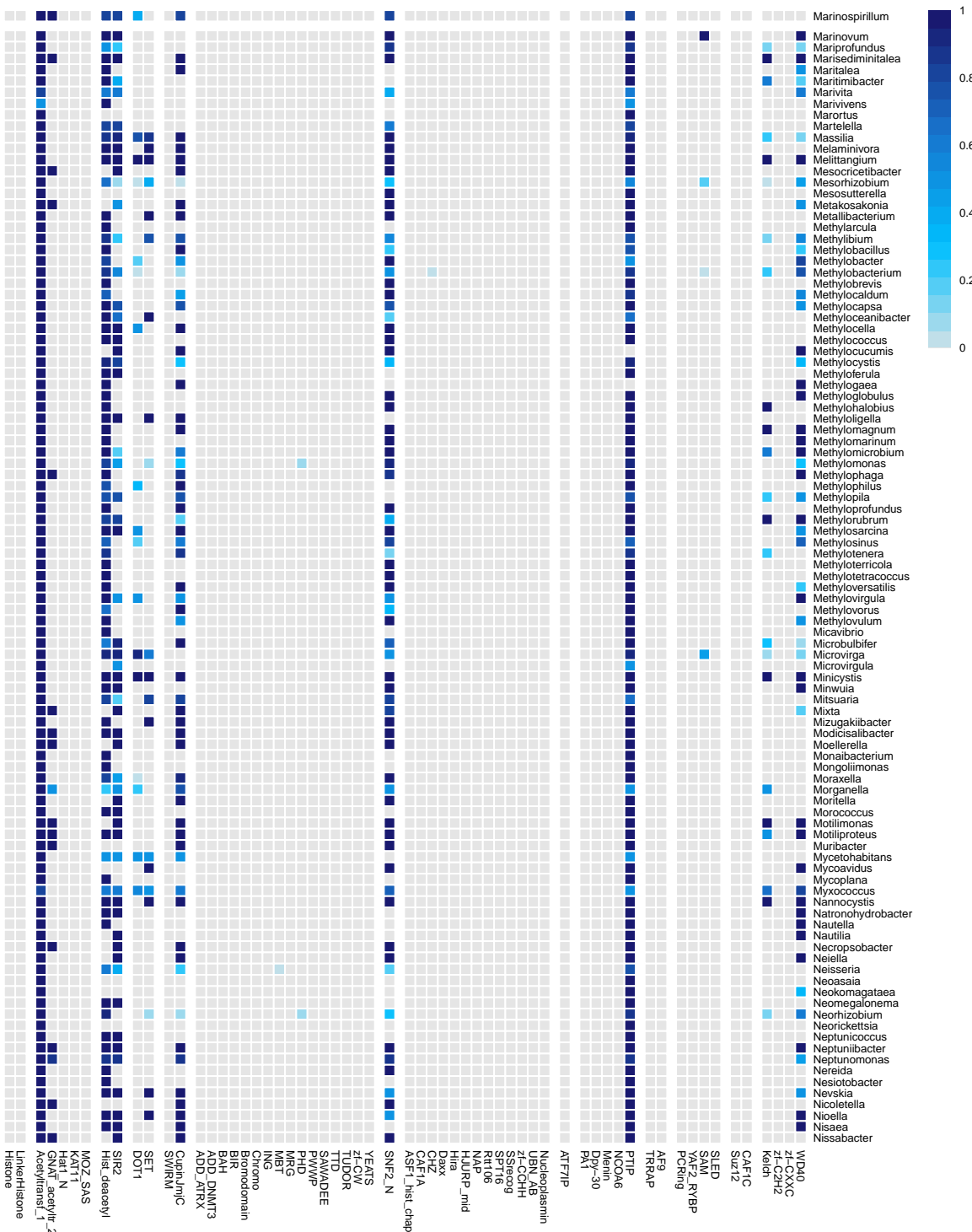
- Marinospirillum
- Marinovum
- Mariprofundus
- Marisedimentalea
- Maritalea
- Marimibacter
- Marivita
- Marivivens
- Marortus
- Martelella
- Massilia
- Melaminivora
- Melittangium
- Mesocricetibacter
- Mesorhizobium
- Mesosutterella
- Metakosakonia
- Metallibacterium
- Methyarcuia
- Methylibium
- Methylobacillus
- Methylobacter
- Methylobacterium
- Methylobrevus
- Methylocaldum
- Methylocapsa
- Methyloceanibacter
- Methylocella
- Methylococcus
- Methylocucumis
- Methylocystis
- Methyloferula
- Methylogaea
- Methyloglobulus
- Methylohalobius
- Methyloligella
- Methyloimagnium
- Methyloimarium
- Methyloimicrobium
- Methyloimonas
- Methylophaga
- Methylophilus
- Methyloplia
- Methyloprofundus
- Methylosuburum
- Methylosarcina
- Methylosinus
- Methylostenora
- Methylostericicola
- Methylostericococcus
- Methyloversatilis
- Methylovirgula
- Methylovorus
- Methylovulum
- Micavibrio
- Microbubifer
- Microvirga
- Microvirgula
- Minicystis
- Minwuia
- Mitsuaria
- Mixta
- Mizugakiibacter
- Modicisalibacter
- Moellerella
- Monaibacterium
- Mongoliomonas
- Moraxella
- Morganelia
- Morifella
- Morococcus
- Motilimonas
- Motiliproteus
- Muribacter
- Mycetohabitans
- Mycocavivirus
- Mycoplasma
- Mycococcus
- Nannocystis
- Natronohydrobacter
- Nautella
- Nautella
- Necropsobacter
- Neiella
- Neisseria
- Neosaxia
- Neokomagataea
- Neomagalania
- Neorhizobium
- Neorickettsia
- Neptunicoccus
- Neptuniibacter
- Neptunomonas
- Nereida
- Nesiotobacter
- Nevskia
- Nicoletella
- Nioella
- Nisaea
- Nissabacter

**Samples (Left):**

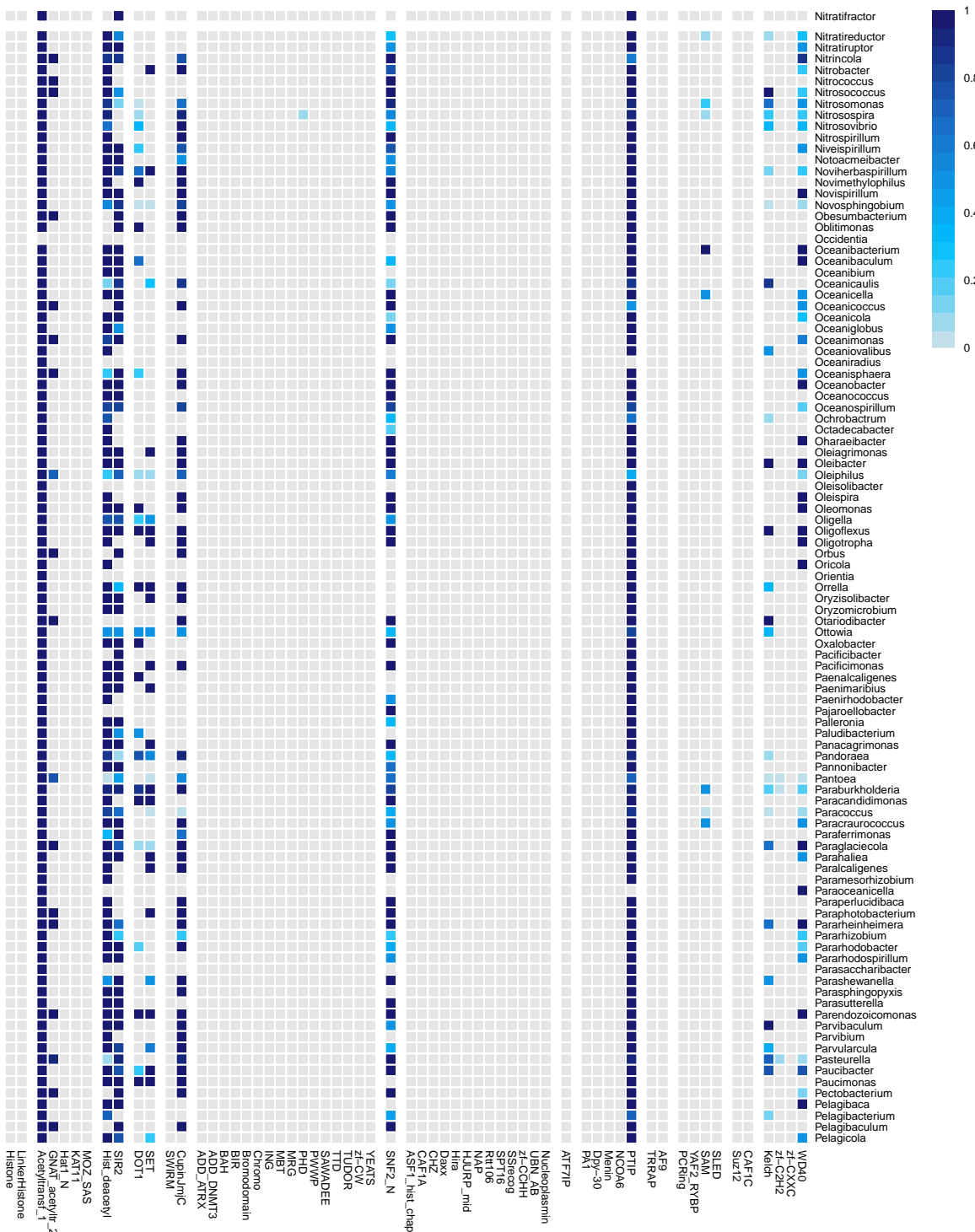
- YEATS
- z1-CW
- TUDOR
- ST10A
- ST10A-DEE
- PMWP
- PHD
- MRG
- MBT
- NG
- DN
- DN-Long
- Burkholderia
- BIR
- BAH
- ADD-DNMT3
- ADD-ATRX
- CapnJunc
- SWIRM
- SFT
- DOT1
- SI2
- Hist-deoxy
- MOZ-SAS
- KAT1
- Hatt\_N
- GAT1
- Acetyltrans\_1
- Acetyltrans\_2
- Acetyltrans\_3
- Acetyltrans\_4
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- Acetyltrans\_98
- Acetyltrans\_99
- Acetyltrans\_100

**Color Scale (Right):**

- 0 (light blue)
- 0.2
- 0.4
- 0.6
- 0.8
- 1 (dark blue)



Gene presence per genus in Proteobacteria (phylum) (8/12) (fraction of species)



Gene presence per genus in Proteobacteria (phylum) (9/12) (fraction of species)



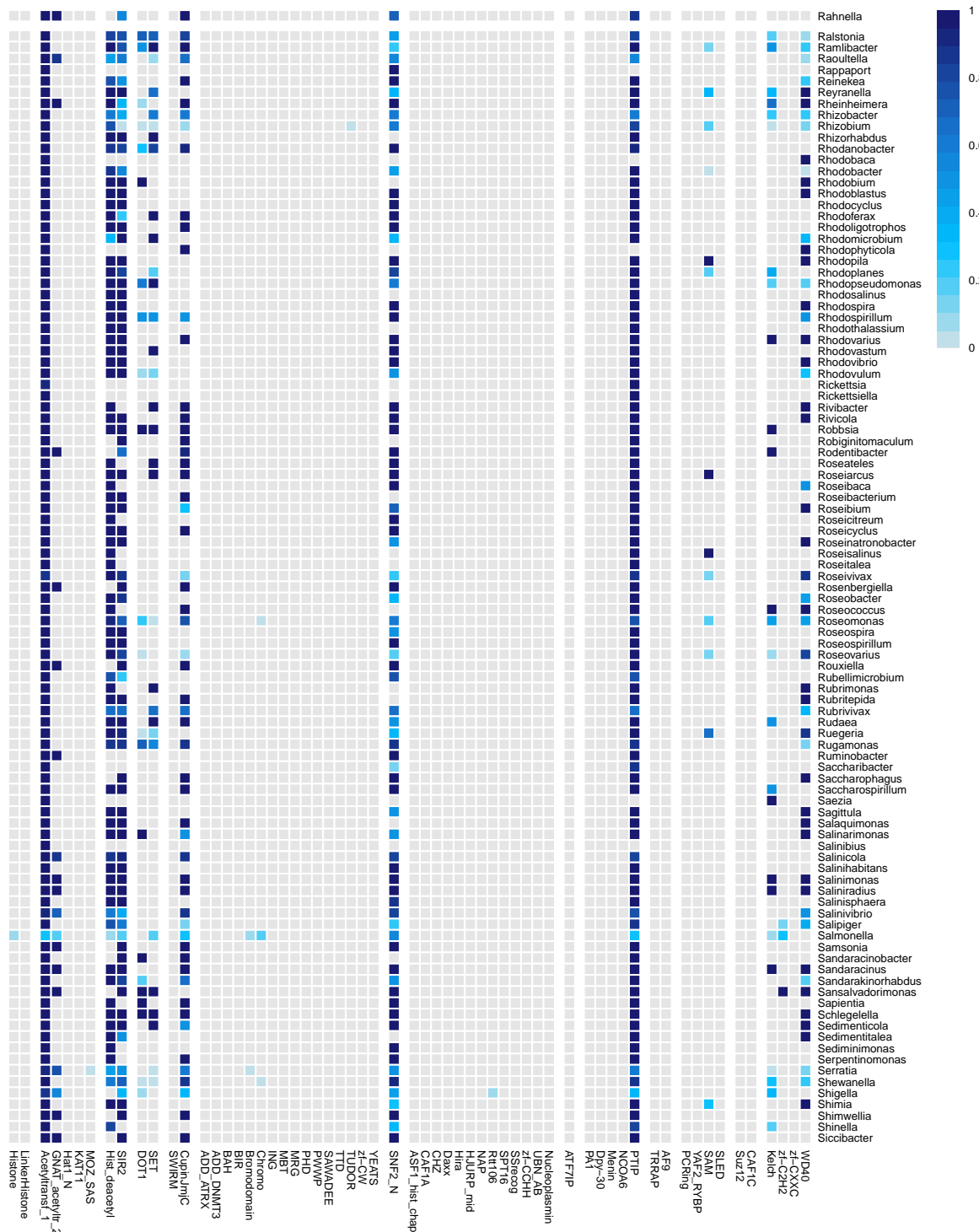
A heatmap visualization showing the relative abundance of various bacterial taxa across numerous samples. The color scale ranges from light blue (0) to dark purple/black (1). Taxa are listed on the right side of the plot, grouped by phylum or class. Samples are represented as columns along the bottom axis.

**Bacterial Taxa (Right Side Labels):**

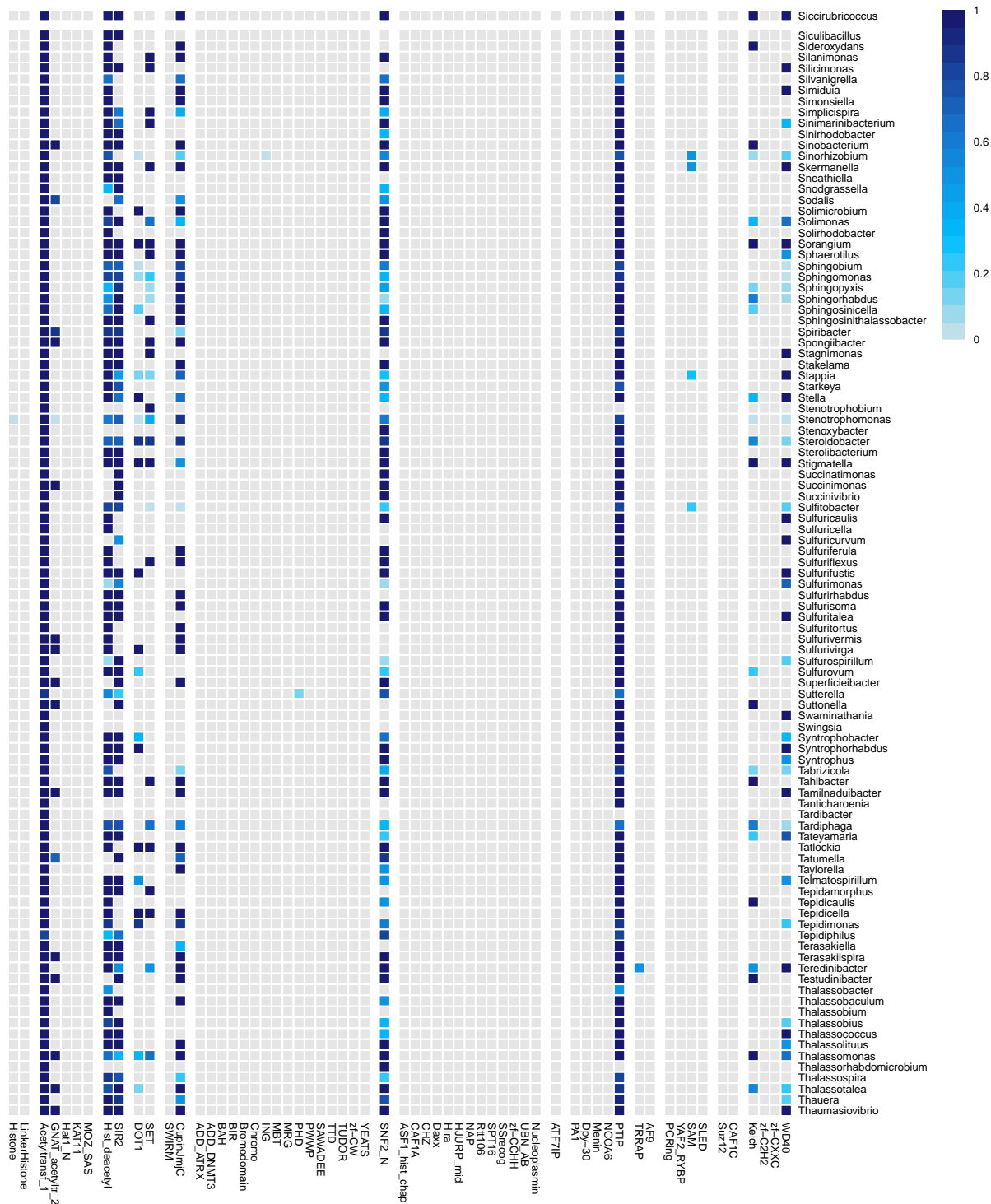
- Rahnella
- Ralstonia
- Ramlibacter
- Raoultella
- Rappaport
- Reinekea
- Reyranella
- Rheinheimera
- Rhizobacter
- Rhizobium
- Rhizorhabdus
- Rhodanobacter
- Rhodobaca
- Rhodobacter
- Rhodobium
- Rhodoblastus
- Rhodocyclus
- Rhodoferrax
- Rhodospirillum rubrum
- Rhododermatophilum
- Rhodopila
- Rhodoplanes
- Rhodopselomonas
- Rhodosalinus
- Rhodospira
- Rhodospirillum
- Rhodothalassium
- Rhodovarius
- Rhodovastum
- Rhodovibrio
- Rhodovulum
- Rickettsia
- Rickettsiella
- Rivibacter
- Rivicola
- Robbia
- Robertsonellaceae
- Rodentibacter
- Roseateles
- Roseiarcus
- Roseibacillus
- Roseibacterium
- Roseibium
- Roseicetone
- Roseicyclus
- Roseinatronobacter
- Roseisalinus
- Roseitalea
- Rosevivax
- Rosenbergiella
- Rosebacteria
- Roseococcus
- Roseomonas
- Roseospora
- Roseospirillum
- Roseoventer
- Rouxiiella
- Rubellimicrobium
- Rubrimonas
- Rubrilinea
- Rubrivivax
- Rudaea
- Ruegeria
- Rugamonas
- Ruminobacter
- Saccaribacter
- Saccharophagus
- Saccharosporillum
- Saezia
- Sagittula
- Salaquimonas
- Salinarmonas
- Salinihiabans
- Salininomads
- Salinariidius
- Salinisphaera
- Salinovibrio
- Salipicea
- Samonella
- Samsonia
- Sandaracinobacter
- Sandaracinus
- Sandarakinorhabdus
- Sansalvadorimonas
- Sapiientia
- Schlegellella
- Sedimenticola
- Sedimentitalea
- Sediminimonas
- Serpentinomonas
- Serratia
- Shewanella
- Shigella
- Shimia
- Shimwellia
- Shinnella
- Siccibacter

**Samples (Bottom Axis Labels - partially visible):**

- Nucleoplasm
- UBN\_AB
- ZI-CCHH
- SSTC9
- RTT06
- HJURP\_mid
- Dmax
- CATFIA
- ASF1\_hst\_chap
- SNR2\_N
- YEATS
- ZI-CW
- TUDOR
- TTD
- SAWADEE
- PMDIP
- MGT
- MKG
- PHD
- ING
- Chromo
- Biomonomain
- BAH
- AAD\_DMUT3
- ADD\_ATTRX
- Guanjunc
- SVKRM
- SET
- DOT1
- Hist\_deacyl
- MOZ\_SAS
- KATI\_7
- KATI\_N
- Acetyltransf\_1
- Linkertstone
- Histone



Gene presence per genus in Proteobacteria (phylum) (11/12) (fraction of species)



Gene presence per genus in Proteobacteria (phylum) (12/12) (fraction of species)





num species per group

