

## **Additional file A-13 - CGP performance on peptidoglycan-related genes on *Prochlorococcus marinus* MIT9313**

### **Movitation and Methods**

In Case study 1, both statistical and inductive CGP methods were evaluated on genes of *Streptococcus agalactiae* 2603 V/R (SA-2603) and *Escherichia coli* K-12 (EC-K12) genomes with a large collection of genome examples incorporating the phyla of Firmicutes (99) and Proteobacteria (222). It is unclear whether the CGP results were favorably biased by the overrepresentation of these genome examples. To investigate the possible bias, an additional experiment was designed to rank genes from the *Prochlorococcus marinus* MIT9313 genomes (PM-MIT9313, 2,269 genes) with all Cyanobacteria genomes removed from the 400 positive genomes examples (including all *Anabaena* spp., *Gloeobacter* spp., *Nostoc* spp., *Prochlorococcus* spp., *Synechococcus* spp., *Synechocystis* spp., *Thermosynechococcus* spp., and *Trichodesmium* spp.). The determination of occurrence matrix, statistical CGP scoring functions, inductive CGP algorithms, and evaluation methodology were identical to Case study 1.

### **Results**

For statistical CGP, the best performing scoring function was *hmss* with high AUC (best AUC: *C*: 0.988, *B*: 0.971, *M*: 0.969). Good inductive CGP performance as measured by AUC were also obtained (*C*:0.999 by *ADTree*; *B*: 0.954 by *SVM/Poly*; *M*: 0.986 by *SVM/Poly*).

### **Discussion**

This experiment attempted to prioritise PM-MIT9313 for discovering genes responsible for peptidoglycan metabolism *without* the representative genomes from the phylum Cyanobacteria. The high AUCs achieved by both methods were similar to the corresponding CGP tasks in SA-2603 and EC-K12 genomes. These results demonstrated that good CGP performances were obtainable by selecting positive genome examples from either closely- or distantly-related genomes.

Table 1: CGP performance on peptidoglycan-related genes (*Prochlorococcus marinus* MIT9313, 2269 genes)

Methods	Validation sets					
	C (8)		B (23)		M (30)	
	AUC	$(\bar{\eta}/\eta_{max})$	AUC	$(\bar{\eta}/\eta_{max})$	AUC	$(\bar{\eta}/\eta_{max})$
Statistical CGP (scoring functions)						
<i>sens</i>	0.887	(2.2/6.65)	0.867	(2.0/4.93)	0.858	(2.0/4.44)
<i>spec</i>	0.262	(0.3/1.33)	0.331	(0.5/1.18)	0.381	(0.7/1.18)
<i>ppv</i>	0.527	(0.8/1.96)	0.594	(1.4/24.7)	0.641	(1.5/37.8)
<i>npv</i>	0.970	(3.6/25.2)	0.953	(3.2/16.4)	0.946	(3.1/15.1)
<i>amss</i>	0.987	(4.6/75.6)	0.970	(4.0/41.5)	0.969	(4.0/43.8)
<i>hmss</i>	0.988	(4.8/87.3)	0.971	(4.0/45.5)	0.969	(4.0/47.3)
<i>OR</i>	0.521	(0.7/1.96)	0.593	(1.4/24.7)	0.638	(1.4/37.8)
<i>chisq</i>	0.980	(4.1/42.2)	0.963	(3.6/23.1)	0.957	(3.5/20.9)
<i>bchisq</i>	0.980	(4.1/42.2)	0.963	(3.6/23.1)	0.957	(3.5/20.9)
<i>F</i>	0.943	(3.3/29.9)	0.906	(2.8/16.4)	0.894	(2.7/15.1)
Inductive CGP (machine learning algorithms)						
<i>NB</i>	0.919		0.878		0.872	
<i>LR</i>	0.968		0.819		0.877	
<i>ADTree</i>	0.999		0.921		0.915	
<i>IBk</i>	0.833		0.952		0.960	
<i>J48</i>	0.993		0.818		0.705	
<i>SMO/Poly</i>	0.937		0.954		0.986	
<i>SMO/RBF</i>	0.995		0.898		0.855	

## Genome examples selected for the statistical CGP experiment

### Positive genome examples (378)

Acidobacteria bacterium Ellin345	Bordetella parapertussis
Acidothermus cellulolyticus 11B	Bordetella pertussis
Acidovorax JS42	Borrelia afzelii PKo
Acidovorax avenae citrulli AAC00-1	Borrelia burgdorferi
Acinetobacter sp ADP1	Borrelia garinii PBi
Aeromonas hydrophila ATCC 7966	Bradyrhizobium japonicum
Agrobacterium tumefaciens C58 UWash	Brucella abortus 9-941
Alcanivorax borkumensis SK2	Brucella melitensis
Alkalilimnicola ehrlichei MLHE-1	Brucella melitensis biovar Abortus
Anabaena variabilis ATCC 29413	Brucella suis 1330
Anaeromyxobacter dehalogenans 2CP-C	Buchnera aphidicola
Aquifex aeolicus	Buchnera aphidicola Cc Cinara cedri
Arthrobacter FB24	Buchnera aphidicola Sg
Arthrobacter aurescens TC1	Buchnera sp
Azoarcus BH72	Burkholderia 383
Azoarcus sp EbN1	Burkholderia cenocepacia AU 1054
Bacillus anthracis Ames	Burkholderia cenocepacia HI2424
Bacillus anthracis Ames 0581	Burkholderia cepacia AMMD
Bacillus anthracis str Sterne	Burkholderia mallei ATCC 23344
Bacillus cereus ATCC14579	Burkholderia mallei NCTC 10229
Bacillus cereus ATCC 10987	Burkholderia mallei SAVP1
Bacillus cereus ZK	Burkholderia pseudomallei 1710b
Bacillus clausii KSM-K16	Burkholderia pseudomallei K96243
Bacillus halodurans	Burkholderia thailandensis E264
Bacillus licheniformis DSM 13	Burkholderia xenovorans LB400
Bacillus subtilis	Campylobacter fetus 82-40
Bacillus thuringiensis Al Hakam	Campylobacter jejuni
Bacillus thuringiensis konkukian	Campylobacter jejuni RM1221
Bacteroides fragilis NCTC 9434	Candidatus Blochmannia floridanus
Bacteroides fragilis YCH46	Candidatus Blochmannia pennsylvanicus
Bacteroides thetaiotaomicron VPI-5482	BPEN
Bartonella bacilliformis KC583	Candidatus Carsonella ruddii
Bartonella henselae Houston-1	Candidatus Pelagibacter ubique HTCC1062
Bartonella quintana Toulouse	Candidatus Ruthia magnifica Cm Calyptogena magnifica
Baumannia cicadellinicola Homalodisca coagulata	Carboxydothermus hydrogenoformans Z-2901
Bdellovibrio bacteriovorus	Caulobacter crescentus
Bifidobacterium adolescentis ATCC 15703	Chlamydia muridarum
Bifidobacterium longum	Chlamydia trachomatis
Bordetella bronchiseptica	Chlamydia trachomatis A HAR-13
	Chlamydophila abortus S26 3

<i>Chlamydophila caviae</i>	<i>Escherichia coli</i> K12
<i>Chlamydophila felis</i> Fe C-56	<i>Escherichia coli</i> O157H7
<i>Chlamydophila pneumoniae</i> AR39	<i>Escherichia coli</i> O157H7 EDL933
<i>Chlamydophila pneumoniae</i> CWL029	<i>Escherichia coli</i> UTI89
<i>Chlamydophila pneumoniae</i> J138	<i>Escherichia coli</i> W3110
<i>Chlamydophila pneumoniae</i> TW 183	<i>Francisella tularensis</i> FSC 198
<i>Chlorobium chlorochromatii</i> CaD3	<i>Francisella tularensis</i> holarktica
<i>Chlorobium phaeobacteroides</i> DSM 266	<i>Francisella tularensis</i> holarktica OSU18
<i>Chlorobium tepidum</i> TLS	<i>Francisella tularensis</i> novicida U112
<i>Chromobacterium violaceum</i>	<i>Francisella tularensis</i> tularensis
<i>Chromohalobacter salexigens</i> DSM 3043	<i>Frankia</i> CcI3
<i>Clostridium acetobutylicum</i>	<i>Frankia alni</i> ACN14a
<i>Clostridium novyi</i> NT	<i>Fusobacterium nucleatum</i>
<i>Clostridium perfringens</i>	<i>Geobacillus kaustophilus</i> HTA426
<i>Clostridium perfringens</i> ATCC 13124	<i>Geobacter metallireducens</i> GS-15
<i>Clostridium perfringens</i> SM101	<i>Geobacter sulfurreducens</i>
<i>Clostridium tetani</i> E88	<i>Gluconobacter oxydans</i> 621H
<i>Clostridium thermocellum</i> ATCC 27405	<i>Gramella forsetii</i> KT0803
<i>Colwellia psychrerythraea</i> 34H	<i>Granulobacter bethesdensis</i> CGDNIH1
<i>Corynebacterium diphtheriae</i>	<i>Haemophilus ducreyi</i> 35000HP
<i>Corynebacterium efficiens</i> YS-314	<i>Haemophilus influenzae</i>
<i>Corynebacterium glutamicum</i> ATCC 13032	<i>Haemophilus influenzae</i> 86 028NP
<i>Bielefeld</i>	<i>Haemophilus somnus</i> 129PT
<i>Corynebacterium jeikeium</i> K411	<i>Hahella chejuensis</i> KCTC 2396
<i>Coxiella burnetii</i>	<i>Halorhodospira halophila</i> SL1
<i>Cytophaga hutchinsonii</i> ATCC 33406	<i>Helicobacter acinonychis</i> Sheeba
<i>Dechloromonas aromatica</i> RCB	<i>Helicobacter hepaticus</i>
<i>Dehalococcoides CBDB1</i>	<i>Helicobacter pylori</i> 26695
<i>Dehalococcoides ethenogenes</i> 195	<i>Helicobacter pylori</i> HPAG1
<i>Deinococcus geothermalis</i> DSM 11300	<i>Helicobacter pylori</i> J99
<i>Deinococcus radiodurans</i>	<i>Hyphomonas neptunium</i> ATCC 15444
<i>Desulfitobacterium hafniense</i> Y51	<i>Idiomarina loihiensis</i> L2TR
<i>Desulfotalea psychrophila</i> LSv54	<i>Jannaschia CCS1</i>
<i>Desulfovibrio desulfuricans</i> G20	<i>Lactobacillus acidophilus</i> NCFM
<i>Desulfovibrio vulgaris</i> DP4	<i>Lactobacillus brevis</i> ATCC 367
<i>Desulfovibrio vulgaris</i> Hildenborough	<i>Lactobacillus casei</i> ATCC 334
<i>Ehrlichia canis</i> Jake	<i>Lactobacillus delbrueckii</i> bulgaricus
<i>Ehrlichia chaffeensis</i> Arkansas	<i>Lactobacillus delbrueckii</i> bulgaricus ATCC
<i>Ehrlichia ruminantium</i> Gardel	BAA-365
<i>Ehrlichia ruminantium</i> str. Welgevonden	<i>Lactobacillus gasseri</i> ATCC 33323
<i>Enterococcus faecalis</i> V583	<i>Lactobacillus johnsonii</i> NCC 533
<i>Erwinia carotovora</i> atroseptica SCRI1043	<i>Lactobacillus plantarum</i>
<i>Erythrobacter litoralis</i> HTCC2594	<i>Lactobacillus sakei</i> 23K
<i>Escherichia coli</i> 536	<i>Lactobacillus salivarius</i> UCC118
<i>Escherichia coli</i> APEC O1	<i>Lactococcus lactis</i>
<i>Escherichia coli</i> CFT073	<i>Lactococcus lactis</i> cremoris MG1363

<i>Lactococcus lactis</i> cremoris SK11	<i>Nitrobacter hamburgensis</i> X14
<i>Lawsonia intracellularis</i> PHE MN1-00	<i>Nitrobacter winogradskyi</i> Nb-255
<i>Legionella pneumophila</i> Lens	<i>Nitrosoccus oceanii</i> ATCC 19707
<i>Legionella pneumophila</i> Paris	<i>Nitrosomonas europaea</i>
<i>Legionella pneumophila</i> Philadelphia 1	<i>Nitrosomonas eutropha</i> C71
<i>Leifsonia xyli</i> xyli CTCB0	<i>Nitrosospira multiformis</i> ATCC 25196
<i>Leptospira borgpetersenii</i> serovar Hardjo-bovis JB197	<i>Nocardia farcinica</i> IFM10152
<i>Leptospira borgpetersenii</i> serovar Hardjo-bovis L550	<i>Nocardoides JS614</i>
<i>Leptospira interrogans</i> serovar Copenhageni	<i>Novosphingobium aromaticivorans</i> DSM 12444
<i>Leptospira interrogans</i> serovar Lai	<i>Oceanobacillus iheyensis</i>
<i>Leuconostoc mesenteroides</i> ATCC 8293	<i>Oenococcus oeni</i> PSU-1
<i>Listeria innocua</i>	<i>Parachlamydia</i> sp UWE25
<i>Listeria monocytogenes</i>	<i>Paracoccus denitrificans</i> PD1222
<i>Listeria monocytogenes</i> 4b F2365	<i>Pasteurella multocida</i>
<i>Listeria welshimeri</i> serovar 6b SLCC5334	<i>Pediococcus pentosaceus</i> ATCC 25745
<i>Magnetococcus</i> MC-1	<i>Pelobacter carbinolicus</i>
<i>Magnetospirillum magneticum</i> AMB-1	<i>Pelobacter propionicus</i> DSM 2379
<i>Mannheimia succiniciproducens</i> MBEL55E	<i>Pelodictyon luteolum</i> DSM 273
<i>Maricaulis maris</i> MCS10	<i>Photobacterium profundum</i> SS9
<i>Marinobacter aquaeolei</i> VT8	<i>Photorhabdus luminescens</i>
<i>Mesorhizobium</i> BNC1	<i>Pirellula</i> sp
<i>Mesorhizobium loti</i>	<i>Polaromonas JS666</i>
<i>Methylibium petroleiphilum</i> PM1	<i>Polaromonas naphthalenivorans</i> CJ2
<i>Methylobacillus flagellatus</i> KT	<i>Porphyromonas gingivalis</i> W83
<i>Methylococcus capsulatus</i> Bath	<i>Propionibacterium acnes</i> KPA171202
<i>Moorella thermoacetica</i> ATCC 39073	<i>Pseudoalteromonas atlantica</i> T6c
<i>Mycobacterium KMS</i>	<i>Pseudoalteromonas haloplanktis</i> TAC125
<i>Mycobacterium MCS</i>	<i>Pseudomonas aeruginosa</i>
<i>Mycobacterium avium</i> 104	<i>Pseudomonas aeruginosa</i> UCBPP-PA14
<i>Mycobacterium avium</i> paratuberculosis	<i>Pseudomonas entomophila</i> L48
<i>Mycobacterium bovis</i>	<i>Pseudomonas fluorescens</i> Pf-5
<i>Mycobacterium bovis</i> BCG Pasteur 1173P2	<i>Pseudomonas fluorescens</i> PfO-1
<i>Mycobacterium leprae</i>	<i>Pseudomonas putida</i> KT2440
<i>Mycobacterium smegmatis</i> MC2 155	<i>Pseudomonas syringae</i> phaseolicola 1448A
<i>Mycobacterium tuberculosis</i> CDC1551	<i>Pseudomonas syringae</i> pv B728a
<i>Mycobacterium tuberculosis</i> H37Rv	<i>Pseudomonas syringae</i> tomato DC3000
<i>Mycobacterium ulcerans</i> Agy99	<i>Psychrobacter arcticum</i> 273-4
<i>Mycobacterium vanbaalenii</i> PYR-1	<i>Psychrobacter cryohalolentis</i> K5
<i>Myxococcus xanthus</i> DK 1622	<i>Psychromonas ingrahamii</i> 37
<i>Neisseria gonorrhoeae</i> FA 1090	<i>Ralstonia eutropha</i> H16
<i>Neisseria meningitidis</i> FAM18	<i>Ralstonia eutropha</i> JMP134
<i>Neisseria meningitidis</i> MC58	<i>Ralstonia metallidurans</i> CH34
<i>Neisseria meningitidis</i> Z2491	<i>Ralstonia solanacearum</i>
<i>Neorickettsia sennetsu</i> Miyayama	<i>Rhizobium etli</i> CFN 42
	<i>Rhizobium leguminosarum</i> bv <i>viciae</i> 3841

Rhodobacter sphaeroides 2 4 1	
Rhodococcus RHA1	Staphylococcus aureus N315
Rhodoferax ferrireducens T118	Staphylococcus aureus NCTC 8325
Rhodopseudomonas palustris BisA53	Staphylococcus aureus RF122
Rhodopseudomonas palustris BisB18	Staphylococcus aureus USA300
Rhodopseudomonas palustris BisB5	Staphylococcus aureus aureus MRSA252
Rhodopseudomonas palustris CGA009	Staphylococcus aureus aureus MSSA476
Rhodopseudomonas palustris HaA2	Staphylococcus epidermidis ATCC 12228
Rhodospirillum rubrum ATCC 11170	Staphylococcus epidermidis RP62A
Rickettsia bellii RML369-C	Staphylococcus haemolyticus
Rickettsia conorii	Staphylococcus saprophyticus
Rickettsia felis URRWXCal2	Streptococcus agalactiae 2603
Rickettsia prowazekii	Streptococcus agalactiae A909
Rickettsia typhi wilmington	Streptococcus agalactiae NEM316
Roseobacter denitrificans OCh 114	Streptococcus mutans
Rubrobacter xylanophilus DSM 9941	Streptococcus pneumoniae D39
Saccharophagus degradans 2-40	Streptococcus pneumoniae R6
Salinibacter ruber DSM 13855	Streptococcus pyogenes M1 GAS
Salmonella enterica Choleraesuis	Streptococcus pyogenes MGAS10270
Salmonella enterica Paratyphi ATCC 9150	Streptococcus pyogenes MGAS10394
Salmonella typhi	Streptococcus pyogenes MGAS10750
Salmonella typhi Ty2	Streptococcus pyogenes MGAS2096
Salmonella typhimurium LT2	Streptococcus pyogenes MGAS315
Shewanella ANA-3	Streptococcus pyogenes MGAS5005
Shewanella MR-4	Streptococcus pyogenes MGAS6180
Shewanella MR-7	Streptococcus pyogenes MGAS8232
Shewanella W3-18-1	Streptococcus pyogenes MGAS9429
Shewanella amazonensis SB2B	Streptococcus pyogenes SSI-1
Shewanella denitrificans OS217	Streptococcus sanguinis SK36
Shewanella frigidimarina NCIMB 400	Streptococcus thermophilus CNRZ1066
Shewanella oneidensis	Streptococcus thermophilus LMD-9
Shigella boydii Sb227	Streptococcus thermophilus LMG 18311
Shigella dysenteriae	Streptomyces avermitilis
Shigella flexneri 2a	Streptomyces coelicolor
Shigella flexneri 2a 2457T	Symbiobacterium thermophilum IAM14863
Shigella flexneri 5 8401	Syntrophobacter fumaroxidans MPOB
Shigella sonnei Ss046	Syntrophomonas wolfei Goettingen
Silicibacter TM1040	Syntrophus aciditrophicus SB
Silicibacter pomeroyi DSS-3	Thermoanaerobacter tengcongensis
Sinorhizobium meliloti	Thermobifida fusca YX
Sodalis glossinidius morsitans	Thermotoga maritima
Solibacter usitatus Ellin6076	Thermus thermophilus HB27
Sphingopyxis alaskensis RB2256	Thermus thermophilus HB8
Staphylococcus aureus COL	Thiobacillus denitrificans ATCC 25259
Staphylococcus aureus MW2	Thiomicrospira crunogena XCL-2
Staphylococcus aureus Mu50	Thiomicrospira denitrificans ATCC 33889
	Treponema denticola ATCC 35405

Treponema pallidum	Xanthomonas campestris 8004
Tropheryma whipplei TW08 27	Xanthomonas campestris vesicatoria 85-10
Tropheryma whipplei Twist	Xanthomonas citri
Verminephrobacter eiseniae EF01-2	Xanthomonas oryzae KACC10331
Vibrio cholerae	Xanthomonas oryzae MAFF 311018
Vibrio fischeri ES114	Xylella fastidiosa
Vibrio parahaemolyticus	Xylella fastidiosa Temecula1
Vibrio vulnificus CMCP6	Yersinia enterocolitica 8081
Vibrio vulnificus YJ016	Yersinia pestis Antiqua
Wigglesworthia brevipalpis	Yersinia pestis CO92
Wolbachia endosymbiont of Brugia malayi	Yersinia pestis KIM
TRS	Yersinia pestis Nepal516
Wolbachia endosymbiont of Drosophila melanogaster	Yersinia pestis biovar Mediaevails
Wolinella succinogenes	Yersinia pseudotuberculosis IP32953
Xanthomonas campestris	Zymomonas mobilis ZM4

### Negative genome examples (17)

Anaplasma marginale St Maries	Mycoplasma hyopneumoniae J
Anaplasma phagocytophilum HZ	Mycoplasma mobile 163K
Aster yellows witches-broom phytoplasma AYW	Mycoplasma mycooides
Mesoplasma florum L1	Mycoplasma penetrans
Mycoplasma capricolum ATCC 27343	Mycoplasma pneumoniae
Mycoplasma gallisepticum	Mycoplasma pulmonis
Mycoplasma hyopneumoniae 232	Mycoplasma synoviae 53
Mycoplasma hyopneumoniae 7448	Onion yellows phytoplasma
	Ureaplasma urealyticum