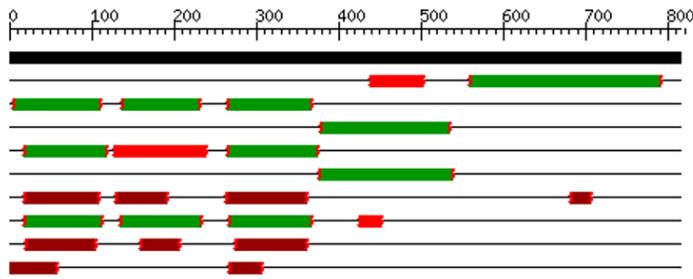


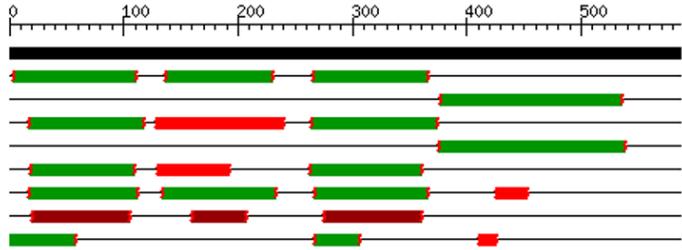
Figure S1. Standardized biofilm formation in ASW medium at 23 h in CUKW and KCC01.

CUKW dgc\_4914 (BFV93\_4914)



- BFV93\_4914
- PF00563.15: EAL domain
- PF13426.1: PAS domain
- PF00990.16: GGDEF domain
- TIGR00229: PAS domain S-box protein
- TIGR00254: diguanylate cyclase (GGDEF) domain
- PF00989.19: PAS fold
- PF08448.5: PAS fold
- PF08447.6: PAS fold
- PF13188.1: PAS domain

KCC01 dgc\_4791 (BFV94\_4790)



- BFV94\_4790
- PF13426.1: PAS domain
- PF00990.16: GGDEF domain
- TIGR00229: PAS domain S-box protein
- TIGR00254: diguanylate cyclase (GGDEF) domain
- PF00989.19: PAS fold
- PF08448.5: PAS fold
- PF08447.6: PAS fold
- PF13188.1: PAS domain

Figure S2. Comparative domain analysis of diguanylate cyclase gene (dgc\_4914) found to be lacking EAL domain in KCC01.

Table S1. BestKeeper analysis for +Cu (n=10).

Target	gyrB	inf2	pfk	maf
geo Mean [CT]	23.26	20.92	26.97	23.97
ar Mean [CT]	23.28	20.93	26.98	23.98
min [CT]	22.18	19.92	26.44	23.22
max [CT]	24.97	22.30	28.08	25.44
std dev [ $\pm$ CT]	0.72	0.60	0.40	0.50
CV [% CT]	3.10	2.89	1.48	2.10
min [x-fold]	-2.12	-2.01	-1.45	-1.69
max [x-fold]	3.27	2.60	2.16	2.77
std dev [ $\pm$ x-fold]	1.65	1.52	1.32	1.42
Pearson coefficient of correlation [r]	0.95	0.96	0.96	0.96
p-value	0.00	0.00	0.00	0.00

Table S2. BestKeeper analysis for low-nutrient (n=11).

Target	gyrB	inf2	pfk	maf
geo Mean [CT]	18.67	16.57	21.35	20.20
ar Mean [CT]	18.70	16.61	21.38	20.20
min [CT]	17.14	15.19	19.55	19.63
max [CT]	20.56	18.35	22.87	20.92
std dev [ $\pm$ CT]	1.04	1.11	1.02	0.27
CV [% CT]	5.58	6.68	4.79	1.34
min [x-fold]	-2.88	-2.61	-3.47	-1.48
max [x-fold]	3.70	3.42	2.88	1.66
std dev [ $\pm$ x-fold]	2.06	2.16	2.03	1.21
Pearson coefficient of correlation [r]	0.97	0.98	0.97	0.66
p-value	0.00	0.00	0.00	0.03

Table S3. Summary of domain comparisons and gene content among CUKW and KCC01 in putative diguanylate cyclase genes.

<b>CUKW GB locus ID</b>	<b>GGDEF</b>	<b>EAL</b>	<b>Other domains</b>	<b>KCC01 GB locus ID</b>	<b>GGDEF</b>	<b>EAL</b>	<b>Other domains</b>
BFV93_0069	Y	Y		BFV94_0069	Y	Y	
BFV93_0081	Y	Y	GAF	BFV94_0081	Y	Y	GAF
BFV93_0114	Y	N		BFV94_0114	Y	N	
BFV93_0133	Y	N	7 TM diverse intracellular signaling	BFV94_0133	Y	N	7 TM diverse intracellular signaling
BFV93_0151	Y	Y	PAS S-box protein, PAS fold, 7TM-DISM extracellular 2	BFV94_0152	Y	Y	PAS S-box protein, PAS fold, 7TM-DISM extracellular 2
BFV93_0158	Y	N	response regulator receiver	BFV94_0159	Y	N	response regulator receiver
BFV93_0161	Y	N	MASE1	BFV94_0162	Y	N	MASE1
FV93_0305	Y	N		BFV94_0306	Y	N	
BFV93_0379	Y	Y	PAS S-box protein, PAS, PAS fold	BFV94_0380	Y	Y	PAS S-box protein, PAS, PAS fold
BFV93_0458	Y	N	7TM intracellular signaling, 7TMR-DISM extracellular 2	BFV94_0459	Y	N	7TM intracellular signaling, 7TMR-DISM extracellular 2
BFV93_0583	Y	N	PAS fold	BFV94_0585	y	N	PAS fold
BFV93_0640	Y	N		BFV94_0642	Y	N	
BFV93_0952	Y	N		BFV94_0952	Y	N	
BFV93_0985	Y	N		BFV94_0985	Y	N	
BFV93_1181	Y	N	Two-component regulator propeller, Y_Y_Y	BFV94_1181	Y	N	Two-component regulator propeller, Y_Y_Y
BFV93_1256	Y	N		BFV94_1257	Y	N	
BFV93_1270	Y	Y	PAS S-box protein, PAS, PAS fold, Two-component regulator propeller, Y_Y_Y	BFV94_1271	Y	Y	PAS S-box protein, PAS, PAS fold
BFV93_1299	Y	N		BFV94_1301	Y	N	
BFV93_1306	Y	Y	HAMP	BFV94_1308	Y	Y	HAMP
BFV93_1340	Y	N	PAS S-box protein	BFV94_1342	Y	N	PAS S-box protein
BFV93_1352	Y	N		BFV94_1354	Y	N	
BFV93_1526	Y	Y		BFV94_1531	Y	Y	
BFV93_1544	Y	Y	PAS S-box protein	BFV94_1548	Y	Y	PAS S-box protein
BFV93_1545	Y	Y		BFV94_1549	Y	Y	
BFV93_1557	Y	N	PAS S-box protein, PAS fold, PAS	BFV94_1561	Y	N	PAS S-box protein, PAS fold, PAS
BFV93_1571	Y	Y	Response regulator receiver, PAS S-box protein, PAS fold, PAS	BFV94_1575	Y	Y	Response regulator receiver, PAS S-box protein, PAS fold, PAS
BFV93_1572	Y	N	Response regulator receiver	BFV94_1576	Y	N	Response regulator receiver
BFV93_1606	Y	Y		BFV94_1610	Y	Y	
BFV93_1663	Y	N	extracellular solute-binding family 3 protein	BFV94_1667	Y	N	extracellular solute-binding family 3 protein
BFV93_1672	Y	N		BFV94_1676	Y	N	
BFV93_1731	Y	N	Two-component regulator propeller, Y_Y_Y	BFV94_1734	Y	N	Two-component regulator propeller, Y_Y_Y
BFV93_1770	Y	N	PAS S-box protein, PAS domain, PAS fold	BFV94_1775	Y	N	PAS S-box protein, PAS, PAS fold
BFV93_1889	Y	Y	HAMP	BFV94_1897	Y	Y	HAMP

BFV93_2053	Y	N		BFV94_2061	Y	N	
BFV93_2221	Y	Y	GAF	BFV94_2226	Y	Y	GAF
BFV93_2222	Y	N		BFV94_2227	Y	N	
BFV93_2401	Y	Y	Two-component regulator propeller, PAS S-box protein, PAS, PAS fold	BFV94_2406	Y	Y	
BFV93_2492	Y	Y		BFV94_2499	Y	Y	
BFV93_2889	Y	Y		BFV94_2901	Y	Y	
BFV93_2981	Y	N		BFV94_2993	Y	N	
BFV93_3092	Y	N		BFV94_3103	Y	N	
BFV93_3197	Y	N	response regulator receiver	BFV94_3207	Y	N	response regulator receiver
BFV93_3261	Y	N	PAS S-box protein, PAS, PAS fold	BFV94_3271	Y	N	PAS S-box protein, PAS, PAS fold
BFV93_3389	Y	N	7TM intracellular signaling, 7TMR-DISM extracellular 2	BFV94_3398	Y	N	7TM intracellular signaling, 7TMR-DISM extracellular 2
BFV93_3474	Y	Y	response regulator receiver	BFV94_3483	Y	Y	response regulator receiver
BFV93_3490	Y	Y		BFV94_3499	Y	Y	
BFV93_3520	Y	Y	PAS domain S-box protein, PAS	BFV94_3529	Y	Y	PAS S-box protein, PAS
BFV93_3580	Y	N	response regulator receiver	BFV94_3589	Y	N	response regulator receiver
BFV93_3921	Y	N		BFV94_3923	Y	N	
BFV93_3984	Y	N		BFV94_3987	Y	N	
BFV93_3992	Y	N		BFV94_3995	Y	N	
BFV93_4148	Y	N	PAS S-box protein, PAS, PAS fold, GAF	BFV94_4150	Y	N	PAS S-box protein, PAS, PAS fold, GAF
BFV93_4905	Y	N		BFV94_4642	Y	N	
BFV93_4914	Y	Y	PAS S-box protein, PAS, PAS fold	BFV94_4790	Y	Y	PAS S-box protein, PAS, PAS fold
BFV93_4965	Y	N		BFV94_4937	Y	N	
BFV93_1545	Y	Y		BFV94_1549	Y	Y	
BFV93_0411	Y	Y		BFV94_0412	Y	Y	
BFV93_3262	Y	N		BFV94_3272	Y	N	
BFV93_1451	Y	Y	PAS S-box protein, PAS, PAS fold	BFV94_1455	Y	Y	PAS S-box protein, PAS, PAS fold