Supporting Information

Table S1. Strains and Plasmids

Plasmids or strains	Description	Antibiotic Resistance	Reference
Strains			
E. coli			
HB101	F ⁻ Δ (gpt-proA)62 leuB6 supE44 ara-14 galK2 lacY 1 Δ (mcrC-mrr) rpsL20 (Str ^R) xyl-5 mtl-1recA13	Str ^R	Promega
XL1-Blue	F'::Tn10 $proA^{+}B^{+}$ $lacIq \Delta(lacZ)$)M15/ $recA1$ $endA1$ $gyrA96$ (NaI ^R) thi $hsdR17$ $(r_{K}^{-}m_{K}^{-+})$ $supE44$ $relA1$ lac	Amp ^R	Stratagene
R. capsulatus			
MT-RBC1 BK-RBC1	$\Delta(petABC::spe)$ $\Delta(petABC::spe) \Delta(recA::gen)$	Spe ^R Spe ^R Gen ^R	(28) This work
Plasmids			
pBluescript II pRK2013 pRK415 pPETI-S pPETI-F pWX1 pWX2	Cloning vector (pBS) Conjugation helper Broad host-range vector 3.4 kb fragment of <i>petABC</i> operon in pBR322 with <i>petB</i> tagged with Strep tag 3.4 kb fragment of <i>petABC</i> operon in pBR322 with <i>petB</i> tagged with Flag tag 1.9 kb of <i>recA</i> region in pBS Δ(<i>recA::gen</i>) allele with a <i>gen</i> cassette inserted between the SmaI and HindIII sites of <i>recA</i> 2 kb <i>gen</i> cassette inactivating <i>recA</i> Δ(<i>recA::gen</i>) in pRK415	Amp ^R Kan ^R Tet ^R Amp ^R Gen ^R	Stratagene (52) (52) (22) (22) This work This work
One Plasmid system			
pBK7	Fragment of NotI linker fused to <i>petB</i> and <i>petC</i> genes in pBS	Amp ^R	This work
pBK8	Fragment of NotI linker fused to <i>petB</i> and	Amp^R	This work

pBK6	petA genes in pBS 4.9 kb $petAB1^{wt}B2^{wt}C$) in pRK415- derivative; $petB2$ is tagged with the Strep	Kan ^R	This work
pBK22	epitope (<i>petB2-S</i>) Fragment of NotI linker fused to <i>petB::F144R and petA</i> in pBS	Amp^R	This work
pBK27	Fragment of NotI linker fused to	Amp^R	This work
pBK32	petB::H217L and petC s in pBS 4.9 kb petAB1 ^{F144R} B2 ^{H217L} C in pRK415- derivative; petB2::H217L is tagged with the Strep epitope (petB2-S)	Kan ^R	This work
Two plasmids system	and sureprepared (person s)		
pMTS1-S	3.4 kb fragment of <i>petABC</i> in pRK415; <i>petB</i> is tagged with the Strep epitope	Kan ^R	(22)
pBK21	(petB-S) 3.4 kb fragment of petABC in pRK415; petB is tagged with the Flag epitope	Tet ^R	This work
pBK18	(petB-F) 3.4 kb fragment of petAB ^{H217L} C in pPETI- F; petB::H217L is tagged with the Flag	Amp^R	This work
pBK19	epitope (<i>petB-F</i>) 3.4 kb fragment of <i>petAB^{F144R}C</i> in pPETI-S; <i>petB::F144R</i> is tagged with the Strep	Amp^R	This work
pBK25	epitope (<i>petB-S</i>) 3.4 kb fragment of <i>petAB</i> ^{F144R} C in pRK415; <i>petB</i> ::F144R is tagged with the	Kan ^R	This work
pBK28	Strep epitope (petB-S) 3.4 kb fragment of petAB ^{H217L} C in pRK415; petB::H217L is tagged with the Flag epitope (petB-F)	Tet ^R	This work

The *petB* is carboxyl-terminally epitope-tagged with Strep (*petB-S*) or Flag (*petB-F*) tags. Str^R, Amp^R, Spe^R, Gen^R, Kan^R, Tet^R correspond to streptomycin, ampicillin, spectinomycin, gentamicin, kanamycin and tetracycline resistance, respectively.

Table S2. Ps and Res colony forming abilities of strains producing mutant cytochrome bc_1 heterodimers. A, one-plasmid system, B two-plasmids system. See Table S1 for strain descriptions.

A.

		Dilution 10 ⁻⁴		Dilution 10 ⁻⁵		Dilution 10 ⁻⁶	
Strain	Medium	*Large (Ps)	Total (Res)	*Large (Ps)	Total (Res)	*Large (Ps)	Total (Res)
pBK6/MT-RBC1	MPYE	47	1725	3	462	0	63
	Med A	60	2818	10	444	1	43
pBK6/BK-RBC1	MPYE	2	1558	1	242	0	38
	Med A	1	1460	0	211	0	32

^{*}Large (Ps⁺) colonies obtained under Ps, and Total (Res), total numbers of colonies obtained under Res growth conditions; numbers indicate colony forming units (cfu)/ml

B.

Strain	Dilution 10 ⁻²		Dilution 10 ⁻³		Dilution 10 ⁻⁴		Dilution 10 ⁻⁵	
	**Large (Ps)	Total (Ps)	Large (Ps)	Total (Ps)	Large (Ps)	Total (Ps)	Large (Ps)	Total (Ps)
pBK25 + pBK28/MT-RBC1								
cyt <i>b</i> -S:F144R+cyt <i>b</i> -F: H217L	0	NC	1	NC	0	NC	0	782
pBK25 + pBK28/BK-RBC1								
cyt <i>b</i> -S:F144R+cyt <i>b</i> -F: H217L	1	NC	0	1158	0	184	0	13

^{**}Large (Ps⁺) colonies obtained under Ps, and Total (Ps) total numbers of colonies (*i.e.*, large and small) obtained under Res growth conditions; numbers indicate colony forming units (cfu)/ml; NC, not counted (too many colonies).