

Additional file 5: Bacterial strains and plasmids used in this work

Strain or plasmid	Important feature (s)	Source or reference
Strain		
<i>E. coli</i> strain DH5 α	F' Φ 80dlacZ Δ M15 Δ (lacZYA-argF)U169 <i>deoR recA1 endA1 hsdR17</i> (<i>r_k⁻</i> , <i>m_k⁺</i>) <i>phoA supE44 λ- thi-1 gyrA96 relA1</i>	Invitrogen
<i>E. coli</i> strain BW29427	<i>dap</i> auxotroph derivative of <i>E. coli</i> strain B2155	K. Datsenko and B. L. Wanner, unpublished
Δ <i>Ecfnr</i> mutant	<i>E. coli</i> K-12 substr. MG1655	K. Jung, unpublished
MSR-1 WT	Wild type R3/S1, but Rif ^r , Sm ^r	(1)
Δ <i>Mgfnr</i>	R3/S1 Δ <i>Mgfnr</i>	This study
MgFnrN27D	R3/S1 MgFnrN27D mutants	This study
MgFnrI34L	R3/S1 MgFnrI34L mutant	This study
MgFnrL98H	R3/S1 MgFnrL98H mutant	This study
MgFnrD153E	R3/S1 MgFnrD153E mutant	This study
Plasmid		
pBBR1MCS-2	Km ^r , mobilizable broad-host-range vector	(2)
pK19mobGII	-	(3)
pAL01	Km ^r , pK19mobGII vector (Km ^r , pMB-1 replicon, <i>gusA</i> , <i>lacZ</i>) containing a 2 kb fragment upstream of <i>mgr4019</i>	(4)
pAL02/2	Gm ^r , pT18mob2 vector containing a 2 kb fragment downstream of <i>mgr4019</i>	(4)
pLYJ87	pBBR1MCS-2 plus <i>nirS</i> promoter and <i>cre</i> fusion	(5)
pLYJ97	pBBR1MCS-2 plus <i>gusA</i> from pK19mobGII	(6)
pOR093	<i>mamX</i> CXXCH (65,104)->AXXAH, pK19mobGII derivative, Km ^r	(7)
pLYJ105	pAL02/2 plus <i>Mgfnr</i> 2-kb upstream region	This study
pLYJ106	pAL01 plus <i>Mgfnr</i> 2-kb downstream region	This study
pLYJ109	pLYJ97 plus <i>Mgfnr</i> promoter region	This study
pLYJ110	pBBR1MCS-2 plus <i>Mgfnr</i> with its own promoter region	This study
pLYJ132	pBBR1MCS-2 plus <i>Mgfnr</i>	This study
pLYJ141	pOR093 plus <i>MgfnrN27D</i>	This study
pLYJ142	pOR093 plus <i>MgfnrI34L</i>	This study
pLYJ143	pOR093 plus <i>MgfnrD153E</i>	This study
pLYJ144	pOR093 plus <i>MgfnrL98H</i>	This study
pLYJ153	pLYJ36 plus <i>Ecfnr</i>	This study

References

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