

Strains	Relevant features	Reference
<i>P. aeruginosa</i>		
PAO1	Wild-type <i>Pseudomonas aeruginosa</i>	Holloway Collection
Δ <i>siaA</i>	PAO1 with a 6 bp deletion in <i>siaA</i>	[1]
Δ <i>siaB</i>	PAO1 with a markerless <i>loxP</i> -site insertion in <i>siaB</i> (PA0171)	This study
Δ <i>siaC</i>	PAO1 with a markerless <i>loxP</i> -site insertion in <i>siaC</i> (PA0170)	This study
Δ <i>siaD</i>	PAO1 with a markerless <i>res</i> -site insertion in <i>siaD</i> (KO0169)	[1]
PW1292	MPAO1 with transposon phoAwp091G03 inserted at bp 311 of the coding region of the <i>siaB</i> gene[[2]
PW1290	MPAO1 with transposon phoAwp09q2A10 inserted at bp 181 of the coding region of the <i>siaC</i> gene	[2]
<i>E. coli</i>		
DH5 α	<i>fhuA2 lac(del)U169 phoA glnV44 Φ80' lacZ(del)M15 gyrA96 recA1 relA1 endA1 thi-1 hsdR17</i>	[3]
S17- λ pir	Tp ^r Sm ^r <i>recA, thi, pro, hsdR</i> -M ⁺ RP4: 2-Tc:Mu: Km ^r Tn7 λ pir	[4]
BL21(DE3) Rosetta T1R	<i>E. coli</i> str. B F ⁻ <i>ompT gal dcm lon hsdS_B(r_B⁻m_B⁻) λ(DE3 [<i>lacI lacUV5-T7p07 ind1 sam7 nin5</i>]) [<i>malB</i>⁺]_{K-12}(λ^S) <i>tonA</i></i>	NTU Protein Production Plattform
Plasmids		
pCR2.1	TOPO TA cloning plasmid, Amp ^r , Kan ^r	Invitrogen
pBBR	Broad host range expression plasmid pBBR1MCS-5, Gm ^r	[5]
pJEM1	Rhamnose inducible broad host range expression plasmid, Kan ^r	[6]
pNIC28-BSA4	Protein production plasmid, N-terminal His6, TEV-cleavable MHHHHHHSSGVDLGTENLYFQ*SM, Kan ^r	[7]
pNIC-CTHF	Protein production plasmid, Cleavable C-terminal His6-FLAG tag, Kan ^r	[7]
pNIC[SiaA]	pNIC28-BSA4 harbouring the partial <i>siaA</i> gene (C-terminal phosphatase domain from amino acid 386-663) for the production of a N-terminal His6-TEV-SiaA allele (SiaA-PP2C).	This study
pNIC[SiaB]	pNIC28-BSA4 for the production of a N-terminal His6-TEC SiaB allele	This study
pNIC[SiaC]	pNIC28-BSA4 for the production of a N-terminal His6-TEV-SiaC allele	This study
pJEM[SiaC]	pJEM1 harbouring the <i>siaC</i> gene for the production of a N-terminal His6-TEV-SiaC allele in <i>Pseudomonas aeruginosa</i> strains	This study
pCre1	Plasmid for the expression of Cre recombinase	[8]
Phage		
E79tv2	Generalised transducing phage	[9]

References

1. Klebensberger, J., Birkenmaier, A., Geffers, R., Kjelleberg, S. & Philipp, B. SiaA and SiaD are essential for inducing autoaggregation as a specific response to detergent stress in *Pseudomonas aeruginosa*. *Environ Microbiol.* 2009;11: 3073–3086.
2. Jacobs, M. A. *et al.* Comprehensive transposon mutant library of *Pseudomonas aeruginosa*. *Proc Natl Acad Sci.* 2003;100: 14339–14344.
3. Bethesda Research Laboratories. *E. coli* DH5 alpha competent cells. *Focus-Bethesda Res Lab.* 1986;8:9.
4. Simon, R., Priefer, U. & Pühler, A. A broad host range mobilization system for in vivo genetic engineering: Transposon mutagenesis in gram negative bacteria. *Bio/Technology* **1**, 784–791 (1983). Lämmli, U. K. Cleavage of Structural Proteins during the Assembly of the Head of Bacteriophage T4. *Nature.* 1970;227: 680–685.
5. Kovach, M. E. *et al.* Four new derivatives of the broad-host-range cloning vector pBBR1MCS, carrying different antibiotic-resistance cassettes. *Gene.* 1995;166: 175-176.
6. Jeske, M. & Altenbuchner, J. The *Escherichia coli* rhamnose promoter rhaP BAD is in *Pseudomonas putida* KT2440 independent of Crp–cAMP activation. *Appl Microbiol Biotechnol.* 2010;85: 1923–1933.
7. Savitsky, P. *et al.* High-throughput production of human proteins for crystallization: The SGC experience. *J Struct Biol.* 2010;172: 3–13.
8. Bailey, J. A. *et al.* Recent segmental duplications in the human genome. *Science.* 2002;297: 1003–1007.
9. Morgan, A. F. Transduction of *Pseudomonas aeruginosa* with a mutant of bacteriophage E79. *J Bacteriol.* 1979;139: 137–140.