

Strains	Genotype	Antibiotic resistance	Origin
<i>Escherichia coli</i> DH5 α	F- 80 <i>lacZ</i> M15 (<i>lacZYA-argF</i>)U169 <i>deoR recA1 endA1 hsdR17 phoA supE44</i> <i>-thi-1 gyrA96 relA1</i>		In vitrogen
<i>B. licheniformis</i> DSM 13 (ATCC 14580)			DSMZ
<i>B. halodurans</i> DSM 18197 (ATCC BAA-125)			DSMZ
<i>Bacillus subtilis</i> 168	<i>trpC2</i>		Anagnostopoulos and Spizizen 1961
BSmrs 111 (168, pDGbceR) BSGY005	168, pDGbceR 168, <i>amyE::PbceA::lacZ</i>	Km ^r Cm ^r	Joseph et al. 2002 Ohki et al. 2003
BSmrs 203 (pDGbceR)	BSGY005, pDGbceR	Cm ^r , Km ^r	This work
BSmrs 293 (Δ bceAB)	BSGY005, <i>PbceA::spc::yttA</i>	Sp ^c , Cm ^r	This work
BSmrs 297 (Δ bceAB, pDGbceR)	293, pDG148bceR	Sp ^c , Cm ^r , Km ^r	This work
BSmrs 294 (bceABBS_{ri})	BSGY005, <i>PbceA::his6-bceABBS::yttA</i>	Tc ^r , Cm ^r	This work
BSmrs 298 (bceABBS_{ri}, pDGbceR)	294, pDGbceR	Tc ^r , Cm ^r , Km ^r	This work
BSmrs 313 (ytsCDBL_{ri})	BSGY005, <i>PbceA::his6-ytsCDBL::yttA</i>	Tc ^r , Cm ^r	This work
BSmrs 318 (ytsCDBL_{ri}, pDGbceR)	313, pDGbceR	Tc ^r , Cm ^r , Km ^r	This work
BSmrs 312 (bceABBh_{ri})	BSGY005, <i>PbceA::his6-bceABBh::yttA</i>	Tc ^r , Cm ^r	This work
BSmrs 317 (bceABBh_{ri}, pDGbceR)	312, pDGbceR	Tc ^r , Cm ^r , Km ^r	This work
Bsmrs 357 (vraDESa_{ri})	BSGY005, <i>PbceA::his6-vraDESa::yttA</i>	Tc ^r , Cm ^r	This work
Bsmrs 361 (vraDESa_{ri}, pDGbceR)	357, pDGbceR	Tc ^r , Cm ^r , Km ^r	This work
Bsmrs 358 (vraFGSa_{ri})	BSGY005, <i>PceA::his6-vraFGSa::yttA</i>	Tc ^r , Cm ^r	This work
Bsmrs 362 (vraFGSa_{ri}, pDGbceR)	358, pDGbceR	Tc ^r , Cm ^r , Km ^r	This work
BSmrs 295 (bceBBSΔloop)	BSGY005, <i>PbceA::his6-bceBBSΔloop::yttA</i>	Tc ^r , Cm ^r	This work

BSmrs 299 (<i>bceBBs</i> Δ loop, pDG<i>bceR</i>)	295, pDG <i>bceR</i>	Tc ^r , Cm ^r , Km ^r	This work
BSmrs 336 (<i>ytsDBI</i> Δ loop)	BSGY005, <i>PbceA::his6-ytsCDBI</i> Δ loop:: <i>yttA</i>		
BSmrs 337 (<i>ytsDBI</i> Δ loop, pDG<i>bceR</i>)	336, pDG <i>bceR</i>		
BSmrs 296 (<i>bceB</i> loop BceBBs_{ri})	BSGY005, <i>PbceA::his6-bceABBs</i> loop BceBBs_{ri} :: <i>yttA</i>	Tc ^r , Cm ^r	This work
BSmrs 300 (<i>bceB</i> loop BceBBs_{ri} , pDG<i>bceR</i>)	296, pDG <i>bceR</i>	Tc ^r , Cm ^r , Km ^r	This work
BSmrs 315 (<i>bceBBs</i> loop YtsDBI_{ri})	BSGY005, <i>PbceA::his6-bceABBs</i> loop YtsDBI_{ri} :: <i>yttA</i>	Tc ^r , Cm ^r	This work
BSmrs 320 (<i>bceBBs</i> loop YtsDBI_{ri} , pDG<i>bceR</i>)	315, pDG <i>bceR</i>	Tc ^r , Cm ^r , Km ^r	This work
BSmrs 314 (<i>bceBBs</i> loop BceBBh_{ri})	BSGY005, <i>PbceA::his6-bceABBs</i> loop BceBBh_{ri} :: <i>yttA</i>	Tc ^r , Cm ^r	This work
BSmrs 319 (<i>bceBBs</i> loop BceBBh_{ri} , pDG<i>bceR</i>)	314, pDG <i>bceR</i>	Tc ^r , Cm ^r , Km ^r	This work
Bsmrs 316 (<i>bceBBs</i> loop YvcSBs_{ri})	BSGY005, <i>PbceA::his6-bceABBs</i> loop YvcSBs_{ri} :: <i>yttA</i>	Tc ^r , Cm ^r	This work
Bsmrs 321 (<i>bceBBs</i> loop YvcSBs_{ri} , pDG<i>bceR</i>)	316, pDG <i>bceR</i>	Tc ^r , Cm ^r , Km ^r	This work

To facilitate understanding strain names, indicated in bold characters, are used throughout the text and tables.

Anagnostopoulos C, Spizizen J. 1961. Requirements for transformation in *Bacillus subtilis*. J Bacteriol **81**: 741-746.

Joseph P, Fichant G, Quentin Y, Denizot F. 2002. Regulatory relationship of two-component and ABC transport systems and clustering of their genes in the *bacillus/clostridium* group, suggest a functional link between them. J Mol Microbiol Biotechnol **4**: 503-513.

Ohki R, Giyanto, Tateno K, Masuyama W, Moriya S, Kobayashi K, Ogasawara N. 2003. The BceRS two-component regulatory system induces expression of the bacitracin transporter, BceAB, in *Bacillus subtilis*. Mol Microbiol **49**: 1135-1144.