

TABLE S1 Bacterial strains, plasmids, and primers used in this study.

	Description	Reference
<i>Streptomyces</i> strains		
<i>S. coelicolor</i> M600	SCP1 ⁻ SCP2 ⁻	Kieser et al. 2000
<i>S. coelicolor</i> H2004	Δ <i>ddlA</i> (SCO5560):: <i>apr</i> , SCP1 ⁻ SCP2 ⁻	Novotna et al. 2012
<i>S. coelicolor</i> H2009	H2004 + pGN8	This study
<i>S. coelicolor</i> H2012	H2004 + pGN17	This study
<i>S. coelicolor</i> H2027	H2004 + pJ10257	This study
<i>S. coelicolor</i> H2077	M600 + pJ10257	Novotna et al. 2012
<i>S. coelicolor</i> H360	H2009 + pMK2	This study
<i>S. coelicolor</i> H361	H2012 + pMK2	This study
<i>S. coelicolor</i> J2175	Δ <i>vanRS</i> :: <i>apr</i> , SCP1 ⁻ SCP2 ⁻	Hutchings et al. 2006
<i>S. coelicolor</i> J3130	Δ <i>femX</i> :: <i>apr</i> , SCP1 ⁻ SCP2 ⁻	Hong et al. 2005
<i>E. coli</i> strains		
ET12567 (pUZ8002)	ET12567 containing helper plasmid pUZ8002	Paget et al. 1999
BW25113 (pIJ790)	BW25113 containing helper plasmid pIJ790	Gust et al. 2003
Plasmids		
pJ773	pBluescript KS(+) containing the apramycin resistance gene and <i>oriT</i> of Plasmid RP4, flanked by FRT sites (Ampicillin ^R)(Apramycin ^R)	Gust et al. 2003
pIJ790	Modified λ RED recombination plasmid pKD20 (Chloramphenicol ^R)	Gust et al. 2003
pIJ6902	integrative (ϕ C31 <i>attP-int</i>) and conjugative (<i>oriT</i> RK2), <i>tipAp</i> expression vector (Apramycin ^R)(Thiostrepton ^R)	Huang et al. 2005
pJ10257	integrative (ϕ BT1 <i>attP-int</i>) and conjugative (<i>oriT</i> RK2), <i>ermEp</i> expression vector (Hygromycin ^R)	Hong et al. 2005
pSET152	ϕ C31 <i>attP-int</i> derived integration vector for the conjugal transfer of DNA from <i>E. coli</i> to <i>Streptomyces</i> spp. (Apramycin ^R)	Bierman et al. 1992
pUZ8002	Non-transmissible <i>oriT</i> -mobilising plasmid (Kanamycin ^R)	Keiser et al. 2000
pHJH5	<i>ddlA</i> gene cloned into pGEM T-easy vector (Ampicillin ^R)	This study
pGN8	<i>ddlA</i> gene cloned into pJ10257 under <i>ermEp</i> control (Hygromycin ^R)	This study
pGN15	<i>vanA</i> gene cloned into pGEM T-easy vector (Ampicillin ^R)	This study
pGN17	<i>vanA</i> gene cloned into pJ10257 under <i>ermEp</i> control (Hygromycin ^R)	This study
pHJH4	derivative of pSET152 containing <i>vanJp-neo</i> (Apramycin ^R)	This study
pMK1	derivative of pSET152 containing <i>tsr</i> (Apramycin ^R)(Thiostrepton ^R)	This study
pMK2	derivative of pSET152 containig <i>tsr</i> and <i>vanJp-neo</i> (Apramycin ^R)(Thiostrepton ^R)	This study
Primers		
<i>ddlA</i> KO F	TCTCGAGGCACCGCGGGCGGGTACTCTCAACGCGATATGATTCCGGGGATCCGTCGACC	
<i>ddlA</i> KO R	GGGAGTCGCCTTCTCTGTGGTCACGACACGAAAGCGTCATGTAGGCTGGAGCTGCTTC	
<i>ddlA</i> KO Test F	TGAAGGAAGTGTATGTCGCGCA	
<i>ddlA</i> KO Test R	TTCCCGGACCAGACAGGAAAC	
<i>ndel-dlIA</i> F	CCGGCATATGATGAGCACCGAGAACCTCCCCCAGA	
<i>pacI-dlIA</i> R	CCGGTTAATTAATCAGCGCAGTCCGGTGGGCCGCC	
<i>ndel-vanA</i> F1	CCGGCATATGATGGCTGAGTCGGACAAGTTGGCCA	
<i>bln-vanA</i> R	CCGGCCTAGGTCATCGGAGCTTCCCCGTCAACGCC	
<i>qvanH</i> F	ACGAGAATCACGGTTTACGG	
<i>qvanH</i> R	CTTGTGGTCAATGCTGATGC	
<i>qvanK</i> F	CCGCAGTTCAAGTACGAGGT	
<i>qvanK</i> R	GGACGTAGAGGTCGTGGAAC	
<i>vanH</i> S1 FOR	TTCGACCTCTATATGAAGCGACGT	
<i>vanH</i> S1 REV	TGAGAGTCGCCTCGACGCCGAAA	
G1680	CACCGACCGGCAGGTGCGCG	
G1681	TCGAGCACCTCGCCCATGTC	

References

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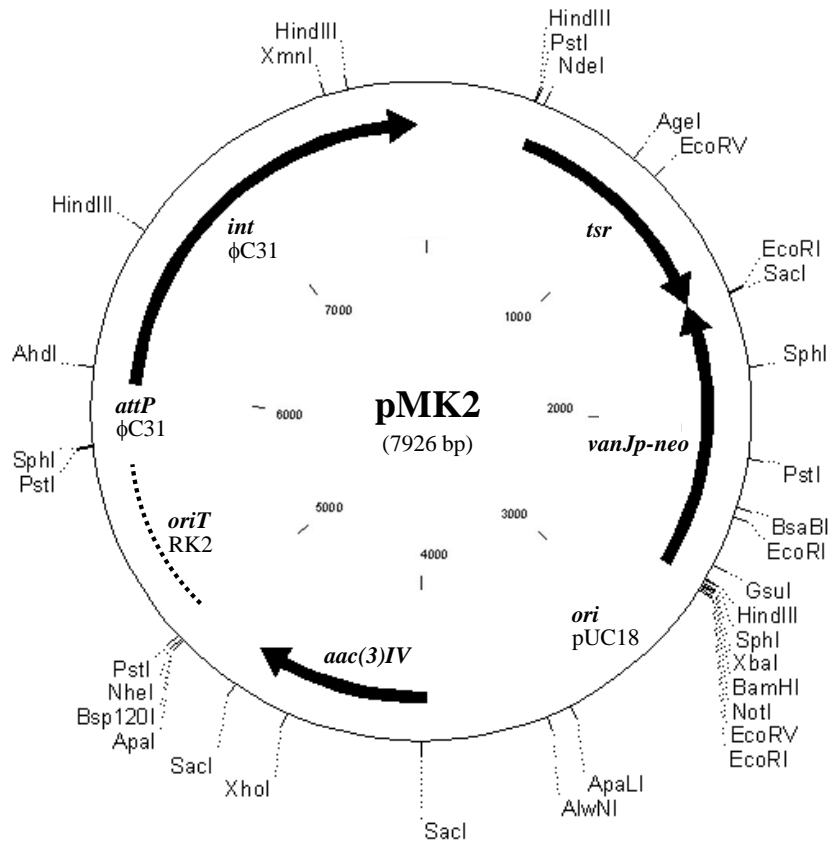


FIG S1 Integrative reporter plasmid pMK2 carrying the *neo*, neomycin/kanamycin resistance gene, expressed under the control of the vancomycin inducible *vanJ* promoter (*vanJp*). *aac(3)IV*, apramycin resistance gene; *tsr*, thiostrepton resistance gene; *oriT* RK2 for conjugation; *attP* and *int* for integration at the ϕ C31 phage attachment site.

construct	D-lac (10 mM)	Vancomycin (mg/L)								
		0	0.0075	0.0150	0.0300	0.0600	0.1250	0.2500	0.5000	1.0000
wild type (H2077)	-									
	+									
Δ <i>ddlA</i> (H102)	-									
	+									
Δ <i>ddlA</i> + <i>ermEp-ddlA</i> (H2009)	-									
	+									
Δ <i>ddlA</i> + <i>ermEp-vanA</i> (H2012)	-									
	+									

FIG S2 D-Lac supplementation alters the ratio of D-Ala-D-Ala:D-Ala-D-Lac-containing PG precursors in strains expressing the VanA ligase, and increases the concentration of vancomycin required for survival of the vancomycin-dependent Δ *ddlA* mutant strain. Phenotypic analysis of *S. coelicolor* wild type (H2077), Δ *ddlA* (H2027) and Δ *ddlA* complemented by the ligases encoded by *ddlA* (H2009) or *vanA* (H2012) expressed from the constitutive *ermEp*. Approximately 10^5 spores of each strain were spotted onto MMCGT plates containing different concentrations of vancomycin in the presence or absence of 10 mM D-Lac, as indicated. The result was scored after 4 days incubation at 30°C. Both complemented strains H2009 and H2012 grow in the absence of vancomycin and are unaffected by supplementation of the media with 10 mM D-Lac. This contrasts with Δ *ddlA* carrying only the empty pIJ10257 vector (H2027) which requires at least 0.25 mg/ml vancomycin for detectable growth in the presence of the D-Lac supplement, and 0.06 mg/ml in its absence. M600 harbouring pIJ10257 (H2077) is shown as a control.