## Characterization of FdmV as an amide synthetase for fredericamycin A biosynthesis in Streptomyces griseus ATCC 43944 Yihua Chen<sup>†</sup>, Evelyn Wendt-Pienkowski<sup>†</sup>, Jianhua Ju<sup>†</sup>, Shuangjun Lin<sup>†</sup>, Scott R. Rajski<sup>†</sup>,

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Running title: FdmV as an amide synthetase

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## SUPPLEMENTAL DATA

Table S1.  ${}^{1}$ H NMR and gCOSY data of FDM C in  $d_{6}$ -DMSOS2Figure S1. Construction and confirmation of the  $\Delta f dmV$  mutant strain SB4027S3Figure S2. Amino acid sequence comparison of FdmV with selected homologs and AS BS4Figure S3. Optimization of FdmV assay conditions with varying pH and DMSO concentrationS5Figure S4. Structures of benzoic acids tested as substrate analogs for FdmV-catalyzed amidationS5Figure S5. FdmV-catalyzed amidation of FDM M-3 to FDM M-6 as analyzed by HPLCS5

Position	FDM C In $a_6$ -DMSO (300 Hz)		
	$\delta_{\rm H} \left( J \left( {\rm Hz} \right) \right)$	gCOSY(H→H)	
1			
2			
3			
4			
5			
6			
7			
8			
9			
10			
11			
OCH3-11	3.94 (3H, s)		
12	6.93 (1H, s)		
13			
14			
15			
16			
17			
18			
19	2.80 (2H, br)	20	
20	2.69 (2H, br)	19	
21			
22	6.75 (1H, s)		
23			
24	4.25 (2H, s)		
25			
26	6.17 (1H, d, 15.5)	27	
27	7.24 (1H, dd, 15.5, 9.0)	26, 28	
28	6.31 (1H, m)	27	
29	6.32 (1H, m)	30	
30	1.85 (3H, d, 4.5)	29	
COOH-1	_a		
OH-3	12.69 (1H, s) <sup>b</sup>		
OH-6	12.85 (1H, s) <sup>b</sup>		
OH-10	_b		
OH-13	_b		
OH-17	_b		

TABLE S1.  $^{1}$ H NMR and gCOSY data of FDMC in  $d_{6}$ -DMSO.PositionFDM C in  $d_{c}$ -DMSO (500 Hz)

<sup>a</sup> not observed

<sup>b</sup> interchangeable.

FIGURE S1. Construction and confirmation of the  $\Delta fdmV$  mutant strain SB4027. (A) Inactivation of fdmV by replacing an internal fragment with the aac(3)IV Am resistance gene as in pBS4059 and construction of the  $\Delta fdmV$  mutant strain SB4027 via a double crossover homologous recombination event. B, BgIII; Bm, BamHI; M, MluI; P, PstI; S, SphI. (B) Confirmation of the genotype of the  $\Delta fdmV$  mutant strain SB4027 by Southern analysis of genomic DNAs digested with MluI and BamHI with the 0.8-kb BgIII-SphI fragment containing fdmU as a probe. Lane 1, S. griseus wild-type; lane 2, recombinant strain resulting from a single crossover event between the wild-type and pBS4059; lane 3, the desired  $\Delta fdmV$  mutant strain SB4027 as predicted from a double crossover event between the wild-type and pBS4059.



FIGURE S2. Amino acid sequence comparison of FdmV with selected homologs and AS B. The residues conserved in AS B active sites are marked with red asterisks. Accession numbers for each of the enzymes are: FdmV, AAQ08933; LlpA, CAM34350; PdmN, ABK58686; GrhP, AAM33677; RubR, AAM97368; OxyD, AAZ78328; TcsG, BAB12569; Ant-Orf1, ABW71832; PhzH, AAF17502; AS B, NP\_415200.

FdmV LlpA PdmN GrhP RubR OxyD TcsG Ant-Off1 PhzH AS B	$(1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) \\ (1) $	MORE G URADDERPEDE VALE AGAN GREDE GERGREVITUE RANDON DE GENERAL DE ALTONO SEDEALTING SINAAA RODIGENE EN D MORE G UPERDIS-EHSTVATIAALLARGED LANTIGECALAR RANDON DE GENERAL VALUE SUN VALUE SUN VALUE REAL DE MSEE G UDERDIS-EHSTVATIAALLARGED LANTIGECALAR RANDON DE GENERAL VALUE SUN VALUE SUN VALUE RANDON DE SANTA MSEE G UDERDIS-ERSTRETISTICO DE USER AGAN ALLE TETGAQANTITE GENERAL VILLO ALTONO DE GENERAL OLIVERTISTICO MSEE G UDERDIS-EDALVET AALLERGEDE E USER AGANA RALEL DE TETGAQANTITE GENERAL VILLO ALTONO DE SANTA MSEE G UDERDIS-EDALVET AALLERGEDE E USER AGANG ALTONO DE GENERAL VILLO ALTONO DE SANTAN RANDON DE SANTA MSEE G UDERDIS-EDALVET AALLERGEDE E USER AGANG ALTONO DE SANTAN RANDON DE SANTAN RANDON DE SANTAN RANDON DE SANTAN MSEE G UDERDIS-EDALVET AALLERGEDE E USER AGANG ALTONO DE SANTAN RANDON DE SANTAN RANDON DE SANTAN RANDON DE SANTAN MSEI G UDERDIS-GORINA CREDOL UT GERALAR RANDON DE SANTAN RANDON DE SANTAN RANDON DE SANTAN RANDON DE SANTAN MSEI G UDERDIS-GANANTITE RANDON DE SANTAN RANDON DE SANTAN RANDON DE SANTAN RANDON DE SANTAN RANDON DE SANTAN MSEI G UDERDIS-GANTAN RANDON DE SANTAN RANDON DE SANTAN RANDON DE SANTAN RANDON DE SANTAN RANDON DE SANTAN MSEI G UDERDIS-ANTANA RANDON DE SANTAN RANDON DE SANTAN RANDON DE SANTAN RANDON DE SANTAN MSEI G UDERDIS-ANTANA RANDON DE SANTAN RANDON DE SANTAN PERSONA DE SANTAN RANDON DE SANTAN MSEI G UDERDIS-ANTANA RANDON DE SANTAN RANDON DE SANTAN D
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FdmV LlpA PdmN GrhP RubR OxyD TcsG Ant-Orf1 PhzH AS B	(325) (319) (323) (326) (324) (319) (316) (326) (324) (324) (291)	TARELE E TANIÈ VARTUR COVERT FONDITI QA ACCER EN L'ALCONTRUIS COLLECTIONE CONTRUCTORE CONT
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FIGURE S3. Optimization of FdmV assay conditions with varying (A) pH (100 mM Tris-HCl) and (B) DMSO concentration.



FIGURE S4. Structures of benzoic acids tested as substrate analogs for FdmV-catalyzed amidation.



FIGURE S5. FdmV-catalyzed amidation of FDM M-3 to FDM M-6 as analyzed by HPLC: (I) completed assay with boiled FdmV as a control and (II) completed assay with FdmV under the optimized assay condition with L-Gln as an ammonia source for 3 hrs. (•) FDM M-3; (•) FDM M-6.

