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# Supporting Information

# Rifamorpholines A-E, Potential Antibiotics from a Locust-Associated

#### Amycolatopsis sp. Hca4

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Name	No. of amino acids	Name of Rif	Proposed function	identity /similar ity	accession no.	
Orf-2	486		hypothetical protein	76/83	SEB34960	
Orf-1	74		hypothetical protein	65/70	EOD68504	
RMPC	1303	RpoC	DNA-dependent RNA polymerase $\beta$ '-subunit	99/99	AAS07761	
RMPB	1167	RpoB	DNA-dependent RNA polymerase $\beta$ -subunit	99/99	AAS07760	
RMP36	419	Orf36	regulatory protein	94/95	AAS07758	
RMPJ	151	RifJ	aminodehydroquinate dehydratase	95/97	AAS07762	
RMP16	420	Orf16c	cytochrome P450 monooxygenase	94/96	AAC01740	
RMP15a	303	Orf15B	putative transketolase B subunit	93/96	AAC01739	
RMP15b	230	Orf15A	putative transketolase A subunit	95/97	AAS07757	
RMP14	272	Orf14	C-27 O-methyltransferase	96/97	AAC01738	
RMP13	424	Orf13c	putative cytochrome P450 monooxygenase	97/97	AAC01737	
RMPR	256	RifR	thioesterase	91/94	AAG52991	
RMP20	403	Orf20c	25-O-acetyltransferase	86/92	AAG52990	
RMP19	500	Orf19c	3-(3-hydroxylphenyl) propionate hydroxylase	91/95	AAG52989	
RMP18	473	Orf18	putative 2,3-dehydratase	96/98	AAG52988	
RMP17	356	Orf17	putative alpha-chain alkanal monooxygenase	91/96	AAG52987	
RMP11	294	Orf11	putative flavin-dependent oxidoreductase	89/94	AAC01735	
RMP10	330	Orf10c	putative dNTP-hexose 3-ketoreductase	88/93	AAC01734	
RMP9	290	Orf9c	dNTP-hexose aminotransferase	93/97	AAC01733	
RMP8	215	Orf8	putative dNTP-hexose 3,5-epimerase	93/96	AAC01732	
RMP7	403	Orf7	putative dNTP-hexose glycosyltransferase	95/97	AAC01731	
RMP6	435	Orf6c	putative dNTP-hexose dehydratase	98/99	AAC01730	
RMP5	421	Orf5c	putative cytochrome P450 monooxygenase	97/98	AAC01729	
RMP4	403	Orf4c	putative cytochrome P450 oxidoreductase	95/98	AAC01728	
RMP3	165	Orf3c	Pyridoxamine 5'-phosphate oxidase	93/97	SEB30227	
RMPQ	241	RifQ	putative tetR-like transcription regulatory protein	95/97	AAC01726	
RMPP	523	RifP	efflux transporter protein	87/92	AAC01725	
RMP2	310	Orf2	putative esterase	89/91	AAC01724	
RMPO	254	RifO	putative regulatory protein	93/96	AAC01723	
RMPN	300	RifN	kanosamine kinase	90/93	AAC01722	
RMPM	232	RifM	phosphatase	96/96	AAC01721	
RMPL	356	RifL	oxidoreductase	94/97	AAS07754	
	200	D'07		06/00	1 1 001 700	

Table S1. Annotation of	the Rifamorpholines biosynthetic gene cluster
No. of	ident

RMPL	263	RifL	aminoquinate dehydrogenase	87/92	AAC01719
RMPH	443	RifH	aminoDAHP sythase	89/91	AAC01718
RMPG	351	RifG	aminodehydroquinate synthase	92/93	AAC01717
RMP1	70	Orf1	hypothetical protein	77/91	AAC01716
RMPF	258	RifF	amide synthase (N-acyl transferase)	93/97	AAC01715
RMPE2	2363	RifE	rifamycin polyketide synthase protein	88/91	AAC01714
RMPE1	1033	RifE	rifamycin polyketide synthase protein	91/94	AAC01714
RMPD2	1163	RifD	rifamycin polyketide synthase protein	85/89	AAC01713
RMPD1	547	RifD	rifamycin polyketide synthase protein	96/98	AAC01713
RMPC	1746	RifC	rifamycin polyketide synthase protein	90/93	AAC01712
RMPB	5023	RifB	rifamycin polyketide synthase protein	91/94	AAC01711
RMPA2	1571	RifA	rifamycin polyketide synthase protein	91/94	AAC01710
RMPA1	3132	RifA	rifamycin polyketide synthase protein	89/93	AAC01710
RMP0	396	Orf0	cytochrome-P450-like protein	96/97	AAC01709
RMP35	71	Orf35	hypothetical protein	94/95	AAS07753
RMPT	259	RifT	putative NADH-dependent dehydrogenase	84/90	AAC01707
RMPS	330	RifS	putative NADH-dependent dehydrogenase	94/96	AAS07752
RMP31	328	Orf31	putative integral membrane protein	83/86	AAS07750
RMP30	191	Orf30	putative membrane protein	91/95	AAS07749
RMP29	421	Orf29	putative secreted protein	96/98	AAS07748
RMP28	391	Orf28	putative secreted protein	95/97	AAS07747
RMP27	394	Orf27	putative secreted protein	96/98	AAS07746
RMP26	328	Orf26	putative lipoprotein	97/98	AAS07745
RMP25	342	Orf25	putative lipoprotein	97/98	AAS07744
RMP24	441	Orf24	putative secreted protein	96/99	AAS07743
RMP23	278	Orf23	putative ABC-transporter permease	99/99	AAS07742
RMP22	250	Orf22	putative ABC-transporter integral membrane protein	99/100	AAS07741
RMP21	391	Orf21	putative ABC transporter ATP-binding protein	98/98	AAS07740
RplI	128	RplI	ribosomal protein L7/L12	93/94	AAS07739
RplJ	184	RplJ	ribosomal protein L10	98/99	AAS07738
Orf+1	215		hypothetical protein	83/86	WP_013222
	315				528
Orf+2	257		alpha/beta hydrolase	95/97	WP_003079
	237				255







Figure S1. Key 2D NMR correlations of 3-5.





chenolin A
(anti-inflammatory activity)

b)







-11.3788





**Figure S6.** HSQC NMR spectrum of **1** in DMSO- $d_6$  (600M Hz)



Figure S8. NOESY NMR spectrum of 1 in DMSO- $d_6$  (600M Hz)



**Figure S10.** <sup>13</sup>C NMR spectrum of **2** in acetone- $d_6$ (150M Hz)



**Figure S12.** HSQC NMR spectrum of **2** in acetone- $d_6$  (600M Hz)



Figure S13. HMBC NMR spectrum of 2 in acetone- $d_6$ (600M Hz)



Figure S14. NOESY NMR spectrum of 2 in acetone- $d_6$  (600M Hz)



**Figure S16.** <sup>13</sup>C NMR spectrum of **3** in acetone- $d_6$  (100M Hz)



**Figure S18.** HSQC NMR spectrum of **3** in acetone- $d_6$  (400M Hz)





-e

-₽

**Figure S20.** NOESY NMR spectrum of **3** in acetone- $d_6$  (400M Hz)



**Figure S22.** <sup>13</sup>C NMR spectrum of **4** in acetone- $d_6$  (150M Hz)



Figure S24. HSQC NMR spectrum of 4 in acetone- $d_6$  (600M Hz)









**Figure S30.** HSQC NMR spectrum of **5** in acetone- $d_6$  (400M Hz)



Figure S32. NOESY NMR spectrum of 5 in acetone- $d_6$  (400M Hz)

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