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Antibiotic evaluation and in vivo analysis of alkynyl Coenzyme A antimetabolites in Escherichia coli

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ABSTRACT

Pantothenamides have been the subject of much study as potential inhibitors of CoA and carrier protein dependent biosynthetic pathways. Based on an initial observation of growth inhibition in Escherichia coli by **3**, we have synthesized a small panel of pantetheine analogues and re-examined the inhibitory properties of this class of antibiotics with an emphasis on understanding the ability of these compounds to act as substrates of native CoA and carrier protein utilizing biosynthetic pathways. Our findings suggest that a secondary structure-activity relationship is an important factor in the antibiotic activity of these compounds.

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Coenzyme A (CoA) is an essential cofactor in living systems, functioning as the major acyl group carrier in numerous metabolic pathways.¹ The bacterial CoA biosynthetic pathway was first elucidated in Escherichia coli, where it was shown that five enzymes (CoaA, CoaB, CoaC, CoaD, and CoaE) are responsible for the conversion of pantothenate (vitamin B₅) to CoA (Fig. 1A).² While this pathway also exists in humans, comparison of the individual enzymes of the prokaryotic and eukaryotic CoA biosynthetic pathways has shown that the two demonstrate very little sequence homology.³ With reports of bacterial resistance to traditional antibiotics becoming increasingly common, the design of selective inhibitors of bacterial CoA biosynthesis presents a potentially novel antibiotic target.⁴

In addition to functioning directly as an acyl group carrier, CoA is also the source of the 4'-phosphopantetheine arm used for acyl group transfer in fatty acid biosynthesis (Fig. 1A).⁵ In this pathway, a phosphopantetheinyltransferase (PPTase) enzyme functions to transfer the 4'-phosphopantetheine arm of CoA to a conserved serine residue of an apo-acyl carrier protein (ACP). The free thiol of this posttranslational modification is then used as the site of acyl-intermediate tethering during the loading, condensation, and reduction reactions necessary for production of fatty acids. ACPs and peptidyl carrier proteins (PCPs) are used similarly in polyketide and nonribosomal peptide biosynthesis.⁶

In the past several years, it has been shown that many PPTases, most notably Sfp from the surfactin synthetase pathway in B. subtilis, have a relaxed substrate specificity which allows for the mod-

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ification of ACPs with CoA analogues in vitro.⁷ When derivatized with fluorescence or affinity tags, this property can be used for the selective visualization and isolation of carrier protein utilizing biosynthetic enzymes.⁸ Our group has had a long-standing interest in using PPTase promiscuity as a strategy for the investigation of primary and secondary biosynthetic pathways in bacterial organisms, which has lead us to investigate methods for the manipulation of intracellular CoA pool as a means of labeling carrier proteins. As CoA analogues cannot cross the cell membrane due to their strong negative charge, we have examined the utility of CoA precursors as in vivo carrier protein labels.⁹

Perhaps the most thoroughly investigated CoA precursors to date have been the antibacterial pantothenamides.¹⁰ This class of antibiotics, typified by N5-Pan (1), has been shown to inhibit E. coli and Staphylococcus aureus growth.¹¹ Originally postulated as inhibitors of the pantothenate kinase (CoaA) enzyme, it has since been shown that these compounds act as competitive substrates of CoaA, and are believed to exert their antibacterial activity through interference with fatty acid biosynthesis by labeling of the E. coli fatty acid ACP (Fig. 1B).^{12,13}

During our own studies of CoA precursors we encountered an interesting phenomenon relevant to the continued development and study of this antibiotic class. Through the synthesis and evaluation of a large number of CoA precursors (henceforth referred to as pantetheine analogues), we identified a single compound (2) capable of modifying the E. coli fatty acid ACP in the native organism (Fig. 1B).¹⁴ Further, these studies showed that compound 2 was non-toxic to E. coli grown in minimal media at concentrations >1 mM, while a structurally similar alkynyl pantetheine analogue (3) possessed cytotoxic activity near equivalent to that of 1

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Figure 1. (A) Pathway for CoA biosynthesis and posttranslational modification of carrier proteins. (B) Pantetheine analogues can act as alternate substrates for these pathways to form CoA antimetabolites. These CoA analogues can be processed by PPTases to form *crypto*-carrier proteins. (C) Structures of pantothenate and pantetheine analogues (1–15) examined in this study.

(Table 3). These observations ran contrary to expectations, as it was anticipated that the most toxic pantetheine analogues would be those which labeled the *E. coli* fatty acid ACP most efficiently, according to the proposed mode of action of these compounds.¹³

These findings lead us to synthesize a small panel of pantetheine analogues in order to examine in detail the influence of chain length (**4–9**), oxidation state (**10–11**), and functionality (**12–15**) on the ability to serve as CoA antimetabolites and inhibit growth in *E. coli* (Fig. 1C).¹⁵ We first tested these compounds as substrates of CoaA, the first enzyme of CoA biosynthetic pathway and the rate-limiting step for the accumulation of CoA metabolites.¹⁶ In past studies, we have correlated good turnover by CoaA with in vivo CoA production and carrier protein modification.¹⁷ As can be seen in Table 1, many of the analogues exhibit good turnover with some (**1–3**, **5**, **10–11**, **13**) approaching the kinetic efficiency of pantothenate. When examining the effect of chain length on turnover of alkynyl pantetheines, **4** shows very poor catalytic efficiency due mainly to its high K_m . This is indicative of weak binding, a conclusion consistent with the crystal structure of *E. coli* CoaA which shows the terminal acid of pantothenate having an important role in binding. This can be replaced by interaction with an amide which is missing in compound **4**.⁴ In contrast, **3** and **5** show catalytic efficiencies in the range of the native substrate, while **6–9** show a decrease in efficiency that correlates to the length of the

Table 1

Kinetic data for CoaA turnover of natural substrate (pantothenate) as well as pantetheine analogues 1-15^a

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Compound #	<i>K</i> _m (μM)	$V_{\rm max}~(\mu{ m M~s^{-1}})$	$k_{\rm cat} ({ m min}^{-1})$	$k_{\rm cat}/K_{\rm m}~({\rm mM}^{-1}~{ m min}^{-1})$			
Pantothenate	35.81 (±6.00)	0.31 (±0.01)	0.76 (±0.03)	21.13 (±1.93)			
1	33.98 (±7.11)	0.34 (±0.02)	0.84 (±0.05)	24.59 (±2.88)			
2	43.51 (±3.4)	0.28 (±0.002)	0.69 (±0.01)	16.1 (±0.53)			
3	36.04 (±6.05)	0.31 (±0.02)	0.77 (±0.04)	21.38 (±2.11)			
4	396.50 (±123.7)	0.26 (±0.05)	0.64 (±0.12)	1.62 (±0.61)			
5	34.04 (±10.72)	0.33 (±0.03)	0.81 (±0.07)	23.78 (±4.23)			
6	40.34 (±12.95)	0.29 (±0.03)	0.72 (±0.07)	17.95 (±3.41)			
7	40.00 (±14.26)	0.29 (±0.03)	0.72 (±0.07)	17.97 (±3.75)			
8	47.80 (±14.62)	0.23 (±0.02)	0.56 (±0.05)	11.70 (±2.22)			
9	48.49 (±10.99)	0.22 (±0.02)	0.54 (±0.04)	11.11 (±1.53)			
10	41.16 (±5.94)	0.38 (±0.02)	0.94 (±0.04)	22.85 (±1.96)			
11	36.91 (±7.52)	0.37 (±0.02)	0.91 (±0.05)	24.65 (±2.88)			
12	34.57 (±6.07)	0.22 (±0.01)	0.53 (±0.03)	15.43 (±1.52)			
13	27.78 (±5.21)	0.26 (±0.01)	0.65 (±0.03)	23.49 (±2.33)			
14	33.70 (±3.98)	0.25 (±0.01)	0.62 (±0.02)	18.29 (±1.20)			
15	57.97 (±12.01)	0.26 (±0.02)	0.64 (±0.04)	10.96 (±1.49)			

^a Values are means of three experiments, standard deviation is given in parentheses.

analogue. This decrease in efficiency is from both binding and turnover, as K_m increases and V_{max} decreases with the growing length of the alkynyl pantetheine substrates. Compounds **8** and **9** highlight another interesting trend seen in this panel, which is the detrimental effect of an amide bond at the natural thiol position of pantetheine on CoaA turnover. This effect is most readily observed by comparison of **12**, **13**, and **14**, which demonstrates a steady decrease in catalytic efficiency with decreasing polarization of the carbonyl (amide < carbamate < ester) at this position (see Table 1).

In comparing what effect degree of unsaturation has on turnover of pantetheine analogues by CoaA, the overall trend appears to indicate marginally improved turnover for fully saturated alkyl-pantetheine analogues. While this effect is slight among those pantetheine analogues terminating in propyl-derived chains (**3**, **10**, **11**), it can be clearly observed upon comparison of **1** to **6**. A similar effect is observed on comparison of amide bond-extended analogues **12** and **15**.

While kinetic values of pantetheine analogues with CoaA are a good predictor of in vivo activity, many other factors, including cell permeability and susceptibility to efflux pumps, impact the performance of antimetabolites when administered to living cells. To analyze the ability of these compounds to be processed by the CoA biosynthetic pathway in vivo and to interact with carrier proteins as CoA analogues, we utilized an in vivo assay.¹⁷ This assav provides a qualitative measure of the ability of pantetheine analogues to be processed by the endogenous CoA biosynthetic enzymes of E. coli by coupling CoA analogue production to the modification of a carrier protein. To facilitate detection and isolate CoA biosynthesis from variables such as carrier protein expression and PPTase promiscuity, E. coli are first transformed with expression plasmids for a carrier protein, in this case the Fren-ACP from the frenocylin polyketide synthase, as well as the PPTase Sfp, which is known to have a very broad substrate specificity. After growth to mid-log phase, the pantetheine analogue (1-15) is added at the same time as IPTG, which induces expression of the reporter system.¹⁵ Compounds that exhibit uptake and processing by the native E. coli CoA biosynthetic pathway produce modified ACPs, which demonstrate a mass shift characteristic of posttranslational modification by each analogue, and can be observed by MALDI-TOF (Table 2).

Having confirmed that the majority of these compounds are capable of formation of CoA analogues in vivo, we sought to correlate our findings with their antibacterial activity in native *E. coli*. To investigate the effects that additives present in the media might

Table 2 MALDI-TOF data for in vivo modification of Fren-ACP by compounds 1-15

Compound #	аро	Modified	Difference	Expected
None (control)	8663	9000 ^a	337 ^a	342 ^a
1	8660	9011	351	351
2	8663	9018	355	358
3	8663.7	8982	318.3	319
4	8661.1	_	_	248
5	8664.9	8996	331.1	333
6	8662.2	9008	345.8	347
7	8665.8	9025.9	360.1	361
8	8663.4	9065	401.6	404
9	8666	9042 & 9350 ^b	376 ^b	376
10	8658	8979	321	321
11	8660	8982	322	323
12	8660	9042	382	380
13	8660	9046	386	381
14	8660	9058	398	396
15	8666	9039	373	376

^a Modification by native CoA of *E. coli* results in the expected mass shift.

^b Compound **9** gives a large peak at 9350 in addition to the expected peak at 9042, possibly due to matrix interactions with the unsaturated activated alkyne.

have on antibiotic activity, we determined the MIC values for **1–15** using *E. coli* K12 grown in both minimal (M9) media, as well as in a richer, 1% tryptone broth which had been used to determine MIC values in an earlier study of pantothenamides.^{11,13}

Inspecting the results, all of the analogues tested showed greater growth inhibition in minimal media compared to rich media (Table 3). These results show a direct correlation between toxicity and CoaA kinetic profile for these compounds. This is to be expected, as it has previously been shown that CoaA is the rate-limiting step for CoA biosynthesis in vivo, and the antibacterial activity of these compounds is believed to be dependent on their in vivo transformation to CoA analogues. The major outliers in this respect are **2** and **13**, which possess good kinetics but do not show inhibition of E. coli at concentrations up to 500 µM even in minimal media. Additional evidence that these compounds act as CoA antimetabolites was provided by the observation that the inhibitory effects of the most toxic members of this panel (1, 3, 5-6, 10-11) were greatly decreased by addition of the CoA precursors pantothenate and β-alanine to the growth medium (Table 3). Among the alkynyl analogues which initially inspired this study (3, 5-9), an increasing MIC value is observed with growing chain length, mirroring the decline in catalytic efficiency observed among this group. Interestingly, among alkyl pantetheine analogues of the same chain-length (3 and 11, 1 and 6), changing the oxidation state from an alkyne to a saturated alkyl chain lowers the MIC by a factor of two to four. However, while 11 is sixfold more active than 1 in minimal media, administration of these same compounds to E. coli grown on rich media shows **11** to be at least $10 \times$ less toxic under these conditions.

The kinetic profiles, in vivo analysis, and inhibitory data generated here all support the previously held hypothesis that the antibiotic activity of pantetheine analogues is due to the production of CoA analogues in vivo.¹² However, the finding that saturated and unsaturated pantetheine analogues demonstrate rates of CoaA turnover within error of one another (i.e., 3 and 11), yet show drastically different MICs suggests that CoA analogue production alone is not sufficient for antibacterial activity. Based on our results it appears that of the pantetheine analogues processed efficiently by CoaA, those terminating in fully saturated alkyl groups are ideal for activity. while substitution by unsaturated alkynyl chains and polar terminal groups on the pantetheine chain (i.e., 2 and 13) results in decreased or no growth inhibition. This suggests a secondary structure-activity relationship for pantetheine analogue inhibition, in which one set of structural characteristics is necessary for biosynthetic processing and formation of CoA or ACP-analogues in vivo, while the identity

Table 3

Minimum inhibitory concentrations of pantetheine analogues **1–15** to *E. coli* grown in different media and effect of additives

Media compound #	M9 MIC (μM)	Tryptone MIC (μM)	M9 + Pan ^a MIC (µM)	M9 + β-Ala ^b MIC (μM)
1	6	50	500	250
2	>500	>500	nd ^c	nd
3	4	>500	>500	>500
4	500	>500	nd	nd
5	7.5	>500	>500	>500
6	15	>500	>500	>500
7	500	>500	nd	nd
8	>500	>500	nd	nd
9	>500	>500	nd	nd
10	3	>500	250	125
11	1	500	500	250
12	>500	>500	nd	nd
13	500	>500	nd	nd
14	>500	>500	nd	nd
15	500	>500	nd	nd

^a Pan, addition of 1 mM pantothenate to growth medium.

^b β -Ala, addition of 1 mM β -alanine to growth medium.

^c nd, not determined.

of the terminal group facilitates interaction of the CoA or ACP-analogue with a biologically relevant target.

In *E. coli*, fatty acid ACP is estimated to be present in the cytosol at concentrations approaching 1 mM. This abundance may explain the observation that **2** is capable of ACP modification in native *E. coli* without toxicity. Why then do alkyl pantetheine analogues **1** and **10–11** show increased inhibitory properties? Based on their kinetics with CoaA, these analogues seem unlikely to modify ACP at substantially higher levels than **2** in vivo. More likely, and in agreement with the secondary structural features of antimicrobial pantetheine analogues observed here, is the hypothesis that the activity of alkyl pantetheine analogues to bind and disrupt the associated lower abundance enzymes of fatty acid biosynthesis. Further elucidation of this process may have important implications for design of new members of this antibiotic class.

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Supplementary data

Supplementary data associated with this article can be found, in the online version, at doi:10.1016/j.bmcl.2008.07.078.

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