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## Communications to the Editor

### Antimicrobial $\alpha,\alpha$ -Dialkylated Amino Acid Rich Peptides with *in-Vivo* Activity against an Intracellular Pathogen

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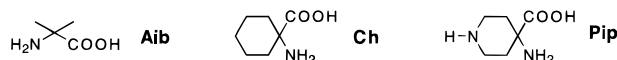
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The development of antibiotic resistance is threatening a return to the preantibiotic era.<sup>1</sup> This threat is hastened by the reliance on only a handful of distinct mechanisms of action among the vast majority of the commercially available drugs.<sup>2</sup> This problem alone is sufficient to warrant development of novel antimicrobial agents with unique modes of action.<sup>3</sup> In addition, new methods of controlling intracellular pathogens are needed because conventional antibiotic therapy and the host's immune response are not very effective against sequestered microorganisms.<sup>4</sup> *Brucella abortus* is an intracellular pathogen which lives and replicates in macrophages and quickly establishes chronic experimental infections in mice.<sup>5</sup> Herein, we report the *in-vitro* antibacterial and *in-vivo* anti-brucellae activity of a series of peptides (**1–10**).

Peptides **2–10** are composed of 50–80%  $\alpha,\alpha$ -dialkylated amino acids. The design of these peptides is loosely based on the natural antimicrobial peptides peptaibols.<sup>6–8</sup> Peptaibols have several  $\alpha$ -aminoisobutyric acid (Aib) residues, are acetylated on the N-terminus, and have an amino alcohol at the C-terminus.<sup>7</sup> *De-novo* peptides that include only a putative amphipathic helix have activity comparable to or greater than that of native antimicrobial peptides.<sup>10</sup> Amphipathic peptides with  $\geq 18$  residues can have very high cytotoxicity,<sup>11</sup> but simply shortening the peptides to 14 residues reduces cytotoxicity and retains much of the antimicrobial

1	Al-14	LysAlaAlaLysLysAlaAlaLysAlaAlaLysLysAlaAla-NH <sub>2</sub>
2	Ai-14	LysAibAibLysLysAibAibLysAibAibLysLysAibAib-NH <sub>2</sub>
3	Ai-11	LysLysAibAibLysAibAibLysLysAibAib-NH <sub>2</sub>
4	Ai-11-ac	Ac-LysLysAibAibLysAibAibLysLysAibAib-NH <sub>2</sub>
5	Ch-14	LysChChLysLysChChLysChChLysLysChCh-NH <sub>2</sub>
6	Ch-11	LysLysChChLysChChLysLysChCh-NH <sub>2</sub>
7	Ch-13	LysChChLysLysChChLysChChLysLysCh-----NH <sub>2</sub>
8	Ch-10	LysLysChChLysChChLysLysCh-----NH <sub>2</sub>
9	Pi-10	AibAibPipLysAibAibPipLysAibAib-NH <sub>2</sub>
10	Pi-10-ac	Ac-AibAibPipLysAibAibPipLysAibAib-NH <sub>2</sub>



activity.<sup>12</sup> Peptides **2–10** were synthesized to test the hypothesis that the Aib or Aib-like residues would stabilize helical conformations and retain biological activity as the peptides were further shortened.<sup>13</sup> In addition, peptides with several  $\alpha,\alpha$ -dialkylated amino acids throughout the sequence were expected to be resistant to enzymatic hydrolysis which is likely to enhance *in-vivo* activity.<sup>14</sup> Combinations of lysine with Aib or 1-amino-1-cyclohexanecarboxylic acid (Ch) and combinations of lysine with Aib and the novel amino acid 4-aminopiperidine-4-carboxylic acid<sup>15</sup> (Pip) were incorporated into peptides **2–10**.

Until recently, the use of Aib and Aib-like residues was limited due to the difficulty of coupling or harsh conditions used for their incorporation. Carpino's development of new coupling methods have allowed simple solid-phase synthesis of Aib or Aib-like rich peptides.<sup>16</sup> Peptide **1** was synthesized using standard Fmoc solid-phase synthesis methods.<sup>12</sup> Peptides **2–10** were synthesized via Fmoc chemistry using preformed acid fluorides on a PAL resin to yield the C-terminus amides. The only synthetic difficulty encountered came in the coupling of the second residue in the Ch-11,14 peptides. Fortunately, the major peptides obtained were also amphipathic and showed good *in-vitro* antibacterial activity. To maximize the number of  $\alpha,\alpha$ -dialkylated amino acids and retain a high level of positive charge

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**Table 1.** Peptide Antibacterial Activity<sup>a</sup> and Percent Helicity

peptide	<i>E. coli</i>	<i>S. aureus</i>	% helicity <sup>b</sup>
Al-14	>178	>178	ND <sup>c</sup>
Ai-14	5.5	11	42
Ai-11	55	>220	35
Ai-11-ac	6.6	212	ND <sup>d</sup>
Ch-13	4.5	2.2	61
Ch-11	5.2	2.6	ND <sup>d</sup>
Ch-10	5.7	2.8	43
Pi-10	7.7	123	43

<sup>a</sup> These MIC (in  $\mu$ M) are corrected for the actual peptide concentration using quantitative amino acid analysis according to ref 12. <sup>b</sup> The percent  $\alpha$ -helix =  $-100([\theta]_{222} + 3000)/33000$ ;  $[\theta]_{222} = [\theta]_{\text{obs}}(\text{MRW}/10\text{lc})$  according to ref 22 taken in 25 mM SDS micelles with 100–200  $\mu$ M peptide. <sup>c</sup> This peptide precipitates when the 25 mM SDS is added, precluding CD spectroscopic measurement. <sup>d</sup> Insufficient quantities of these peptides were made for CD spectroscopic measurements.

**Table 2.** Peptide-Mediated *B. abortus* Reductions in Chronically Infected BALB/c Mice

treatment (dose)	% bacterial reduction	significance <sup>a</sup>
Ai-14 (500 $\mu$ g)	73	<0.05
Ai-11 (500 $\mu$ g)	63	<0.05
Ai-11-ac (500 $\mu$ g)	30	<0.05
Pi-10 (500 $\mu$ g)	90	<0.05
Al-14 (1 mg)	−6	NS

<sup>a</sup> NS means not significant.

on the polar face of these peptides, the lysine-like  $\alpha$ , $\alpha$ -dialkylated amino acid Pip is introduced. This residue with the  $\gamma$ -amine protected as the Boc derivative allows the synthesis of an amphipathic peptide containing no natural amino acids. We have not yet attempted to synthesize a peptide devoid of natural amino acids because of the perceived problems in the synthesis and characterization. A peptide with only achiral  $\alpha$ , $\alpha$ -dialkylated amino acids would show no CD spectra because there would be no preference for a left- or right-handed helix.

The replacement of Ala residues in Al-14 with Aib residues in Ai-14 has a substantial effect on the biological activity. MIC assays were done with *Escherichia coli* and *Staphylococcus aureus*.<sup>12</sup> Ai-14 is at least 8–16 times more active than Al-14. Al-14 precipitates out of 25 mM SDS micelles, so a comparison of secondary structure is precluded. Shortening Ai-14 to make Ai-11 showed diminished but substantial activity against *E. coli* and no activity against *S. aureus*. While Aib helps to promote helical conformations, it is not a very hydrophobic residue. The Ch residue was introduced to determine whether greater hydrophobic character would increase the activity of peptides with otherwise similar designs. Unfortunately, sufficient amounts of the analogous peptide sequence were not produced due to coupling problems with the second Ch residue from the C-terminus. Ch-13 had essentially the same activity as its closest Aib peptide counterpart, Ai-14, but Ch-10 with only 10 residues had increased activity over Ch-13 and much higher activity than its closest Aib peptide counterpart, Ai-11. Pi-10 with only 10 residues and only two natural amino acids also showed good antibacterial activity against *E. coli* and moderate activity against *S. aureus*.

*In-vivo* peptide activity was tested using BALB/c mice infected with a virulent strain of *B. abortus*. Both the humoral and cell-mediated immune responses developed 3–4 weeks after the infection of the mice.<sup>5</sup> Both of these

responses are needed to control the infection, and seldom do they eliminate the infection because the organism is hidden within the macrophages.<sup>17–19</sup> The mice were treated intravenously with peptide, and 24 h later the spleens of the treated and control (saline) animals were cultured for viable organisms.<sup>20</sup> Percent bacterial reduction was determined by subtracting the total number of organisms in the treated spleens from the total number of organisms in the control spleens. This difference was divided by the total number of organisms in the control spleens and multiplied by 100. Infected mice treated with 100  $\mu$ g of Ch-10, Ch-13, or Pi-10 showed no significant differences compared to the controls. However infected mice treated with 100  $\mu$ g of the Ai series show some reductions compared to the controls, but the reductions were not statistically significant. Noninfected mice were unharmed, but infected mice treated with a single 500  $\mu$ g dose of Ch-10 or Ch-13 died immediately after treatment. As shown in Table 2, this dose of Pi-10 and the Ai peptides resulted in significant reductions of bacterial load compared to the controls. *B. abortus* is one of the few bacteria that are resistant to the direct antimicrobial effect of these peptides.<sup>12,21</sup> We speculate that the peptides are selectively lysing the infected macrophages exposing a portion of the *B. abortus* load to the immune system, which is then destroying the released bacteria. *In-vitro* peptide testing with infected and noninfected macrophages shows a selective destruction of infected macrophages.<sup>21</sup>

The selective destruction of a wide range of bacteria by amphipathic peptides relative to mammalian cells has been attributed to distinct differences in the cell membranes.<sup>12</sup> Bacteria have negatively charged phospholipids on the exterior membrane. Normal mammalian cells do not. Thus, one potential reason for the selective destruction of *B. abortus*-infected macrophages could be a greater negative charge on the exterior membrane of infected macrophages. Future studies are directed toward defining what membrane differences result in the selective destruction of infected macrophages.

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**Supporting Information Available:** *B. abortus* normal macrophage (Table A) and *B. abortus*-infected macrophage (Table B) susceptibility to peptide treatment, synthesis of 1-Boc-4-Fmoc(amino)piperidine-4-carboxylic acid, and a representative synthesis of an acid fluoride (5 pages). Ordering information is given on any current masthead page.

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