



# Control of the cytotoxicity of dansylated polytheonamide mimic, an artificial peptide ion channel, by modification of the N-terminal structure



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## ABSTRACT

We demonstrate that the cytotoxicity of dansylated polytheonamide mimic (**2**) is controlled by chemical modification of its N-terminal structure. Dansylated polytheonamide mimic (**2**) is an ion channel peptide which displays potent cytotoxicity against P388 mouse leukemia cells ( $IC_{50} = 12$  nM). To modulate its cytotoxicity, three analogues of **2**, possessing distinct N-terminal structures with different hydrophobicities, were synthesized and their cytotoxicities were evaluated. This focused structure–activity relationship study unveiled that the cytotoxicity of **2** is enhanced 10-fold by simply changing its N-terminal 5,5-dimethyl-2-oxohexanamide to the more hydrophobic palmitamide. The data obtained here provide new understanding for the functional control of the artificial ion channel peptide **2**.

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Polytheonamide B (**1**) is an ion channel forming peptide that displays extraordinarily potent cytotoxicity against P388 mouse leukemia cells ( $IC_{50} = 0.098$  nM).<sup>1</sup> The 48-mer peptide **1** possesses a D,L-alternating amino acid sequence with numerous non-proteinogenic amino acids (Fig. 1).<sup>2,3</sup> This unusual peptide sequence folds into a  $\beta^{6,3}$ -helix in a hydrophobic environment;<sup>4</sup> the resulting tubular structure is believed to function as a transmembrane ion channel in a lipid bilayer.<sup>5</sup> Motivated by these unusual features, we launched a research program to investigate the structure and function of **1**, and recently accomplished the total synthesis<sup>6</sup> and a structure–activity relationship study of **1**.<sup>7</sup> Furthermore, artificial peptide **2**, designated as dansylated polytheonamide mimic, was designed as a simplified analogue of **1**. The synthesis and functional analysis of **2**<sup>8</sup> were recently achieved.<sup>9,10</sup> Mimic **2**, which differs from the parent natural product **1** by six amino acid residues, was demonstrated to form an ion conducting channel in a lipid bilayer and exhibited potent cytotoxicity against P388 mouse leukemia cells ( $IC_{50} = 12$  nM), although its toxicity was approximately 100-times weaker than that of **1**.<sup>8</sup>

To create artificial peptides with more potent activities, we decided to enhance the cytotoxicity of **2** by structural modification. Here we report the synthesis and biological evaluation of N-terminal modified analogues of **2**, and discuss the strong correlation between the cytotoxicity and overall hydrophobicity. These

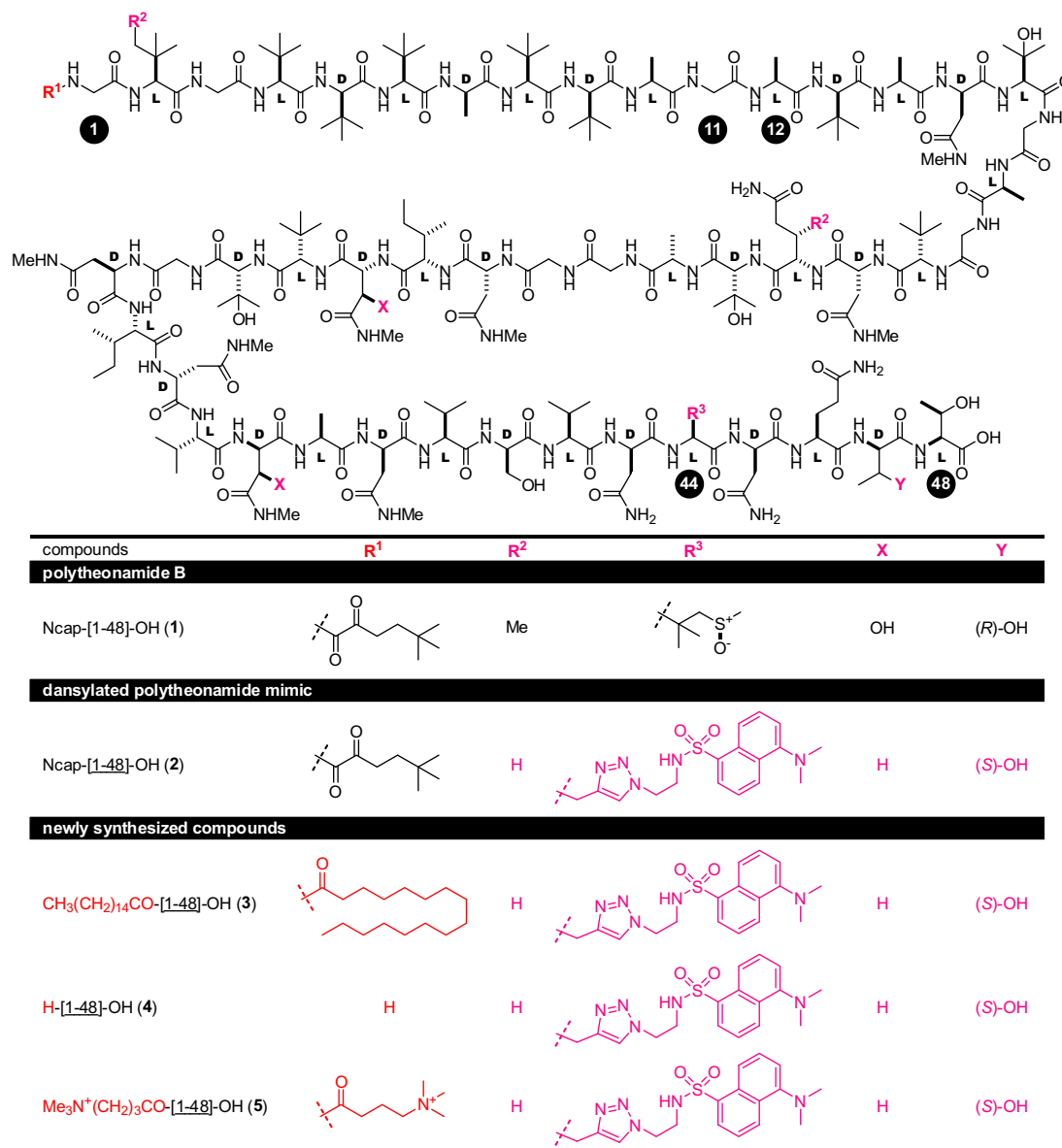
results provide a new design principle for functional control of artificial cytotoxins based on large polypeptide structures.

The 48 amino acid sequence of dansylated polytheonamide mimic **2** is capped at the N-terminus with 5,5-dimethyl-2-oxohexanoate (Ncap). The hydrophobic Ncap of **2** was hypothesized to be the critical substructure for its cytotoxicity because the hydrophobic moiety would be required for effective insertion into the cell membrane, where **2** would exert its cytotoxicity by forming an ion channel. To undertake experimental verification of this hypothesis, we synthesized palmitamide **3** (Fig. 1), amine **4**, and trimethylammonium derivative **5**, which were intended to possess different hydrophobicities.

Three analogues **3–5** were chemically constructed through fully-protected 48-mer peptide **10** (Scheme 1), which was prepared by condensation between residue 1–11 (**7**) and residue 12–48 (**9**). Construction of 37-mer peptide **9** was attained by a single automated solid-phase peptide synthesis as previously reported.<sup>8</sup> Fmoc-protected 11-mer peptide **7** was synthesized from Fmoc-Gly-Wang resin **6**<sup>11</sup> using Fmoc-based chemistry<sup>12</sup> with HATU/HOAt activation.<sup>13</sup> Automated peptide chain elongation and cleavage from the resin produced desired peptide **7** (20% overall yield from **6**), which was converted to the corresponding thioester **8** using a reagent combination of  $HS(CH_2)_2CO_2Et$ , HOBT, and  $N,N'$ -diisopropylcarbodiimide in 82% yield. Then, the two fragments **8** and **9** were coupled under Ag-mediated coupling conditions.<sup>14</sup> When thioester **8** and amine **9** were treated with  $AgNO_3$  and HOObt in DMF/THF at 50 °C, the full-length 48-mer polyamide **10** was obtained. Finally, peptide **10** was treated with

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**Figure 1.** Structures of polytheonamide B (1), dansylated polytheonamide mimic (2), and newly synthesized analogues 3–5. The sequences are abbreviated as R-[X-Y]-Z (R = functional group attached to the N-terminal amine, X = N-terminal residue number, Y = C-terminal residue number, Z = functional group attached to the C-terminal carboxylic acid). The residue numbers of 1 are written as plain text (R-[X-Y]-Z) and those of 2–5 are underlined (R-[X-Y]-Z).

piperidine/DMF to remove the N-terminal Fmoc group, then subsequently exposed to TFA/*i*-Pr<sub>3</sub>SiH<sup>15</sup>/H<sub>2</sub>O (95:2.5:2.5) to realize global deprotection (the eight Tmb, four Tr, and three *t*-Bu groups), successfully delivering amine 4 (20% overall yield from 37-mer peptide 9). Palmitamide 3 and trimethylammonium derivative 5 were transformed from 4 by applying the corresponding mixed anhydrides in 57 and 23% yields, respectively.

Having efficiently synthesized 3–5, the hydrophobicities of polytheonamide B (1), mimic 2, and the three analogues 3–5 were compared by accessing their octanol/water partition coefficients (log*P*) using reversed phase HPLC.<sup>16</sup> First, the retention times of the standard samples (acetoanilide, thymol, biphenyl, and phenanthrene) with known log*P* values (1.0, 3.3, 4.0, and 4.5, respectively)<sup>17</sup> were measured using an ODS column and gradient elution of *i*-PrOH/water as mobile phase.<sup>18</sup> Next, polytheonamide B (1) and peptides 2–5 were eluted under the same HPLC conditions to deduce their log *P* values by comparison of their retention times with the standard samples.<sup>19</sup> As shown in Table 1, the log *P*

value of the parent natural product 1 (4.5) was larger than that of polytheonamide mimic 2 (3.8), while replacement of 5,5-dimethyl-2-oxohexanamide of 2 with palmitamide in 3 increased the log*P* value from 3.8 to 4.9. On the other hand, amine 4 and trimethylammonium derivative 5 had lower hydrophobicity than 2 (log*P* = 3.2 and 2.6, respectively). Significant differences in the log*P* values of peptides 2–5, whose structures are same except at the N-terminus, clarified that the N-terminal structure is the crucial factor in controlling the total polarity of this series of peptides.

Next, the cytotoxicities of 1 and mimics 2–5 against P388 mouse leukemia cells were evaluated using the XTT method (Table 1).<sup>20,21</sup> Although they were less toxic than the exceptionally bioactive natural product 1, all the mimics exhibited potent cytotoxicities. Most importantly, palmitamide 3 had 10-fold stronger toxicity (IC<sub>50</sub> = 1.0 nM) than the original mimic 2, and thus was only 10-times less toxic than 1. Not unexpectedly, amine 4 and trimethylammonium derivative 5 displayed 2-fold and 7-fold lower cytotoxicities than 2 (IC<sub>50</sub> = 25 nM and 83 nM, respectively).<sup>8</sup>

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