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Down-regulating Proteolysis to Enhance Anticancer Activity of Peptide Nanofibers

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Abstract

Nanofibers of short peptides are emerging as a promising type of agents for inhibiting cancer cells. But the proteolysis of peptides decreases the anticancer efficacy of the peptide nanofibers. Here we show that decreasing the activity of proteasomes enhance the activity of peptide nanofibers for inhibiting cancer cells. Based on the structure of galactin-3, we designed a heptapeptide, which self-assembles to form nanofibers. The nanofibers of the heptapeptide exhibit moderate cytotoxicity to three representative cancer cell lines (HeLa, MCF-7, and HepG2), largely due to the proteolysis of the peptides. Using a clinically approved proteasome inhibitor, bortezomib, to treat the cancer cells significantly decreases the proteolysis of the peptides and enhances the activity of the peptide nanofibers for inhibiting the cancer cells. This work illustrates a promising approach for enhancing the anticancer efficacy of peptide nanofibers by modulating intracellular protein degradation machinery, as well as provides insights for understanding the cytotoxicity of aberrant protein or peptide aggregates in complicated cellular environment.

Graphical Abstract

Here we report the design and synthesis of a heptapeptide (1) based on the structure of galactin-3, which can self-assemble into nanofibers. While the proteolysis of the peptides leads to a moderate cytotoxicity of the molecule on cancer cells, co-incubation of a proteasome inhibitor BTZ with the peptide significantly decreases the proteolysis of the peptide and enhances the activity of the peptide nanofibers for inhibiting the cancer cells.

Supporting information for this article is available on the WWW



Keywords

anticancer; nanofibers; self-assemble; peptide; proteolysis

As the consequence of reaching thermodynamic equilibrium, self-assembly autonomously reorganizes components to form ordered structures. When small organic molecules self-assemble in solutions, the assemblies of the molecules usually build up the three dimensional networks to encapsulate water to result in a hydrogel.^[1] These small molecules, consequently, are classified as hydrogelators. Share common features such as π - π interactions, hydrogen bonding, hydrophobic interactions, charge interactions, and other non-covalent interactions, the hydrogelators self-assemble and usually form nanofibers. These nanofibers, having the similar morphology as endogenous nanofibers that are normal (e.g., actin filaments and microtubules) or aberrant (e.g., amyloid) cellular structures, are emerging as a new class of molecular entities for controlling the fate of cells.^[2]

Among the small molecules being explored for self-assembly, one attracting class of candidates is peptides. Peptide based hydrogelators,^[3] possessing inherent biological activities and structural diversity, are accessible in large quantity. Inspired by proteins that self-assemble into nanofibers to serve as critical components of cellular processes, research focusing on the self-assembly of peptides is attracting increasing attentions from scientists. [3a, 3d, 3g, 3h, 3k, 4] The supramolecular assemblies of peptides are potential candidates for biological applications such as 3D cell culture,^[5] wound healing^[6] and tissue engineering.^[7] Studies on the cytotoxicity of nanofibers formed by peptides reveals that nanofibers are able to inhibit cancer cells selectively,^[8] by interacting with the cancer cells intracellularly or pericellularly. In addition, the self-assembly of a peptide amphiphile, which contained a cytotoxic peptide fragment, efficiently delivers the cytotoxic peptide to cancer cells and inhibit the cancer cells.^[9] Recently, Yang et al.^[10] and Wells et al.^[11] reported nanofibers formed by the self-assembly of small molecules to inhibit cancer cells. Our group also has been exploring the anticancer properties of nanofibers on cancer cells and reported that the supramolecular assemblies of small molecules behaved differently from the molecule monomers and the assemblies can interact with cells and kill cancer cells.^[2d, 8] While nanofibers of peptides are becoming an active research frontier for developing anticancer agents, one of the less explored aspects is how the stability of peptides in biological

environment and the proteolysis of these peptides affect the activities of the peptide nanofibers against cancer cells.

In this study, encouraged by the previous studies on the inhibition of cancer cells by peptide nanofibers, we design and synthesize a peptide (Nap-Phe-Phe-His-Phe-Asn-Pro-Arg, 1 (Scheme 1)), and examine how reducing proteolysis affects the activity of the peptide nanofibers of 1 for inhibiting cancer cells. We choose to follow the structure of galectin-3^[12] because we want to increase the interaction of nanofibers and cell surface heparan sulfate for cellular uptake.^[13] Since Nap-Phe-Phe is an excellent motif to enable self-assembly and hydrogelation,^[14] we connect the peptide fragment His-Phe-Asn-Pro-Arg from galectin-3 to Nap-Phe-Phe for making the hydrogelator 1. The hydrogelator 1 self-assembles into nanofibers. The existence of nanofibers in cancer cells would induce cell death. Cell assays show that the co-incubation 1 with bortezomib (BTZ), a proteasome inhibitor,^[15] significantly increases the overall cytotoxicity of 1 in cancer cells. The analysis of cell lysates in the presence of BTZ indicates that the inhibition of proteasome, indeed, reduces the degradation of the peptides, supporting that more nanofibers exist in the cancer cell to lead to higher cytotoxicity. This work illustrates a promising approach for enhancing the anticancer efficacy of peptide nanofibers by modulating intracellular protein degradation machinery, as well as provides insights for understanding the cytotoxicity of aberrant protein or peptide aggregates in complicated cellular environment.

After making the hydrogelator **1** by solid phase peptide synthesis (SPPS),^[16] we purify the molecule by high-performance liquid chromatography (HPLC), with the overall yield of **1** about 80%. Molecules are characterized by ¹H-NMR (Figure S1–S3). Stable hydrogel forms when **1** (0.4 wt%) dissolves in water at pH7.4. As shown in Figure 1A, transmission electron microscopy (TEM) image of the hydrogel formed by **1** contains uniform and twisted nanofibers with a diameter at 10 ± 2 nm and a well-defined pitch at 52 ± 2 nm. We also test the hydrogelation of **1** at 0.4 wt% together with 1 equivalent of lactose. As shown in Figure 1, a more stable hydrogel is formed when adding 1 eq. of lactose. While the TEM image shows that there are nanofibers with a diameter at 10 ± 2 nm existed in the hydrogel of **1** and lactose, the lengths of pitch of the nanofibers become less uniform. This result indicates that **1** is able to interact with lactose, albeit moderately.

Rheological test of the hydrogels further elucidates the viscoelastic properties of the hydrogels formed by the heptapeptide. Figure 1C and 1D show the strain and frequency dependence of storage modulus (G') and loss modulus (G") of the hydrogels formed by **1** alone or in combination with 1 equivalent of lactose. The G' values of both the hydrogels are larger than their G" value; the G' and G" of both the hydrogels are nearly independent of frequency. These features confirm the viscoelastic features of the hydrogels. Hydrogel formed by **1** together with lactose shows a higher G' than that of the hydrogel of **1**, with the maximum storage modulus G' at 36 Pa and indicating that the hydrogel is stronger than the hydrogel formed by hydrogelator **1** alone. The value of critical strain of the hydrogel formed by **1** alone. Since part of the sequence of peptide **1** is originated from the saccharide binding domain (FHFNPR) of galactin-3,^[17] this peptide should have certain binding affinity to

carbohydrate such as lactose. So we prepare the hydrogel of **1** with 1 eq. of lactose to verify the interactions between **1** and lactose. The slight increase of the strength of the hydrogel agrees with the interaction between lactose and **1**. These rheological results indicate that the addition of lactose affects the viscoelasticity of the hydrogel and the self-assembling ability of the **1**. Such an observation also supports that there are interactions between the nanofibers of **1** and lactose and agrees with that the sequence in molecule **1** originates from the carbohydrate interacting motif in galactin-3.

After the gelation test of the hydrogelator, we test the cytotoxicity of **1** against HeLa (cervical cancer), MCF-7 (breast cancer), and HepG2 (liver cancer) cells. While 1 exhibits the highest cytotoxicity on MCF-7, with the IC₅₀ at 51 μ M (48 h), its cytotoxicity against HepG2 is lower (IC₅₀ = 233 μ M) than against MCF-7 cells (Figure S4), and 1 shows the lowest cytotoxicity against HeLa cells (IC₅₀ > 500 μ M). We then co-incubated 1 with BTZ^[18] to treat the aforementioned three cancer cell lines. The results (Figure S5) shows that the addition of BTZ significantly increases the cytotoxicity of **1** against the cancer cells, decreasing the IC₅₀ of 1 against MCF-7, HepG2, and HeLa cells to 38, 56, and 80 μ M, respectively. As shown by the summarized cytotoxicity of BTZ, 1 alone, or the combination of **1** and BTZ against the cancer cells (Figure 2A), BTZ alone at 100 nM barely shows cytotoxicity on the cancer cells, 1 at 200 μ M exhibits mild cytotoxicity against HeLa cells, with the cell viability of 79 %, and the combination of 1 (200 μ M) and BTZ (100 nM) decreases the viability of the HeLa cells to 38 %. Similarly, the cell viability of MCF-7 cells is 43 % after being treated by 1 at 200 µM (TEM image containing nanofibers shown in Figure S6), and the co-incubation of 1 and BTZ decreases the cell viability of MCF-7 cells to 29 %. The combination also decreases the cell viability of HepG2 cells from 57 % (treated by 1 alone) to 34 %. Comparing that it significant increases the cytotoxicity of 1 to HeLa cells, BTZ only moderately increases the cytotoxicity of 1 against MCF-7 and HepG2 cells.

To verify whether the decrease of proteolysis results in enhanced cytotoxicity of **1** against the cancer cells, we examine the amount of **1** remained in the cell lysates or in culture medium after the cancer cells are incubated with **1** alone or the mixture of **1** and BTZ for 1 h (Figure 2B). Without the addition of BTZ, most of the hydrogelator **1** inside the cells degrades within 1 hour, with the remaining **1** to be 14 %, 33%, and 39% in HeLa, HepG2, and MCF-7 cells, respectively. This result explains the higher cytotoxicity of **1** on MCF-7 and HepG2 cells since more molecules of **1** present in those two cancer cell lines. The addition of BTZ together with **1** increases the percentages of remaining **1** to be 33%, 44%, and 50% in HeLa, HepG2, and MCF-7 cells, respectively. This result explains the significant increase of cytotoxicity of **1** by combining it with BTZ against HeLa cells. That is, there are more molecules of **1** remaining after the co-incubation of BTZ, which leads to more nanofibers, and thus resulting in higher cytotoxicity.

To elucidate the reason that the hydrogelator **1** shows cytotoxicity to cancer cells and exhibits higher cytotoxicity in the presence of BTZ, we synthesize two control molecules: **2** (Nap-Phe-Phe-Asn) and **3** (Nap-Phe-Phe-Glu), as shown in Scheme 2. Containing arsparagine (N) and the self-assembly motif,^[19] **2** is a part of **1**; and **3** contains a negative

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charged residue (glutamic acid, E) and the self-assembly motif. **2** exhibits little cytotoxicity against HeLa, MCF-7 and HepG2 cells, except starting to inhibit the cells at 500 μ M (Figure 3). Molecule **3** hardly exhibits cytotoxicity against the three cancer cell lines (Figure 3). TEM images of **1**, **2**, and **3** at 500 μ M indicate that the three molecules can form nanofibers at this concentration, which explains the cytotoxicity to cancer cells at 500 μ M (Figure S7). These results suggest that the interaction of the assemblies of **1** with cell surface glycans is important because such interactions likely facilitate the nanofibers enter the cells.

We also co-incubate the molecule 2 and 3 with BTZ to treat the cancer cells, which barely increases the cytotoxicity of 2 and 3 (Figures 4A, 4B, S8, and S9). We incubate the HeLa cells with 2 (or 3) alone or in combination with BTZ for 1 h and examine the molecules remained in cell lysate or culture medium (Figure 4C, 4D), which indicates that both 2 and 3 degrade slower than 1. Intracellular concentrations of 1, 2, and 3 (Table 1) after treating HeLa cells with 1, 2, and 3 (200 μ M, 1 h) show that 1 has a much higher intracellular concentration than the incubation concentration, but 2 (or 3) shows much lower intracellular concentrations. This result agrees with that (i) 1 is able to interact with glycans on cells for cellular uptake and (ii) 1 likely self-assembles inside the cells to reduce the exocytosis of the peptides.

In conclusion, here we present a promising approach to enhance the anticancer activity of nanofibers formed by peptide by co-incubating the peptides with a proteasome inhibitor, BTZ. The overall cytotoxicity of peptide **1** is significantly increased after adding BTZ in the cell culture, since BTZ inhibits proteasome and increases the amount of nanofibers inside cells. The results suggest that the cellular uptake and intracellular fate of the peptides are important factors for designing peptide nanofibers for inhibiting cancer cells. Comparing to the use of D-amino acid to increase the long-term bio-stability of peptide materials,^[20] this approach should be a useful alternative because L-peptides are more likely to exhibit higher affinity to cellular targets than D-peptides do, which may improve the specificity of the nanofibers against cancer cells. Moreover, these results imply that increasing proteolysis may be an effective way to reduce the cytotoxicity of the aggregates of aberrant proteins.

Experimental Section

Materials

The reagents such as N, N-diisopropylethylamine (DIPEA), O-benzotriazole-N,N,N',N'tetramethyluronium-hexafluorophosphate (HBTU), were purchased from ACROS Organics USA. All other amino acid derivatives including Fmoc-Phe-OH, Fmoc-His(Trt)-OH, Fmoc-Asn(Trt)-OH, Fmoc-Pro-OH, Fmoc-Arg(pbf)-OH were purchased from GL Biochem (Shanghai) Ltd. The synthesis of hydrogelator **1** is based on solid phase peptide synthesis (SPPS) and has been reported previously. The crude compound that acquired from SPPS is purified by HPLC and the overall yield of the synthesis is about 80%. ¹H-NMR and LC-MS spectra are used to characterize the product.

Instrumentation and characterization

We used Waters Delta600 HPLC system, which equipped with an XTerra C18 RP column and an in-line diode array UV detector to purify the product. LC-MS machine we used was a Waters Acquity Ultra Performance LC with Waters MICROMASS detector. Hydrogen nuclear magnetic resonance (NMR) spectra were recorded on a Varian Unity Inova 400 with DMSO as solvent. We used Morgagni 268 transmission electron microscope to take Transmission electron microscope (TEM) images. MTT assay for cell cytotoxicity was test on DTX880 Multimode Detector.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Figure 1.

(A, B) TEM images of the hydrogels (inset: optical images) formed by (A) 0.4 wt% **1**; (B) 0.4 wt% **1** and 1 eq. lactose in water at pH7.4 (Scale bar = 100 nm); (C, D) The (C) strain and (D) frequency dependence of dynamic storage modulus G' and loss modulus G'' of the gels formed by **1** at 0.4 wt% or the gel formed by **1** at 0.4 wt% and 1 eq. lactose at pH7.4 in water.



Figure 2.

(A) Cell viability of HeLa, MCF-7 and HepG2 cells incubated with 1 (200 μ M), BTZ (100 nM) or the mixture of 1 and BTZ for 48 h. (B) Percentage of 1 remained in HeLa cell or in culture medium; MCF-7 cell or in culture medium; HepG2 cell or in culture medium after being treated with 200 μ M of 1 alone or together with 100 nM BTZ for 1 h.



Figure 3.

(A–C) Cell viability of (A) HeLa, (B) MCF-7, (C) HepG2 cells incubated with **2**; (D–F) Cell viability of (D) HeLa, (E) MCF-7, (F) HepG2 cells incubated with **3** for 3 days.



Figure 4.

(A, B) Cell viability of HeLa, MCF-7 and HepG2 cells incubated with (A) 2 (200 μ M), BTZ (100 nM) or the mixture of 2 and BTZ; (B) 3 (200 μ M), BTZ (100 nM) or the mixture of 3 and BTZ for 48 h; (C) Percentage of 2 remained in HeLa cell or in culture medium after being treated with 200 μ M of 2 alone or together with 100 nM BTZ for 1 h; (D) Percentage of 3 remained in HeLa cell or in culture medium after being treated with 200 μ M of 3 alone or together with 100 nM BTZ for 1 h; (D) Percentage or together with 100 nM BTZ for 1 h.



Scheme 1.

Chemical structure of a hydrogelator and the proposed mechanism for its nanofibers to inhibit cancer cells.



Scheme 2. Chemical structures of molecule 2 and 3.

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Table 1

Intracellular concentrations of the hydrogelators in HeLa cells.

Intracellular concentration (µM) ^[a]	Average	st.dev
1	296.8	3.61
2	23.5	0.82
3	36.4	1.01

[a] The cell lysates of HeLa cells are collected after 1h incubation with 200 μ M of hydrogelators at 37 °C.