Evaluation of novel cyclic analogues of apelin

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Abstract. Apelin regulates various cell signaling processes through interaction with its specific cell-surface receptor, APJ, which is a member of a seven transmembrane G protein-coupled receptor superfamily. To develop a novel apelin analogue, we synthesized cyclic analogues of minimal apelin fragment RPRLSHKGPMF (apelin-12), and evaluated their bioactivities in a recombinant human APJ-expressed cell line. Three cyclic analogues were synthesized: cyclo apelin-12 (C1) in combination with amino-terminal to carboxy-terminal, cycloureap apelin-12 (C3) in combination with amino-terminal and amino acid side chain at positions 7, and cyclic apelin-12 (C4) in combination with amino acid side chain at positions 7 to carboxy-terminal. All cyclic analogues exhibited dose-dependent inhibitory effects against forskolin-induced cyclic adenosine monophosphate (cAMP) accumulation, and the maximal effects were almost abolished by pertussis toxin (PTx) treatment. Moreover, they could modulate the intracellular signaling pathways composed of Akt and extracellular signal-regulated kinase 1/2 (ERK1/2) serine/threonine protein kinases in PTx-sensitive manner. This is the first approach to apply cyclization on apelin, and these results provide the basis for the development of drug-like apelin analogues.

Introduction

Apelin is an endogenous ligand peptide of APJ (a putative receptor protein related to the angiotensin-type 1 receptor, AT1) (1), which is a member of a seven transmembrane G protein-coupled receptor superfamily. Preproapelin synthesized as a 77 amino acid polypeptide is cleaved to multiple short forms of different sizes. Peptides of 12, 13, 17 and 36 amino acids in length including C-terminal portion are known to be biologically active (2,3). The APJ has a 31% amino acid sequence homology with the AT1, but does not display specific binding to angiotensin II, which is the ligand of AT1 and plays a central role in blood pressure and water and electrolyte homeostasis. Both apelin and APJ are expressed in a wide range of tissues, such as the cardiovascular and the central nervous system (4,5). Exogenous apelin administration alters cardiovascular function, blood pressure, body temperature, body fluid and behaviors involved in food intake and water intake (4,5). In addition, APJ was also identified as a human immunodeficiency virus type I (HIV-1) coreceptor, and apelin was shown to block HIV-1 entry through APJ (3,6).

Currently, there are growing lines of evidence obtained from clinical studies that several human cardiovascular diseases are accompanied by changes in apelin and/or APJ expression in cardiovascular tissues (5). In animal studies, it was shown that apelin exerts vasodilation and positive inotropic actions ex vivo and in vivo (7-10). Apelin activates endothelial nitric oxide synthase (eNOS) and consequently stimulates nitric oxide (NO) release from the vascular endothelial cells, indicating the possible protective effect of apelin-APJ system on cardiovascular diseases through NO-dependent mechanism (9). In addition, the administration of apelin also ameliorated prognosis for heart function under pathological condition, such as experimental ischemia and reperfusion injury (11-14). Thus, it has been implicated that apelin plays an important role in the maintenance of cardiovascular homeostasis, and APJ has a potential to be a novel therapeutic target of cardiovascular dysfunction (4,5). However, there are no available chemical small compounds for APJ.

The cyclization can be applied to a variety of peptides to improve their agonist or antagonist potency, including bioactivity, selectivity and bioavailability (15). We therefore made an attempt to synthesize cyclic apelin analogues based on apelin-12 fragment, which was identified as a minimum unit ligand for APJ, and evaluated their bioactivities in cell-based assays. APJ was demonstrated to exhibit apelin-dependent inhibition of adenylyl cyclase via PTx-sensitive G, proteins, when expressed in recombinant cell lines (16-18). Moreover, in these cells, it was also shown that apelin induced the activation of the PI3K/Akt and MEK1/ERK1/2 signaling pathways. In the present study, we validated the potency of cyclic analogues of apelin by using these cellular signal indicators in vitro.
Materials and methods

Peptides and reagents. Peptides were synthesized by Peptide Institute, Inc. (Osaka, Japan). The sequence data of synthetic peptide are shown below: pE-apelin-13: pGlut-Arg-Pro-Arg-Leu-Ser-His-Lys-Gly-Pro-Met-Pro-Phe; cyclo apelin-12 (1-12): cyclo [Arg-Pro-Arg-Leu-Ser-His-Lys-Gly-Pro-Met-Pro-Phe] (1-12); cycloleucine apelin-12 (1-7): cycloleucine [Arg-Pro-Arg-Leu-Ser-His-Lys-Gly-Pro-Met-Pro-Phe] (1-7); cyclo apelin-12 (7-12): cyclo [Arg-Pro-Arg-Leu-Ser-His-Lys-Gly-Pro-Met-Pro-Phe] (7-12). The characters used for the amino acids are the standard IUPAC. ‘pE’ and ‘pGlu’ represent a pyroglutamic acid residue. Cyclo apelin-12 (1-12), cycloleucine apelin-12 (1-7) and cyclo apelin-12 (7-12) are referred to as simply C1, C3 and C4, respectively. The number in the brackets indicates positions of amino acids of which terminal-group and side chain are covalently bound. 

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Establishment of stable cell lines. HEK293-T cells were grown in growth medium: Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% heat-inactivated fetal bovine serum (FBS), non-essential amino acids, 100 U/ml penicillin and 100 μg/ml streptomycin at 37°C (5% CO2 and 95% humidity). Cells were transfected with the coding region of the human APJ gene expressed pcDNA3.1/Zeo(+) (Invitrogen) using TransFast (Promega). Stable transformants (so-called APJ/293-T) were selected and propagated with 200 ng/ml zeocin (Invitrogen). To confirm the protein expression of APJ, cells were solubilized in Laemmli's sample buffer and subjected to immunoblot analysis (see below). HEK293-T cells stably transfected with pcDNA3.1/Zeo(+) (so-called mock/293-T) were used to verify non-specific action of synthetic analogues.

CAMP assay. Cells were plated at 20,000 cells per well into type I collagen-coated 96-multiwell plates and allowed to adhere overnight in 90 μl of growth medium supplemented with 20 mM HEPES buffer at 37°C (5% CO2 and 95% humidity). For PTx treatment, cells were incubated with 100 ng/ml of PTx for 12-15 h before assay. On the day of assay, the cell cultured plates were equilibrated at room temperature for 20 min and then 5 μl of aliquots of 1.95 mM Ro-20-1724 were added to the cultured medium at final concentration of 100 μM and incubated at room temperature for another 20 min. After incubation, 5 μl of aliquots of different doses of peptide analogues containing 0.1 mM forskolin were loaded 5% of the final volume (forskolin; at final concentration of 5 μM) and incubated at room temperature for 10 min. After stimulation, the reaction was stopped by adding of 5-fold diluted lysis buffer 1A, a component of the cAMP enzyme-immunoassay (EIA) system (Amersham Biosciences), and then, 100 μl aliquots of cell lysate served as cAMP measurement according to the manufacturer's instructions. The sample absorbance was measured on Wallac ARVOX Multilabel Counter (Perkin-Elmer, Tokyo, Japan) and analyzed using WorkOut Ver. 1.5 software (Perkin-Elmer).

Immunoblot analysis. Cells were plated at 150,000 cells per well into poly-d-lysine-coated 24-multiwell plates and allowed to adhere overnight in 90 μl of growth medium at 37°C (5% CO2 and 95% humidity). In the next assay, the cultured medium was replaced to 495 μl of serum-free DMEM and the cells were further incubated for 24 h. For PTx treatment, cells were incubated with 100 ng/ml of PTx for 12 h before assay. On the day of assay, 5 μl aliquots of peptide analogues were added to the cultured medium and incubated at 37°C. After stimulation, the plates were left to stand in ice water for a few min, and then, the medium was removed. The cells were washed with ice-cold PBS, followed by immediate solubilization with 100 μl of Laemmlli's sample buffer containing 100 mM dithiothreitol. After sonication and treatment at 99°C for 3 min, the cell lysates were centrifuged at 4°C for 5 min and the super-natants were subjected to SDS-PAGE (10% polyacrylamide) and transferred to a nitrocellulose membrane (Protoran; Schleicher&Suell, Dassel, Germany). Filters were blocked with skim milk and then blotted with primary antibodies. After incubation, blots were incubated with the corresponding secondary antibodies, followed by visualization via chemiluminescent detection using Immobilon Western HRP Substrate (Millipore Corp., Billerica, MA, USA), and data images were captured with LAS-3000 (Fujifilm, Tokyo, Japan).

Data analysis. Duplicates or triplicates were performed and at least three independent trials with similar results were done for each set of experiments. Non-linear regression analysis of dose-response data was performed using Prism4 (GraphPAD Software, San Diego, CA, USA). One-way ANOVA and Bonferroni's multiple comparison test with 95% confidence were adapted for the evaluation of the significance of the differences between groups. The data are presented as means ± standard error (SE) of at least three independent experiments.

Results

We synthesized three cyclic apelin analogues using apelin-12 as a template (Fig. 1). The structure of synthetic peptides...
were followed: cyclo apelin-12 (C1) in combination with amino-terminal to carboxy-terminal, cyclourea apelin-12 (C3) in combination with amino-terminal and amino acid side chain at positions 7, and cyclic apelin-12 (C4) in combination with amino acid side chain at positions 7 to carboxy-terminal. The amino-terminal pyroglutamylated apelin-13 (pE-apelin-13) was also synthesized for use as a reference drug for cell-based assays.

To evaluate the bioactivities of the synthetic analogues in cell-based assays, we established recombinant cell lines and assessed a specific stable expression of APJ in HEK293-T cell transfected with human APJ gene. Immunoreactive band of the APJ (~42 kDa) was predominantly detected in the APJ/293-T cell line by immunoblotting using an anti-APJ antibody, while not in the mock/293-T cell line (Fig. 2A). First, we performed the cAMP assay using the APJ/293-T cell line to evaluate the bioactivities of the synthetic analogues for cAMP accumulation. (A) APJ stable expression in APJ/293-T cells was determined by immunoblotting using anti-APJ antibody. Upper panel, immunoreactive band of APJ is detected in APJ/293-T (right lane) but not in mock/293-T (left lane) cells. Bottom panel, immunoreactive band of β-actin in APJ/293-T (right lane) is comparable to that observed in mock/293-T (left lane) cells. Molecular weight (MW) markers are indicated in kilodaltons (kDa) on the left of panel. (B) Dose-response curves of forskolin-induced cAMP accumulation in APJ/293-T cells for apelin and cyclic analogues. (C) Effects of apelin (100 nM) and cyclic analogues (10 μM) on cAMP accumulation in APJ/293-T cells were almost abolished by PTx treatment. (D) No significant effects of apelin (100 nM) and cyclic analogues (10 μM) on cAMP accumulation were observed in mock/293-T cells. The X axis is the negative logarithm of peptide concentration and the Y axis is the % of cAMP accumulation of forskolin control. Values are expressed as means ± SE of at least three independent experiments.

<table>
<thead>
<tr>
<th>pE-apelin-13</th>
<th>pGlut-Arg-Pro-Arg-Leu-Ser-His-Lys-Gly-Pro-Met-Pro-Phe</th>
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<tr>
<td>apelin-12</td>
<td>Arg-Pro-Arg-Leu-Ser-His-Lys-Gly-Pro-Met-Pro-Phe</td>
</tr>
<tr>
<td>C1</td>
<td>[Arg-Pro-Arg-Leu-Ser-His-Lys-Gly-Pro-Met-Pro-Phe]</td>
</tr>
<tr>
<td>C3</td>
<td>[Arg-Pro-Arg-Leu-Ser-His-Lys-Gly-Pro-Met-Pro-Phe]</td>
</tr>
<tr>
<td>C4</td>
<td>[Arg-Pro-Arg-Leu-Ser-His-Lys-Gly-Pro-Met-Pro-Phe]</td>
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</table>

Figure 1. Structure of synthesized apelin and cyclic analogues. The combined amino acids in cyclic analogues are represented by line in the alignment of the sequences of apelin-12.
cell line, when exposed to different doses of peptide analogues, and determined the relative potency (Fig. 2B and Table I). All synthetic peptides showed dose-dependent inhibitory effects on forskolin-induced cAMP accumulation in the APJ/293-T cells. The effective concentration ranges of every cyclic analogue shifted to higher concentration than that of pE-apelin. The EC50 (median effective concentration) value of pE-apelin-13 was 0.38±0.04 nM, in quantitative agreement with previous reports (19), yielding the maximal inhibition of ~10% at 100 nM. The EC50 value of C1 was 493.17±88.76 nM (relative 1,275-fold increase to pE-apelin-13), yielding the maximal inhibition of ~20% at 10 μM. The EC50 values of C3 and C4 were 144.27±5.25 nM (relative 373-fold increase to pE-apelin-13) and 78.62±14.88 nM (relative 203-fold increase to pE-apelin-13), respectively, yielding the similar maximal inhibition of ~10% at 10 μM in an experimental range of concentration of peptides. To determine whether the effects of pE-apelin-13 and cyclic analogues were specific for APJ, we subsequently investigated the PTx-sensitivity and the non-specific action in the mock/293-T cells. The inhibitory effects of pE-apelin-13 (at 100 nM) and three cyclic analogues (at 10 μM, each) on forskolin-induced cAMP accumulation were almost abolished in APJ/293-T cells with PTx pre-treatment (Fig. 2C). No significant inhibitory effects of any peptides, at the same concentration as mentioned above, were observed in the mock/293-T cells (Fig. 2D). Taken together, we showed that all cyclic analogues were API agonists.

Secondly, to assess whether cyclic analogues could modulate the intracellular signaling pathways, we examined the activation of Akt and ERK1/2 molecules by the immunoblot analysis using antibodies against phosphorylated Akt (Ser473) and ERK1/2 (Thr202/Tyr204), which are known to reflect its activation. In the APJ/293-T cells, the phosphorylation levels of both Akt and ERK1/2 were rapidly increased within 2 min after exposure to pE-apelin-13 (at 1 μM) (Fig. 3A). The enhanced phosphorylation level of Akt was sustained during the experimental period (0-30 min), whereas that of ERK1/2, by contrast, was transient during the short period (2-5 min) and subsequently depressed close to the basal level after 15 min. Exposure to the cyclic analogues (at 1 μM) also rapidly induced both Akt and ERK1/2 phosphorylation and showed similar time-course manner to pE-apelin-13. Though a slight increment of the phosphorylation levels of ERK1/2 compared with exposure to pE-apelin-13, all cyclic analogues increased the phosphorylation levels of Akt to a similar extent compared with pE-apelin-13 after exposure. We also tested the dose-dependent effects of pE-apelin-13 and cyclic analogues on the activation of these molecules. The APJ/293-T cells were exposed to different doses of peptides in a concentration range from 1 nM to 1 μM for 5 min, followed by immunoblot analysis. As shown in Fig. 3B, the phosphorylation level

<table>
<thead>
<tr>
<th>Peptide</th>
<th>EC50 (nM)</th>
<th>Relative EC50</th>
<th>Maximal inhibition (%)</th>
</tr>
</thead>
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<tr>
<td>pE-apelin-13</td>
<td>0.38±0.04</td>
<td>1</td>
<td>10.0±1.2</td>
</tr>
<tr>
<td>C1</td>
<td>493.17±88.76</td>
<td>1275</td>
<td>19.3±2.1</td>
</tr>
<tr>
<td>C3</td>
<td>144.27±5.25</td>
<td>373</td>
<td>13.1±2.0</td>
</tr>
<tr>
<td>C4</td>
<td>78.62±14.88</td>
<td>203</td>
<td>11.3±1.3</td>
</tr>
</tbody>
</table>

Table I. Activity profile of synthetic peptides from inhibitory effects on cAMP accumulation.
of Akt was not changed at 1 nM pE-apelin-13 compared with none and vehicle, but was increased to a similar extent in a concentration range from 10 nM to 1 μM. In contrast, the phosphorylation levels of ERK1/2 were increased in a pE-apelin-13 dose-dependent manner. Exposure to the cyclic analogues at high concentrations (C1; 1 μM, C3 or C4; from 100 nM to 1 μM) induced the increments of the phosphorylation levels of both Akt and ERK1/2 in a dose-dependent manner, whereas at low concentrations, they did not increase. The induced phosphorylation of both Akt and ERK1/2 in APJ/293-T cells was almost abolished with PTx pretreatment (Fig. 4A), and no effects of any peptide (at 1 μM) were observed in the mock/293-T cells (Fig. 4B). These signaling assays demonstrated the potency of all cyclic analogues to activate both Akt and ERK1/2 in APJ-dependent and PTx-sensitive manner.

Discussion

APJ is thought to be a novel therapeutic target of cardiovascular dysfunction. The ameliorative effectiveness of apelin administration was demonstrated in pathological animal models, however, the effective dose was too high (12). Moreover, it is evident that exogenous apelin might be unstable and rapidly hydrolyzed by various peptidases including ACE2 (20). Therefore, it is necessary to modify apelin fragment to raise the efficacy as a drug and its physiological stability.

Several investigators have reported the structure-activity relationship (SAR) analysis of linear apelin analogues at human APJ. Medhurst et al. (19) reported that ‘RPRL’ (Arg-Pro-Arg-Leu) motif in the amino-terminus of apelin-13 was an important motif for binding to APJ. On the other hand, Fan et al. (21) demonstrated that the positive charge and hydrophobic residues of apelin-13, such as Arg and Pro, were very important for its functional potency and binding affinity. They also observed the secondary structure of apelin-36 and apelin-13 using Circular dichroism spectroscopy. Both peptides were shown to take unordered structure in aqueous solution. Despite these detailed analyses, there is no available pharmacophore model of apelin at APJ, because of its high dynamic conformational diversity. In this situation, one might expect that the cyclization, another SAR analysis method, is one of the most useful approaches to design drug-like apelin analogues.

Cyclization applies conformational restriction to a peptide, resulting in the improvement of its stability and bioactivity. There are several successful applications of cyclization to a peptic ligand. For example, the bioactive cyclic analogues of somatostatin-derived peptides were synthesized (22,23). The cyclic hexapeptide of somatostatin was shown to contain pharmacophore for somatostatin type 2 receptor (sst2), allowing the discovery of highly active pharmacological agents (15). It was also reported that the N-terminal cyclic motilin partial peptides were identified as neutral motilin agonist (24), and that the bioactivity of the minimal histogranin tetrapeptide for dopamine D3 receptor was potentiated by amino acid replacement and cyclization (25). These studies showed that the cyclization could fix a peptide ligand in rigid conformation so as to activate or inactivate its specific receptor. Subsequent active core exploration will contribute to a pharmacophore search from non-peptidic small compound library. Furthermore, bioactive peptic analogue will provide a therapeutic importance for the target receptor at an early stage in the drug development process. However, there is no report to apply cyclization to apelin yet.

In this study, we attempted to generate the cyclic apelin analogues. Both apelin-12 and pE-apelin-13 are thought to be the minimal active units of apelin. The amino-terminal pyroglutamyl residue in pE-apelin-13 is considered to contribute to its stability, but is not essential for its functional potency and binding at APJ (3,26). Therefore, we selected the apelin-12 as a template for cyclic analogues, and assessed the effect of cyclization on apelin-12. We successfully synthesized three cyclic analogues of apelin-12 fragment and subsequently evaluated their bioactivities in a recombinant human APJ-expressed cell line. In our cell-based assays, we used pE-apelin-13 as a reference drug because of its well-established bioactivity in vitro assays (16-19,21,26). From the cAMP assay, we identified the functional agonistic activities of all
cyclic apelin analogues. Importantly, these analogues could induce the activation of Akt and ERK1/2 in PTX-sensitive manner.

The Akt and MAPK (mitogen-activated protein kinase) family are key molecules that mediate G protein-coupled receptor-dependent signal transduction involved in the control of cell cycle progression, proliferation, migration, apoptosis and survival (27,28). In various cell lines and primary neurons, it was shown that apelin-dependent activation of PI3K/Akt and MAPK signaling pathways resulted in cell proliferation, migration and survival (17,18,29,30). Moreover, apelin was shown to produce direct cardioprotective actions against myocardial ischemia-reperfusion injury of mouse via the PI3K/Akt and MEK1/ERK1/2 signaling pathways (12). In this study, the activation of Akt and ERK1/2 induced by cyclic analogues suggested the potential activities of cyclic apelin analogues as a therapeutic agent in cardiovascular diseases as well as endogenous apelin.

In conclusion, by using cyclization methods, we established novel cyclic apelin analogues as the effective agonist for APJ. These cyclic apelin analogues may be useful for early validation of APJ and its target disease in a drug discovery. For this purpose, further evaluation will be needed on the stability and binding affinity of these peptides. The characterization of cyclic apelin analogues may provide insight into APJ-related cardiovascular diseases.

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