

such as interneurons, astrocytes and/or microglia that also express CXCR4 receptors (Tanabe *et al.*, 1997; Klein *et al.*, 1999). A hypothetical model of gp120-induced enhancement of GDP frequency is shown in Fig. 7.

GDPs are network-driven membrane oscillations characterized by recurrent membrane depolarization with superimposed fast action potentials. They are large synaptic events recorded in rat hippocampal slices between postnatal days P0 to P10 (Ben-Ari *et al.*, 1989). GDPs are synchronized locally within a temporal frame of several hundreds of milliseconds and recur at a frequency of approximately 0.1 Hz (Ben-Ari, 2001; Kasyanov *et al.*, 2004). They constitute the first synaptic pattern observed in the connections that are established around the time of birth with an extensive growth at the second postnatal week (Gaiarsa *et al.*, 1992; Gomez-Di Cesare *et al.*, 1997). During this period of intensive growth, GDPs provide most of the ongoing activity in pyramidal neurons and interneurons (Khazipov *et al.*, 1997). Because GDPs resulting from the excitatory effects of GABA can increase the levels of $[Ca^{2+}]_i$ through voltage-gated Ca^{2+} channels and NMDA receptor channels (Leinekugel *et al.*, 1995; Leinekugel *et al.*, 1997) and because application of gp120 increases the levels of $[Ca^{2+}]_i$, the persistent increase of GDPs mediated by gp120 may cause neuronal damage as a result of excessive increase of $[Ca^{2+}]_i$, leading to retardation of brain development and deterioration in cognitive function. Interestingly, hippocampal-dependent integrative functions – learning and memory – emerge later at the end of the first postnatal

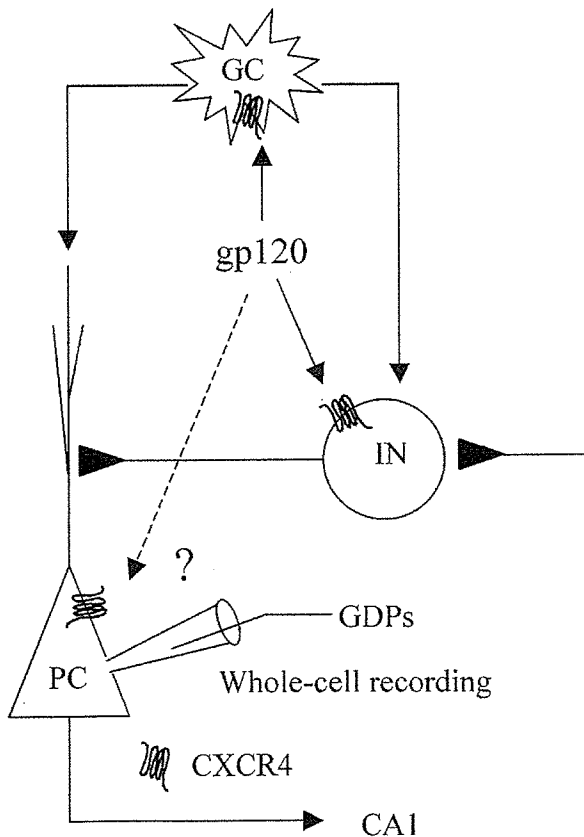


FIG. 7. A hypothetical model of gp120-induced enhancement of GDP frequency. Gp120 may act on CXCR4 receptors expressed on interneurons (IN) and increase GABA release. Gp120 may also act on glial cell (GC) CXCR4 receptors and enhance glial cell secretion of cytokines/chemokines and other bioactive molecules. These bioactive substances may target interneurons and/or CA3 pyramidal cells (PC) and up-regulate interneuron secretion of GABA and/or enhance the sensitivity of CA3 pyramidal cells to GABA, or both.

month (Altman *et al.*, 1973; Rudy & Stadler-Morris, 1987), suggesting that GDPs occur at an early period during which the hippocampus does not perform the integrative functions. Thus, the persistent increase of GDP frequency by gp120 during the first postnatal week may disturb the subsequent synaptogenesis and hippocampal integrative functions such as learning and memory (Ben-Ari, 2001), leading to cognitive decline as seen in paediatric AIDS patients with HE.

In the immature brain, transient elevations of $[Ca^{2+}]_i$ are important in many aspects of development at the network level. These include proliferation (LoTurco *et al.*, 1995), neuronal migration (Komuro & Rakic, 1996), and motility of axonal and dendritic growth cones (Kater *et al.*, 1988). Many of these Ca^{2+} -dependent processes require repeated, periodic changes of $[Ca^{2+}]_i$. As the GDPs are recurrent synaptic events that involve large populations of neurons (Garaschuk *et al.*, 1998; Garaschuk *et al.*, 2000), they may represent a prime candidate to promote neuronal development and to 'synchronize' development over large neuronal populations. However, a sustained elevation of $[Ca^{2+}]_i$ may cause neural cell damage. It has been shown in rat primary hippocampal neuronal cultures that gp120 produced a delayed onset, long-lasting intracellular Ca^{2+} oscillation (Lo *et al.*, 1992) and elevated intracellular free Ca^{2+} levels (Lo *et al.*, 1992; Pandey & Bolsover, 2000). We have also observed that gp120 produced a sustained increase of $[Ca^{2+}]_i$ in primary rat hippocampal-cortical neuronal cultures (Fig. 3). Consistent with these previous observations, we found that gp120 produced a delayed onset in the increase of GDP frequency. As membrane depolarization during GDPs increases $[Ca^{2+}]_i$ through the activation of voltage-dependent Ca^{2+} channels in interneurons and pyramidal cells in the hippocampus (Leinekugel *et al.*, 1995), the long-lasting enhancement of GDP frequency induced by gp120 may result in excessive increase of $[Ca^{2+}]_i$, leading to neuronal dysfunction and/or death. Thus, elevation of $[Ca^{2+}]_i$ in the neurons as a result of gp120-induced increase of GDP occurrence may play an important role in the pathogenesis of HIV-1-associated neurological symptoms seen in infants and children.

Chemokine receptor CXCR4, a coreceptor for HIV-1 entry into the target cells, is expressed in human and rat hippocampal glial and neuronal cells (Lavi *et al.*, 1997; Meucci *et al.*, 1998; Banisadr *et al.*, 2002). The functional significance of CXCR4 expression in hippocampal neurons is not fully understood. It has been shown that the brains of mouse embryos without CXCR4 receptors have a severe abnormality in the development of cerebellar morphology (Ma *et al.*, 1998; Zou *et al.*, 1998). Activation of neuronal CXCR4 receptors by a physiological agonist, SDF-1 α , modulates synaptic transmission in developing rat cerebellum (Limatola *et al.*, 2000). In contrast to its physiological role, CXCR4 has been shown to play a key role in mediating gp120 cytotoxicity in both immune cells (Herbein *et al.*, 1998) and neurons (Meucci *et al.*, 2000), even in the absence of the cell surface glycoprotein CD4 (Hesselgesser *et al.*, 1998). Activation of CXCR4 triggers a transient increase of $[Ca^{2+}]_i$ (Zheng *et al.*, 1999), which is believed to participate in gp120 neurotoxicity (Haughey & Mattson, 2002). Thus, the CXCR4 has been speculated as the primary target of gp120 in mediating neuronal toxicity (Bachis & Mocchetti, 2004). We found that bath application of T140, a CXCR4 receptor antagonist, blocked gp120-induced enhancement of GDP firing frequency. This blockade indicates the involvement of the chemokine receptor, CXCR4, in gp120 enhancement of GDP firing. This finding is further supported by experimental results that bath application of the only CXCR4 receptor agonist, SDF-1 α , mimicked the effects of gp120 on GDPs. It is worth pointing out that SDF-1 α decreased GDP frequency during the bath perfusion period, an effect that was not observed during bath application of gp120. The potential

mechanism(s) underlying this difference remains to be determined. It may result that SDF-1 α and gp120 engage different mechanisms in stimulating and/or regulating GABA release following their binding to G-protein-coupled CXCR4 receptors expressed on interneurons, glial cells and/or CA3 pyramidal cells. It may also be the consequence that the glial cells, stimulated by gp120 or SDF-1 α , release a different spectrum of cytokines and/or chemokines that influence GABA-mediated synaptic events or change the feature of 'pacemaker' cells in the immature hippocampus, or both.

Several groups have reported the effects of HIV-1 gp120 on signal transduction pathways in the CNS (Zorn *et al.*, 1990; Schneider-Schaulies *et al.*, 1992; Wyss-Coray *et al.*, 1996). Gp120 activates PKC (Zorn *et al.*, 1990) and induces PKC translocation differentially in rat primary neuronal cultures (Ushijima *et al.*, 1993). Based on these observations, we reasoned that gp120 may enhance GDP firing through PKC. Our results showed that gp120-induced enhancement of GDP occurrence was blocked by a PKC antagonist, H7, demonstrating the involvement of PKC in the gp120-induced increase of GDPs. In addition, we found that the adenylyl cyclase stimulator forskolin mimicked the effects of gp120 on GDPs, suggesting the possible involvement of cAMP-dependent kinase in gp120-associated signaling as well. Taken together, our results revealed that HIV-1 gp120 enhances GDPs via the chemokine receptor, CXCR4, in neonatal rat hippocampal slices and this enhancement was mediated through PKA/PKC-dependent pathway.

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Abbreviations

EPSPs, excitatory postsynaptic potentials; GDPs, giant depolarizing potentials; HAD, HIV-1-associated dementia; HE, HIV-1-associated encephalopathy; HIV-1, human immunodeficiency virus type one; MPs, mononuclear phagocytes; PKA, cAMP-dependent protein kinase; PKC, protein kinase C.

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Development of a linear type of low molecular weight CXCR4 antagonists based on T140 analogs†

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A linear type of several low molecular weight CXCR4 antagonists were developed based on T140 analogs, which were previously found to be strong CXCR4 antagonists that block X4-HIV-1 entry and have inhibitory activities against cancer metastasis/progression and rheumatoid arthritis.

Introduction

A system of a chemokine receptor, CXCR4, and its endogenous ligand, stromal cell-derived factor-1 (SDF-1/CXCL12), has multiple important functions in normal physiology involving the migration of progenitors during embryologic development of the cardiovascular, hemopoietic and central nervous systems.¹ The CXCL12/CXCR4 system has been also recognized to be involved in several pathologic conditions, such as HIV infection,² cancer metastasis/progression³ and rheumatoid arthritis (RA).⁴ First, CXCR4 was identified as a co-receptor that is used in the entry of T cell line-tropic (X4-) HIV-1 into T cells.² Second, it is found that the CXCL12/CXCR4 system is involved in the metastasis of several types of cancers, including breast cancer, pancreatic cancer, melanoma, prostate cancer, kidney cancer, neuroblastoma, non-Hodgkin's lymphoma, lung cancer, ovarian cancer, multiple myeloma, chronic lymphocytic leukemia, acute lymphoblastic leukemia and malignant brain tumor,³ and that this system might determine the metastatic destination of tumor cells. For instance, Müller *et al.* reported that CXCR4 is highly expressed in human breast cancer cells, while CXCL12 is highly expressed in lymph nodes, bone marrow, lung and liver, which represent the primary metastatic destinations of breast cancer, and that breast cancer metastasis can be significantly inhibited by neutralization using anti-CXCR4 antibodies in mice.^{3a} Third,

Nanki *et al.* reported that the memory T cells highly express CXCR4 and the concentration of CXCL12 is extremely high in the synovium of RA patients, and that CXCL12 stimulates migration of the memory T cells and inhibits T cell apoptosis followed by T cell accumulation in the RA synovium.^{4a} Taken together, CXCR4 is thought to represent an important therapeutic target.⁵ Thus, several antagonists directed against CXCR4 have been developed. We previously found a 14-mer peptide, T140, which specifically antagonizes CXCR4,⁶ and that Arg², L-3-(2-naphthyl)alanine (Nal)³, Tyr⁵ and Arg¹⁴ constitute the biologically critical residues of T140 (Fig. 1).⁷ Recently, its potent analogs, 4F-benzoyl-TN14003 and 4F-benzoyl-TE14011, possessing increased stability in serum and liver homogenate, were developed by introduction of a *p*-fluorobenzoyl group, which was defined as a new pharmacophore, into the *N*-terminus.⁸ 4F-benzoyl-TN14003 and 4F-benzoyl-TE14011 showed strong anti-HIV activity *in vitro*, anti-metastatic activity against breast cancer^{3b} and melanoma^{3c} and anti-RA activity in experimental model mice.^{4b} Furthermore, T140-related analogs exhibited significant inhibition against CXCL12-induced migration/activation/invasion of small-cell lung cancer cells,^{3b} acute lymphoblastic leukemia cells^{3c} and pancreatic cancer cells^{3c,d} *in vitro*. Molecular-size reduction of T140 based on the above four critical residues (Arg × 2, Nal and Tyr) led to discovery of a low molecular weight CXCR4 antagonist with a cyclic pentapeptide template, FC131.⁹ In this paper, identification of the enhanced pharmacophore involving an electron-deficient aromatic ring at the *N*-terminus of 4F-benzoyl-TN14003 and 4F-benzoyl-TE14011, such as a *p*-fluorobenzoyl or *p*-trifluoromethylbenzoyl moiety, prompted us to develop novel linear-type low molecular weight CXCR4 antagonists. By combining substructure units of

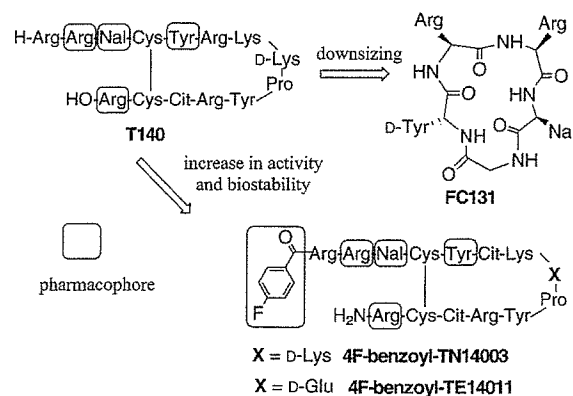


Fig. 1 Development of bio-stable CXCR4 antagonists, 4F-benzoyl-TN14003 and 4F-benzoyl-TE14011, and a downsized antagonist, FC131. Nal = L-3-(2-naphthyl)alanine, Cit = L-citrulline.

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the above four critical residues (Arg × 2, Nal and Tyr) that were used in the development of FC131, in addition to the above

electron-deficient aromatic ring, several compounds were designed and synthesized.

Biological results and discussion

Biological activities of the present synthetic compounds were evaluated by two assays: the 3-(4, 5-dimethylthiazol-2-yl)-2, 5-diphenyltetrazolium bromide (MTT) assay based on the inhibition of X4-HIV-1 (HIV-1_{IIIB})-induced cytopathogenicity in MT-4 cells by test compounds (anti-HIV activity)¹⁰ and a blocking assay based on displacement of CXCL12 binding to CXCR4 by test compounds (binding affinity for CXCR4).¹¹ Initially, three tripeptide mimetics containing amide bonds and/or reduced amide bonds, 1–3 were designed based on the sequence of Arg¹–Arg²–Nal³ in the *N*-terminal region of T140 (Fig. 1 and 2) and synthesized using solution-phase techniques involving amide bond-forming condensation and reductive amination reactions. In this study, (*S*)-(-)-1-(1-naphthyl)ethylamide, which was used in another CXCR4 antagonist KRH-1636,¹² was introduced with the view to enhancement of biostability. Compounds 2 and 3 showed significant anti-HIV activity, while compound 1 did not exhibit activity until the 100 μM concentration, suggesting that a reduced amide bond possessing the conformational flexibility might be more suitable for the interaction of CXCR4 (Table 1).

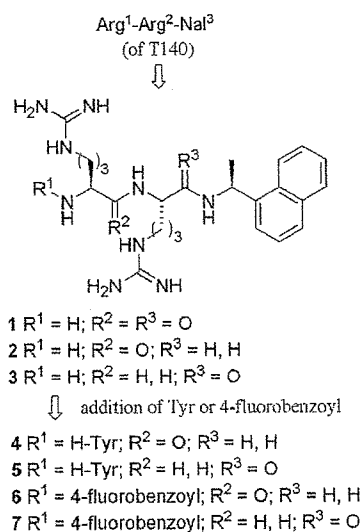


Fig. 2 Development of tri- and tetrapeptide mimetics with CXCR4-antagonistic activity.

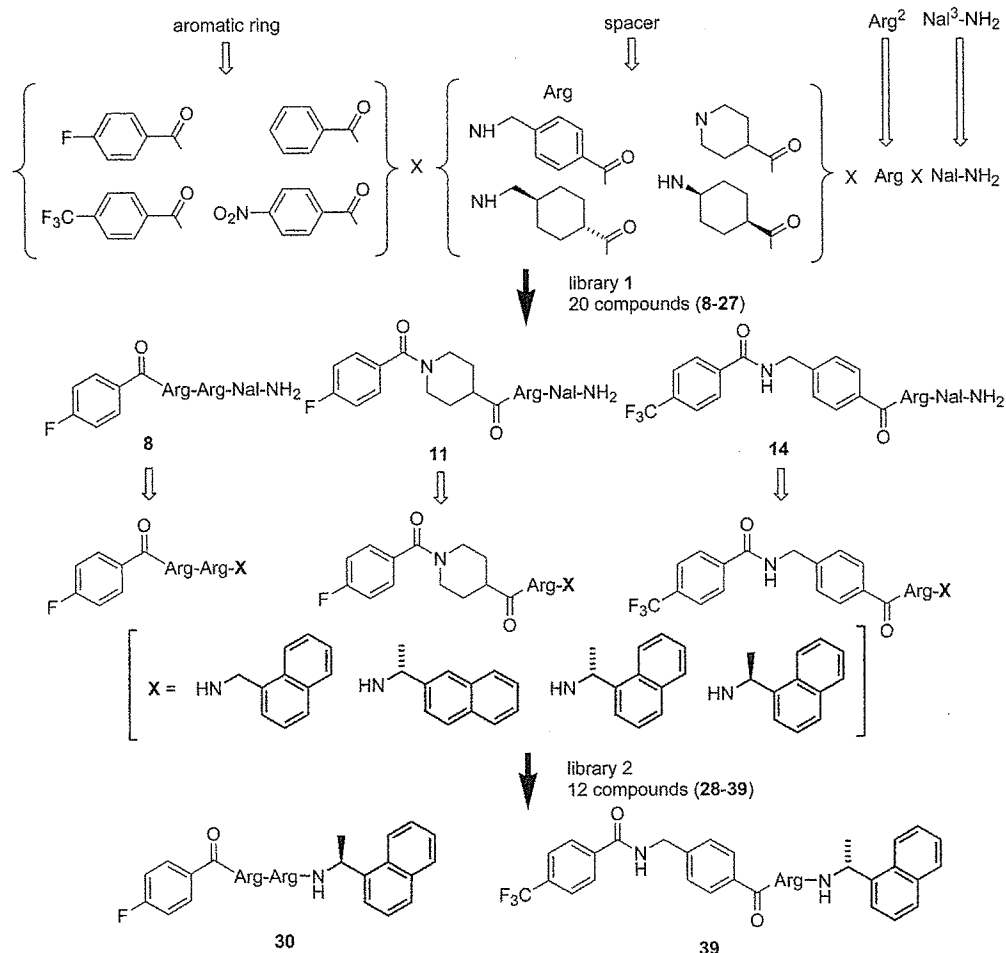


Fig. 3 Design of tripeptide library containing three pharmacophores of the *N*-terminal region of 4F-benzoyl-TN14003 and 4F-benzoyl-TE14011 (aromatic ring, Arg² and Nal³) and the development of new leads.

Table 1 Cytotoxicity, anti-HIV activity and inhibitory activity against CXCL12 binding to CXCR4 of the synthetic compounds

| Compound | CC ₅₀ /μM ^a | EC ₅₀ /μM ^b | IC ₅₀ /μM ^c |
|----------|-----------------------------------|-----------------------------------|-----------------------------------|
| 1 | >100 | >100 | 0.32–1 |
| 2 | >100 | 52 | 0.32–1 |
| 3 | >100 | 46 | 0.32–1 |
| 4 | >100 | 22 | 0.090 |
| 5 | >100 | 26 | 0.30 |
| 6 | >100 | 11 | 0.32–1 |
| 7 | >100 | 1.7 | >1 |
| 8 | >100 | 45 | 0.30 |
| 11 | >100 | 7.7 | >1 |
| 14 | >100 | 6.0 | >1 |
| 30 | >100 | 61 | >1 |
| 39 | 66 | 7.4 | >1 |
| FC131 | >100 | 0.073 | 0.0032 |
| T140 | >10 | 0.026 | 0.0045 |
| AZT | >100 | 0.014 | |

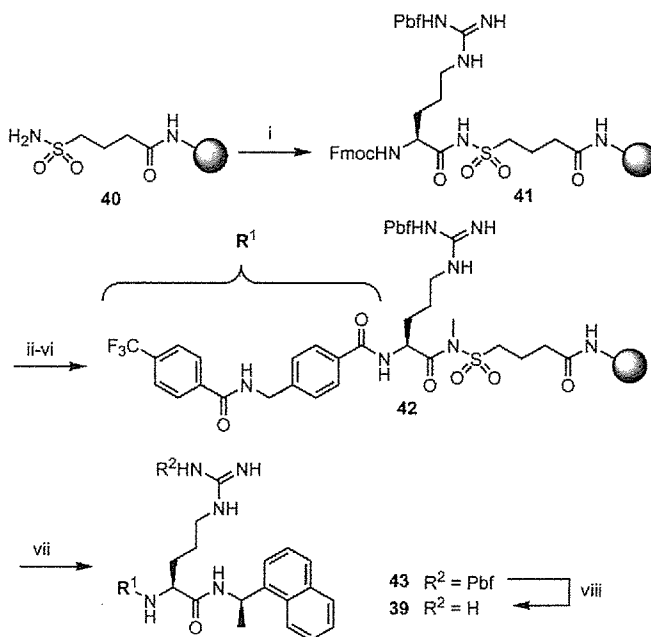
^a CC₅₀ values are based on the reduction of the viability of mock-infected MT-4 cells. Since the cytotoxicity of T140 was previously evaluated as CC₅₀ > 40 μM, further estimation at high concentrations was omitted in this study. ^b EC₅₀ values are based on the inhibition of HIV-induced cytopathogenicity in MT-4 cells. ^c IC₅₀ values are based on the inhibition of [¹²⁵I]-CXCL12 binding to CXCR4 transfectants of CHO cells. All data are the mean values for at least three independent experiments.

Thus, we synthesized two tetrapeptide mimetics, **4** and **5**, where a Tyr residue was added in the *N*-terminus of compounds **2** and **3**, respectively, based on the sequence of the FC131 sequence. Compounds **4** and **5** showed approximately twice stronger anti-HIV activity than compounds **2** and **3**, indicating that an *N*-terminal addition of a Tyr residue is effective for an increase in anti-HIV activity. Furthermore, compounds **4** and **5** exhibited stronger binding affinity for CXCR4, compared to compounds **1**–**3**. Next, we synthesized *p*-fluorobenzoylated tripeptide mimetics, **6** and **7**, based on the *N*-terminal sequence of 4F-benzoyl-TN14003 and 4F-benzoyl-TE14011. As a result, *p*-fluorobenzoylation caused an increase in anti-HIV activity. Compound **7** showed strong anti-HIV activity, suggesting that introduction of a reduced amide bond between two Arg residues is more suitable than that between Arg and naphthalenylethylamine. However, binding affinity of compound **7** for CXCR4 could not be exhibited until the 1 μM concentration, and compound **6** is weaker than compounds **4** and **5** in terms of binding affinity for CXCR4, although anti-HIV activity of compounds **6** and **7** is stronger than that of compounds **4** and **5**. This discrepancy might be caused by the difference between the interactive site of HIV and the binding site of CXCL12 on CXCR4.¹³

Since hit compounds with significant anti-HIV activity were found among several compounds that were synthesized using solution-phase techniques, we attempted to prepare more compounds by solid-phase synthesis: A tripeptide library containing three pharmacophores of the *N*-terminal region of 4F-benzoyl-TN14003 and 4F-benzoyl-TE14011 (aromatic ring, Arg² and Nal³) and the *C*-terminal carboxy amide was designed (Fig. 3). Since Arg¹ is not an indispensable residue for high activity, it was replaced by several spacers involving conformationally constrained units, such as 4-piperidinecarboxylic acid and 4-(aminomethyl)benzoic acid. Use of this library involving 20 synthetic compounds, which was constructed by solid-phase peptide synthesis (Fig. 3, library 1), led to the discovery of

lead compounds for anti-HIV agents, **11** and **14**, although these compounds did not show significant binding affinity for CXCR4 until the 1 μM concentration. Compound **8**, which contains Arg¹ based on the original 4F-benzoyl-TN14003 and 4F-benzoyl-TE14011 sequence, also exhibited moderate anti-HIV activity and significant CXCR4-binding affinity. These results suggest that Arg¹ can be replaced by conformationally restricted units in terms of anti-HIV activity. The other compounds that were contained in library 1 did not show significant anti-HIV activity until the 100 μM concentration.

Next, in due consideration of an increase in biostability, focused library of compounds with the *C*-terminal substituted amide was constructed based on the structures of compounds **8**, **11** and **14** by solid-phase techniques using Kenner's sulfonamide safety-catch linker¹⁴ (Fig. 3, library 2): *C*-terminal Nal-amide of compounds **8**, **11** and **14** was replaced by several amides possessing various naphthalene units. The synthetic scheme for compound **39** is shown as a representative in Scheme 1. Compounds **30** and **39** showed moderate and strong anti-HIV activity, respectively, although each compound did not show significant CXCR4-binding affinity until the 1 μM concentration. Anti-HIV potency of compounds is not always in proportion to binding affinity for CXCR4, especially in case of these small compounds, since there is a significant difference between the interactive site of HIV and the binding site of CXCL12 on CXCR4. There is a great interest in this result: compound **39**, possessing (*R*)-(+)-1-(1-naphthyl)ethylamine in the *C*-terminus, is stronger than compound **38**, possessing (*S*)-(–)-1-(1-naphthyl)ethylamine in the



Scheme 1 Reagents: (i) Fmoc-Arg(Pbf)-OH, DIPEA, PyBOP, CHCl₃; (ii) 20% (v/v) piperidine-DMF; (iii) Fmoc-(4-aminomethyl)benzoic acid, DIPCDI, HOBT, DMF; (iv) 20% (v/v) piperidine-DMF; (v) 4-trifluoromethylbenzoic acid, DIPCDI, HOBT, DMF; (vi) TMSCHN₂, hexane, THF; (vii) (*R*)-(+)-1-(1-naphthyl)ethylamine, DMF, reflux; (viii) thioanisole, TFA; Pbf = 2,2,4,6,7-pentamethyl-dihydrobenzofuran-5-sulfonyl, DIPEA = *N,N*-diisopropylethylamine, PyBOP = benzotriazole-1-yl-oxy-tris-pyrrolidino-phosphonium hexafluorophosphate, DIPCDI = *N,N*-diisopropylcarbodiimide, HOBT = *N*-hydroxybenzotriazole.

C-terminus, which is a common structure unit with KRH-1636. Compound 39 is thought to be a useful lead possessing chemically modified N- and C-terminal ends. The other 10 compounds that were contained in library 2 did not show significant anti-HIV activity until the 100 μ M concentration.

In summary, several compounds that were synthesized based on pharmacophores of T140 analogs showed significant anti-HIV activity and binding affinity for CXCR4. According to these results, two types of libraries based on the N-terminal region of 4F-benzoyl-TN14003 and 4F-benzoyl-TE14011 were constructed to find effective lead compounds. Linear-type low molecular weight compounds obtained in this study are thought to be useful leads for chemotherapy of AIDS, cancer and RA.

Acknowledgements

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Development of a ^{111}In -labeled peptide derivative targeting a chemokine receptor, CXCR4, for imaging tumors

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Abstract

The chemokine receptor CXCR4 is highly expressed in tumor cells and plays an important role in tumor metastasis. The aim of this study was to develop a radiopharmaceutical for the imaging of CXCR4-expressing tumors *in vivo*. Based on structure–activity relationships, we designed a 14-residue peptidic CXCR4 inhibitor, Ac-TZ14011, as a precursor for radiolabeled peptides. For ^{111}In -labeling, diethylenetriaminepentaacetic acid (DTPA) was attached to the side chain of D-Lys⁸ which is distant from the residues indispensable for the antagonistic activity. In-DTPA-Ac-TZ14011 inhibited the binding of a natural ligand, stromal cell-derived factor-1 α , to CXCR4 in a concentration-dependent manner with an IC_{50} of 7.9 nM (Ac-TZ14011: 1.2 nM). In biodistribution experiments, more ^{111}In -DTPA-Ac-TZ14011 accumulated in the CXCR4-expressing tumor than in blood or muscle. Furthermore, the tumor-to-blood and tumor-to-muscle ratios were significantly reduced by coinjection of Ac-TZ14011, indicating a CXCR4-mediated accumulation in tumor. These findings suggested that ^{111}In -DTPA-Ac-TZ14011 would be a potential agent for the imaging of CXCR4 expression in metastatic tumors *in vivo*.

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Keywords: CXCR4; Peptide radiopharmaceutical; Indium-111; Metastatic tumor

1. Introduction

Chemokines are a family of small proteins (8–14 kDa) that chemoattract leukocytes by binding to cell surface receptors, chemokine receptors [1]. The chemokine receptor family, which belongs to a superfamily of seven transmembrane domain G-protein-coupled receptors, comprises 18 members [2]. In 1996, one member, CXCR4, was identified as a coreceptor for the entry of T-cell line-tropic HIV-1 [3]. Since then, this receptor has attracted considerable attention as a pathogenic factor or a therapeutic target for HIV infection. Recent studies indicated that CXCR4 and its ligand, stromal cell-derived factor-1 (SDF-1), play an important role also in tumor metastasis [4–8]. Müller et al. [4] reported that CXCR4 was highly expressed in breast cancer and SDF-1 was highly expressed in organs

representing the first destinations of metastasis. Moreover, they demonstrated that neutralization with anti-CXCR4 monoclonal antibody significantly inhibited the metastasis of breast cancer cells in mice. Similar results were obtained in other types of cancer [5–8]. These findings suggest that CXCR4 is a potential target for the *in vivo* imaging of metastatic tumors.

We have previously demonstrated that a peptide with anti-HIV-1 activity, T22 ([Tyr^{5,12}, Lys⁷]-polyphemusin II), is an inhibitor of CXCR4 that blocks the entry of T-cell line-tropic HIV-1 mediated by this receptor. T22 is an 18-residue peptide amide, which was previously found by us based on an analysis of the structure–activity relationships of self-defense peptides of horseshoe crabs, tachyplesin and polyphemusin [9,10]. On the basis of the structure of T22, we designed and synthesized several downsized analogs, 14-residue peptides [11,12]. Among them, T140 showed the greatest inhibitory effect on the binding of an anti-CXCR4 monoclonal antibody to CXCR4 and the strongest inhibitory

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activity against HIV-1 entry [12]. The aim of this study was to develop a radiolabeled T140 derivative as an imaging agent for metastatic tumors. Considering that the three residues on the restricted backbone (L-3-(2-naphthyl)alanine (Nal)³, Tyr⁵ and Arg¹⁴) and the single residue in the flexible region (Arg²) form the intrinsic pharmacophore of T140 [13–15], we designed a 14-residue peptidic inhibitor, Ac-TZ14011, as the precursor for radiolabeled peptides (Fig. 1). This precursor contains the above four residues which are necessary for the inhibitory activity against CXCR4. Furthermore, for site-selective conjugation of radiolabels, Ac-TZ14011 has a single amino group (D-Lys⁸), which is distant from the pharmacophore, and the carboxyl group of Arg¹⁴ of Ac-TZ14011 is protected via amidation for stability in vivo [16,17].

¹¹¹In constitutes one of the most useful radionuclides for the radiolabeling of peptides for diagnostic applications in nuclear medicine. Diethylenetriaminepentaacetic acid (DTPA) is still an attractive chelating agent with which to prepare ¹¹¹In-labeled peptides since it provides ¹¹¹In-labeled peptides with highly specific activity. In addition, the development of a monoreactive DTPA derivative has provided an easy and efficient way to prepare DTPA-conjugated peptides [18,19]. In this study, DTPA-Ac-TZ14011 was prepared using a monoreactive DTPA derivative and coordinated with nonradioactive In or radioactive ¹¹¹In. Furthermore, the antagonistic activity of In-DTPA-Ac-TZ14011 and in vivo behavior of ¹¹¹In-DTPA-Ac-TZ14011 were investigated and the applicability of ¹¹¹In-DTPA-Ac-TZ14011 as a radiopharmaceutical for imaging tumors was evaluated.

2. Materials and methods

2.1. Reagents and chemicals

¹¹¹InCl₃ (74 MBq/ml in 0.02N HCl) was kindly supplied by Nihon Medi-Physics (Nishinomiya, Japan). 9-Fluorenylmethoxycarbonyl (Fmoc)-protected amino acids and 4-(2',4'-dimethoxyphenylaminomethyl)phenoxy (SAL) resin were purchased from Watanabe Chemical Industries (Hiroshima, Japan) or Calbiochem-Novabiochem Japan (Tokyo, Japan). 1-*tert*-Butyl hydrogen 3,6,9-tris((*tert*-butoxycarbonyl)methyl)-3,6,9-triazaundecanedioic acid (mDTPA) was synthesized as reported previously [18]. All the other chemicals were purchased from either Nacalai Tesque (Kyoto, Japan) or Wako Pure Chemical Industries (Osaka, Japan). Ion spray mass spectra (IS-MS) were obtained with the API III model (PerkinElmer Sciex Instruments, Thornhill, Canada). Cellulose acetate electrophoresis (CAE) strips were run in veronal buffer (pH 8.6, *I*=0.06) at a constant current of 0.8 mA for 40 min. Thin-layer chromatography (TLC) analyses were performed with silica plates (Silica gel 60, Merck, Darmstadt, Germany) with 10% aqueous ammonium chloride–methanol (1:1) as the developing solvent.

2.2. Synthesis of In-DTPA-Ac-TZ14011

Fig. 2 shows the scheme for the synthesis of In-DTPA-Ac-TZ14011. A protected peptide was constructed using Fmoc-based solid-phase synthesis on SAL resin and its N-terminus was acetylated. After being treated with thioanisole/trifluoroacetic acid (TFA) in the presence of *m*-cresol and 1,2-ethanedithiol, the crude peptide was air-oxidized and purified by reversed-phase HPLC (RP-HPLC). RP-HPLC was carried out with a Cosmosil 5C18-AR column (20×250 mm, Nacalai Tesque) eluted with a linear gradient of 10–30% acetonitrile in 0.1% aqueous TFA in 30 min at a flow rate of 7 ml/min. Fractions containing the peptide were collected, and the solvent was removed by lyophilization to afford Ac-TZ14011 as a white powder. IS-MS calcd for C₉₂H₁₄₄N₃₅O₁₉S₂ [M+H⁺]: *m/z* 2107.1, found: *m/z* 2107.4.

DTPA-Ac-TZ14011 was prepared by mDTPA conjugation. Briefly, to a solution of mDTPA (19 mg, 30.8 μmol) in acetonitrile (350 μl) were added *N*-hydroxysuccinimide (3.74 mg, 32.3 μmol) and *N,N*-dicyclohexylcarbodiimide (6.67 mg, 32.3 μmol) at 0°C, and the mixture was incubated overnight at room temperature. After cooling to 0°C again, 200 μl of Ac-TZ14011 (10.2 mg, 3.65 μmol) in a mixture of acetonitrile and phosphate-buffered saline (pH 7.4) (1:1) was added to the reaction mixture and incubated overnight at room temperature. After treatment with 95% TFA, the crude peptide was purified by RP-HPLC under the same conditions as above. IS-MS calcd for C₁₀₆H₁₆₅N₃₈O₂₈S₂ [M+H⁺]: *m/z* 2482.2, found: *m/z* 2482.9.

Fifty microliters of DTPA-Ac-TZ14011 (610 μg, 0.20 μmol) in 0.1 M acetic acid was reacted with 25 μl of nonradioactive InCl₃·4H₂O (64.5 μg, 0.22 μmol) in 0.02N HCl for 30 min at room temperature. Subsequent

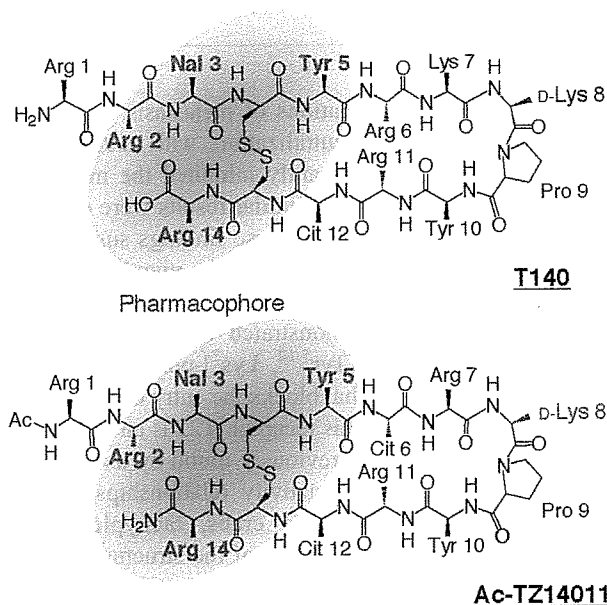


Fig. 1. Structures of T140 and Ac-TZ14011. There are four amino acid residues indispensable for the antagonistic activity (blue residues) which formed the pharmacophore. Nal: L-3-(2-naphthyl)alanine, Cit: L-citrulline. Ac-TZ14011 has a single amino group (D-Lys⁸) for site-selective conjugation of radiolabels, which is distant from the pharmacophore.

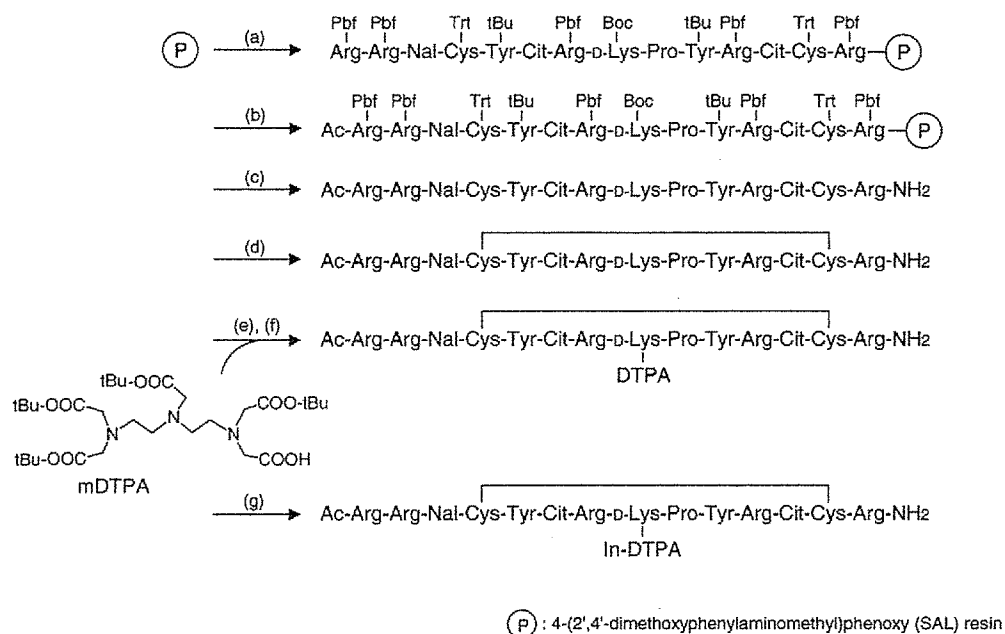


Fig. 2. Synthesis of In-DTPA-Ac-TZ14011. Reagents: (a) stepwise elongation; (b) acetic anhydride, pyridine; (c) trifluoroacetic acid, thioanisole, *m*-cresol, 1,2-ethanedithiol; (d) air oxidation; (e) *N*-hydroxysuccinimide, *N,N*-dicyclohexylcarbodiimide; (f) trifluoroacetic acid; (g) $\text{InCl}_3 \cdot 4\text{H}_2\text{O}$.

purification by RP-HPLC was carried out with a Hydro-sphere C18 column (4.6×250 mm, YMC, Kyoto, Japan) eluted with 17% acetonitrile in 0.1% aqueous TFA at a flow rate of 1 ml/min. Fractions containing the peptide were collected, and the solvent was removed by lyophilization to afford In-DTPA-Ac-TZ14011 as a white powder. IS-MS calcd for $\text{InC}_{106}\text{H}_{161}\text{N}_{38}\text{O}_{28}\text{S}_2$ [$\text{M}+\text{H}^+$]: m/z 2594.2, found: m/z 2594.3.

2.3. Synthesis of ^{111}In -DTPA-Ac-TZ14011

$^{111}\text{InCl}_3$ (3.7 MBq) in 0.02N HCl (100 μl) was added to DTPA-Ac-TZ14011 (10 μg) in 0.1 M acetic acid (200 μl), and the mixture was incubated for 30 min at room temperature. Then, ^{111}In -DTPA-Ac-TZ14011 was separated from DTPA-Ac-TZ14011 by RP-HPLC under the same conditions used for the purification of In-DTPA-Ac-TZ14011. The radiochemical purity of ^{111}In -DTPA-Ac-TZ14011 was determined by TLC, CAE and RP-HPLC.

2.4. Binding assay

The binding assay was performed according to the procedure of Hesselgesser et al. [20] with a slight modification. The stable CXCR4-transfected Chinese hamster ovary (CHO) cell lines were prepared by transfection with cDNA encoding alanine scanning mutants in pcDNA3 (Invitrogen, Carlsbad, CA, USA) using lipofectamine (GIBCO, Rockville, MD, USA) and selection in neomycin (G418 500 mg/ml; GIBCO). The expression of CXCR4 on the surface of each transfectant was measured by flow cytometry. CXCR4-transfected CHO cell lines were suspended in the binding buffer (Ham's F-12 containing 20 mM HEPES and 0.5% BSA) and placed in siliconized tubes (5×10^5 cells/120 μl /tube). Binding reactions were

performed on ice for 1 h in the presence of [^{125}I]SDF-1 α (PerkinElmer Life Sciences, Boston, MA, USA) and various concentrations of peptides. Cells were separated from the buffer by centrifugation through a dibutylphthalate/olive oil mixture. After removal of the water and oil layer, cell-associated radioactivity was measured. The 50% inhibitory concentration (IC_{50}) of peptides was determined based on inhibition of the binding of SDF-1 α to CXCR4-transfected CHO cells.

2.5. Calcium fluorimetry

Calcium fluorimetry was performed as described previously [21]. CXCR4-transfected CHO cell lines were placed in wells of a microtiter tray (3×10^4 cells/100 μl /well) and incubated for 1 day at 37°C in a CO_2 incubator. The cells were loaded with 5 μM of Fura-2-AM (Dojindo Laboratories, Kumamoto, Japan), 2.5 mM probenecid (Sigma, St Louis, MO, USA) and 20 mM HEPES (pH 7.4) in Ham's F-12 (80 μl /well) for 1 h at 37°C. After the cells were incubated with various concentrations of T140 analogs for 3 min, recombinant human SDF-1 α (PeproTech EC, London, UK) was added. Changes in intracellular Ca^{2+} concentrations were measured by a spectrofluorometer (96-well Fluorescence Drug Screening System, Hamamatsu Photonix, Hamamatsu, Japan) using a modified version of the Fura-2 method [22]. The IC_{50} of peptides was determined based on the inhibition of Ca^{2+} mobilization induced by SDF-1 α through CXCR4.

2.6. Biodistribution study in tumor-bearing mice

Animal experiments were conducted in accordance with our institutional guidelines and were approved by the Kyoto University Animal Care Committee. Athymic nude BALB/c

mice (8 weeks old, female) were inoculated subcutaneously with CXCR4-expressing pancreatic carcinoma cells, AsPC-1 [23,24]. When tumors were approximately 0.5 cm in diameter, the animals were intravenously injected with ^{111}In -DTPA-Ac-TZ14011 (25–30 kBq). The biodistribution of radioactivity was monitored at 1, 6 and 24 h postinjection. Groups of five mice were used for the experiments. Organs of interest were excised and weighed, and the radioactivity counts were determined with a well counter (ARC380CL, Aloka, Tokyo, Japan). For the *in vivo* blocking experiment, mice were coinjected with Ac-TZ14011 (10 mg/kg).

2.7. Statistical analysis

Statistical analysis was performed by using the unpaired *t*-test. $P < .05$ was considered to be statistically significant.

3. Results and discussion

T140 and its analogs have one disulfide bond and maintain an antiparallel β -sheet structure connected by a type II' β -turn with D-Lys⁸-Pro⁹ at the (*i*+1) and (*i*+2) positions, and the side chain of D-Lys⁸ is distant from the pharmacophore for the antagonistic activity [14,15]. Therefore, we designed Ac-TZ14011 as a mother compound that contains the residues indispensable for the antagonistic activity and has a single amino group of D-Lys⁸ for site-selective conjugation of DTPA (Fig. 1). In calcium fluorimetric assays, this compound showed strong inhibitory activity equal to that of T140 (Table 1). To assess the effect of the conjugation of ^{111}In -DTPA with Ac-TZ14011 on the antagonistic activity toward CXCR4, nonradioactive ^{111}In -DTPA-Ac-TZ14011 was synthesized (Fig. 2). In binding assays with CXCR4, ^{111}In -DTPA-Ac-TZ14011 maintained strong inhibitory activity although its IC_{50} value was slightly larger than that of Ac-TZ14011 (Table 1). This result indicated the validity of the chemical design of ^{111}In -DTPA-Ac-TZ14011 based on structure–activity relationships.

In RP-HPLC analyses, ^{111}In -DTPA-Ac-TZ14011 and DTPA-Ac-TZ14011 showed well-separated peaks as shown in Fig. 3. After purification by RP-HPLC under the same conditions, ^{111}In -DTPA-Ac-TZ14011 was obtained with high radiochemical purity (over 96%) as determined by TLC, CAE and RP-HPLC. The radioactivity pharmacoki-

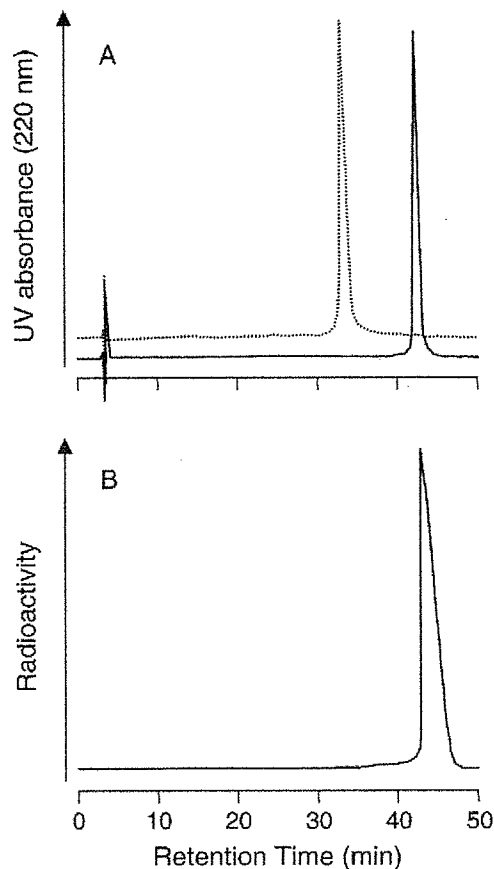


Fig. 3. Reversed-phase HPLC profiles of nonradioactive ^{111}In -DTPA-Ac-TZ14011 (solid line), DTPA-Ac-TZ14011 (broken line) (A) and ^{111}In -DTPA-Ac-TZ14011 (B).

netics of ^{111}In -DTPA-Ac-TZ14011 was evaluated in nude mice bearing the CXCR4-expressing pancreatic carcinoma AsPC-1 (Table 2). ^{111}In -DTPA-Ac-TZ14011 showed a rapid clearance from the blood and a marked accumulation and retention in the liver, kidney and spleen. The accumulation of radioactivity was greater in the tumor than in the blood or muscle (Table 2). In mice, CXCR4 mRNA is highly expressed in various lymphoid tissues and cells such as spleen, thymus, lymph node, bone marrow and leukocytes [25,26]. Thus, since liver and spleen are concerned with the immune system, the accumulation of ^{111}In -DTPA-Ac-TZ14011 in these organs should be mediated by CXCR4-binding. In fact, coinjection of Ac-TZ14011 significantly reduced the accumulation in the liver by over one-tenth and in the spleen by over one-third. This marked reduction of radioactivity in the liver and spleen on the coinjection of Ac-TZ14011 caused the high levels of radioactivity in the blood and consequently increased the accumulation of radioactivity in organs which were small in size and/or did not take up much radioactivity (Table 2). The accumulation in the tumor was also increased by coinjection of Ac-TZ14011, but the tumor-to-blood and tumor-to-muscle ratios were significantly reduced (Table 2). Since there is very little or no CXCR4 in the muscle [26],

Table 1

Antagonistic activity of T140 derivatives

| | IC_{50} (nM) | |
|------------------------------------|-------------------------------------|--|
| | SDF-1 α binding ^a | Ca^{2+} mobilization ^b |
| ^{111}In -DTPA-Ac-TZ14011 | 7.9 | ND ^c |
| Ac-TZ14011 | 1.2 | 2.6 |
| T140 | ND ^c | 2.2 |

^a Values are the concentrations for 50% inhibition of the binding of [^{125}I]SDF-1 α to CXCR4.

^b Values are the concentrations for 50% inhibition of Ca^{2+} mobilization induced by SDF-1 α through CXCR4.

^c Not determined.

Table 2
Biodistribution of radioactivity after intravenous injection of ^{111}In -DTPA-Ac-TZ14011 in nude mice bearing pancreatic carcinoma, AsPC-1

| | 1 h | 6 h | 24 h | 1 h+ Ac-TZ14011 ^a |
|------------------------|-------------|-------------|-------------|---------------------------------|
| Blood ^b | 0.39 (0.06) | 0.05 (0.01) | 0.03 (0.01) | 2.06** (0.61) |
| Liver ^b | 27.0 (2.9) | 25.2 (2.0) | 19.3 (2.5) | 1.95** (0.25) |
| Kidney ^b | 50.9 (4.3) | 43.4 (6.3) | 29.5 (5.7) | 45.4 (6.8) |
| Spleen ^b | 8.22 (0.70) | 7.57 (0.54) | 5.83 (0.99) | 2.66** (1.21) |
| Pancreas ^b | 0.15 (0.03) | 0.05 (0.01) | 0.05 (0.02) | 1.07** (0.46) |
| Muscle ^b | 0.17 (0.05) | 0.07 (0.02) | 0.07 (0.01) | 1.35** (0.26) |
| Tumor ^b | 0.51 (0.08) | 0.20 (0.03) | 0.14 (0.03) | 1.70** (0.27) |
| T/B ratio ^c | 1.31 (0.14) | 4.05 (0.79) | 5.65 (2.89) | 0.88* (0.31) |
| T/M ratio ^d | 3.17 (0.99) | 4.43 (1.89) | 3.23 (1.08) | 1.31** (0.41) |

Each value represents the mean (S.D.) for five animals.

^a Coinjection with unlabeled Ac-TZ14011 (10 mg/kg).

^b Expressed as % injected dose per gram.

^c Tumor-to-blood ratio.

^d Tumor-to-muscle ratio.

* $P < .05$, comparison between ^{111}In -DTPA-Ac-TZ14011 with or without unlabeled Ac-TZ14011 at 1 h.

** $P < .005$, comparison between ^{111}In -DTPA-Ac-TZ14011 with or without unlabeled Ac-TZ14011 at 1 h.

tumor-to-muscle ratios reflect target-to-nontarget ratios. Thus, the reduction in the tumor-to-muscle ratio caused by the coinjection of Ac-TZ14011 indicated that ^{111}In -DTPA-Ac-TZ14011 accumulated in the tumor through CXCR4. On the other hand, coinjection of Ac-TZ14011 did not alter the levels of ^{111}In -DTPA-Ac-TZ14011 in the kidney, suggesting a nonspecific accumulation. This is consistent with previous findings that CXCR4 mRNA levels expressed in the kidney were very low [25,26]. Recent studies indicated that an electrostatic interaction between positively charged peptides and the negatively charged surface of renal proximal tubular cells plays an important role in the reabsorption of peptides into proximal tubular cells [27–29]. Since five Arg residues are contained in the peptide ^{111}In -DTPA-Ac-TZ14011, the highly positive charge would cause a greater nonspecific accumulation in the kidney even compared to other ^{111}In -DTPA peptides [28,30,31]. Due to its accumulation in nontarget organs, ^{111}In -DTPA-Ac-TZ14011 may be unavailable as a radiopharmaceutical for screening small tumors, particularly in the kidneys and their surroundings.

It was reported that CXCR4 expression could be a powerful predictive factor for prognosis (recurrence, metastasis or survival rate) in colorectal cancer [32,33], malignant melanoma [34] and osteosarcoma [35]. Therefore, a CXCR4 imaging agent would be a new type of radiopharmaceutical for predicting the prognosis of cancer patients. CXCR4 also represents a novel target for tumor therapy, and some CXCR4 inhibitors have been investigated as anti-metastatic agents [36–39]. These agents showed positive effects in suppressing tumor metastasis; however, they would also have deleterious effects on normal physiological functions since CXCR4 plays a crucial role in numerous biological processes [2]. Therefore, in vivo imaging of CXCR4 expression could be a potential method for determining

the dose of anti-metastatic agents and for monitoring their therapeutic efficacy.

In conclusion, we designed ^{111}In -DTPA-Ac-TZ14011 based on the structure–activity relationships of peptidic CXCR4 inhibitors. In-DTPA-Ac-TZ14011 showed strong inhibitory activity against the binding of CXCR4 to an endogenous ligand. Furthermore, the accumulation of ^{111}In -DTPA-Ac-TZ14011 in the CXCR4-expressing tumor was greater than that in the blood or muscle, being mediated by this receptor. These findings suggest that ^{111}In -DTPA-Ac-TZ14011 is a potential radiopharmaceutical for the imaging of CXCR4 expression in metastatic tumors in vivo for predicting the prognosis of cancer patients and for monitoring the therapeutic efficacy of anti-metastatic agents.

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Brief Articles

Identification of a New Class of Low Molecular Weight Antagonists against the Chemokine Receptor CXCR4 Having the Dipicolylamine–Zinc(II) Complex Structure

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Several low molecular weight nonpeptide compounds having the dipicolylamine–zinc(II) complex structure were identified as potent and selective antagonists of the chemokine receptor CXCR4. These compounds showed strong inhibitory activity against CXCL12 binding to CXCR4, and the top compound exhibited significant anti-HIV activity. Zinc(II)–dipicolylamine unit-containing compounds proved to be useful and attractive lead compounds for chemotherapy of these diseases as nonpeptide CXCR4 antagonists possessing the novel scaffold structure.

Introduction

CXCR4 is a chemokine receptor that transduces signals of its endogenous ligand, CXCL12/stromal cell-derived factor-1 (SDF-1).^{1–4} CXCR4 is classified into 7TMGPCR and plays a physiologically critical role by the action of CXCL12 in the migration of progenitors during embryologic development of the cardiovascular, hemopoietic, central nervous systems, etc. In addition, CXCR4 was previously identified as a coreceptor that is used by X4-HIV-1 in its entry into T cells⁵ and has recently been proven to be involved in several problematic diseases, including HIV infection, metastasis of several types of cancer,^{6–8} leukemia cell progression,^{9,10} rheumatoid arthritis (RA).^{11,12} Thus, CXCR4 is thought to be a great therapeutic target to overcome these diseases, and several inhibitors directed against CXCR4 have been developed to date.^{13–22} We previously found a highly potent CXCR4 antagonist, T140, which is a 14-mer peptide with a disulfide bridge, and its downsized derivative, FC131, which has a cyclic pentapeptide scaffold structure (Table 1).^{18–21} Although reduction of the peptide character based on these peptides is underway,^{23,24} we would like to discover novel CXCR4 antagonists having nonpeptide structures, since few nonpeptide compounds with low molecular weight have been reported, such as AMD3100 series^{14,16} and KRH-1636.²² Previously, anthracene derivatives having two sets of zinc(II)–2,2′-dipicolylamine (Dpa) complex were identified as the first chemosensors that can selectively bind and sense phosphorylated peptide surfaces.²⁵ In the present study, we have found several aromatic compounds having the zinc(II)–Dpa structure to be a new class of low molecular weight CXCR4 antagonists.

Experimental Section

Chemistry. Synthesis of Bis(dipicolylamine)-*p*-xylene–Zn Complexes. Aromatic compounds having the zinc(II)–Dpa structure were previously synthesized as reported elsewhere.^{25–28}

For a comparative study, bis(3,3′- and bis(4,4′-dipicolylamine)-*p*-xylenes, **24** and **25**, respectively, were synthesized by treatment of *p*-xylenediamine with the corresponding pyridinecarbaldehydes and sodium triacetoxyborohydride [NaBH(OAc)₃] (Figure 1).²⁹ Zinc(II) complexation in the preparation of **18** and **19** was performed by treatment of bis(dipicolylamine)-*p*-xylenes with NaOH to afford salt-free compounds, followed by addition of aqueous zinc nitrate [Zn(NO₃)₂].

Biological Assays. Calcium mobilization,³⁰ [¹²⁵I]CXCL12 binding (oil cushion method),³⁰ and anti-HIV²³ assays were performed as reported previously.

Molecular Modeling Calculations. Molecular modeling calculations were performed using SYBYL program (version 7.0, TRIPOS Inc.). Energy minimizations were performed using Tripos force field. The lowest energy conformation was obtained by random search method.

Biological Results and Discussion

Several aromatic compounds having the zinc(II)–dipicolylamine structure were prepared and surveyed for CXCR4-antagonistic activity based on inhibitory activity against Ca²⁺ mobilization induced by CXCL12 stimulation through CXCR4. The structures and CXCR4-antagonistic activity of these compounds are shown in Tables 1–3. Positive controls **8** (T140), **9** (FC131), and **10** (KRH-1636) showed strong antagonistic activity. Compound **2**, which has two sets of the [bis(pyridin-2-ylmethyl)amino]methylene unit with zinc(II) complexation at the para-position of benzene, showed potent CXCR4-antagonistic activity (IC₅₀ = 0.1 μM). Compound **1**, which has a piece of this unit, did not show any activity until 1 μM. It suggests that two sets of this unit are required for binding to CXCR4. Compound **3**, which has two sets of this unit at the meta-position of benzene, showed lower activity than compound **2**, suggesting that the presence of this unit at the para-position is critical for

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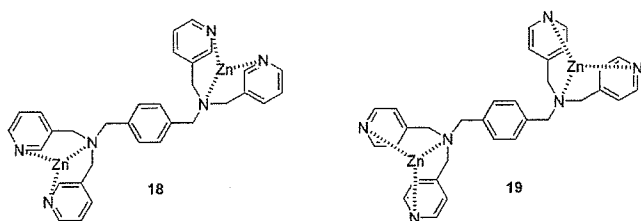
^{||} St. Marianna University.

[⊥] National Institute of Infectious Diseases.

Table 1. Structures and CXCR4-Antagonistic Activity of Aromatic Compounds Having the Zinc(II)–Dipicolylamine Structure (I)

| Compd. (No.) | Structure | IC ₅₀ (μM) ^a |
|--------------|-----------|------------------------------------|
| 1 | | > 1 |
| 2 | | 0.10 |
| 3 | | 0.49 |
| 4 | | 0.35 |
| 5 | | 0.12 |
| 6 | | 0.46 |
| 7 | | 0.10 |
| 8 | | 0.0036 |
| 9 | | 0.036 |
| 10 | | 0.040 |

^a IC₅₀ values are based on the inhibition against Ca²⁺ mobilization induced by CXCL12 stimulation through CXCR4. All data are the mean values for at least two independent experiments.

**Figure 1.** Structures of the zinc(II)–bis(3,3'- and –bis(4,4'-dipicolylamine)-*p*-xylylene (**24** and **25**) complexes, **18** and **19**, respectively.

strong CXCR4-antagonistic activity. Biphenyl compounds having two sets of this unit at the 3,3'- and 4,4'-positions, **4** and **5**, respectively, exhibited significantly high antagonistic activity, although the compound with the 4,4'-positions substituted (**5**) is stronger than that having the 3,3'-positions substituted (**4**). It seems to be important that two sets of the [bis(pyridin-2-ylmethyl)amino]methylene unit with zinc(II) complexation are

Table 2. Structures and CXCR4-Antagonistic Activity of Aromatic Compounds Having the Zinc(II)–Dipicolylamine Structure (II)

| Compd. (No.) | Structure | IC ₅₀ (μM) |
|--------------|-----------|-----------------------|
| 11 | | > 1 |
| 12 | | 0.18 |
| 13 | | 0.77 |
| 14 | | 0.24 |
| 15 | | 0.75 |
| 16 | | > 1 |
| 17 | | > 1 |

Table 3. Structures and CXCR4-Antagonistic Activity of Aromatic Compounds Having the Zinc(II)–Dipicolylamine Structure and Zinc-Free Compounds

| compd | structure | IC ₅₀ (μM) |
|-----------|---------------------------------|-----------------------|
| 18 | shown in Figure 1 | >10 |
| 19 | shown in Figure 1 | >10 |
| 20 | zinc-free analogue of 2 | >10 |
| 21 | zinc-free analogue of 5 | >10 |
| 22 | zinc-free analogue of 7 | >10 |
| 23 | zinc-free analogue of 14 | >10 |
| 24 | zinc-free analogue of 18 | >10 |
| 25 | zinc-free analogue of 19 | >10 |

located at 180° of each other as in compounds **2** and **5**, which both have almost the same potency. Furthermore, replacement of the biphenyl unit of compounds **4** and **5** by [2,2']bipyridinyl did not cause significant change in CXCR4-antagonistic activity, since the [2,2']bipyridinyl compounds **6** and **7** showed almost the same potency as the biphenyl compounds **4** and **5**, respectively. It may be a matter of course that a [2,2']bipyridinyl compound having only one [bis(pyridin-2-ylmethyl)amino]methylene unit with zinc(II) complexation (**11**) did not show any activity until 1 μM, as compound **1** did not. A naphthyl compound (**12**) and an anthracenyl compound (**14**), which have two sets of this unit at the 1,4-positions and at the 9,10-positions,

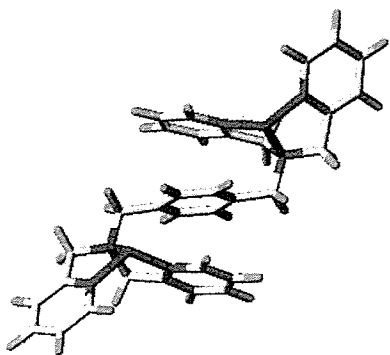


Figure 2. The structure calculated by molecular modeling of compound **2** having the zinc(II)–Dpa structure. Nitrogen atoms, blue; hydrogen atoms, sky blue; zinc atoms, red.

respectively, showed high activity. However, it suggests that the addition of one and two benzene moieties to compound **2** caused a slight decrease in activity. Unexpectedly, an anthracenyl compound having only one [bis(pyridin-2-ylmethyl)amino]methylene unit with zinc(II) complexation (**13**) exhibited not so high but significant antagonistic activity. It may be reasonable that the anthracenyl compound **15**, which has two sets of this unit at the 1,8-positions, did not show high antagonistic activity, since it is thought to be important that these units are located 180° from each other. In addition, terphenyl compounds **16** and **17** did not show any significant activity until 1 μM , suggesting that it might be unsuitable that two sets of this unit be located far apart from each other.

Next, to investigate whether the location of the nitrogen atom in the pyridine ring of the [bis(pyridin-2-ylmethyl)amino]methylene unit is critical for expression of CXCR4-antagonistic activity, compounds **18** and **19**, which contained the [bis(pyridin-3-ylmethyl)amino]methylene and [bis(pyridin-4-ylmethyl)amino]methylene units, respectively, with zinc(II) complexation, were synthesized. Neither compound **18** nor **19** showed any significant activity, although antagonistic activity was estimated to 10 μM . This result suggested that the location of the nitrogen atom in the pyridine ring is important either for formation of active conformation or for stable complexation with zinc(II).

Molecular modeling simulation analysis showed that the bis-(3,3'-dipicolylamine) and bis(4,4'-dipicolylamine)-*p*-xylene Zn complexes **18** and **19** did not converge as well as bis(Dpa)–Zn complex **2** did (Figure 2). Values of the coordinate bond lengths between zinc and nitrogen atoms (of the pyridine rings/of the tertiary amine) in the bis(Dpa)–Zn complex **2** are 1.93–1.95 Å, according to molecular modeling calculations. Complex **2** forms a stable conformation, having π – π stacking among three aromatic rings, as shown in Figure 2. However, values of bond lengths between zinc and nitrogen atoms of the pyridine rings in the complexes **18** and **19** would be ca. 2.9 and ca. 4.2 Å, respectively, which are relatively long. In addition, the proton atoms at the positions 2 and 2' in the pyridine rings might interfere with the zinc atom to prevent the molecules from forming a stable coordinate conformation.

Furthermore, to verify the indispensability of zinc(II) complexation, zinc-free analogues of **2**, **5**, **7**, and **14**, compounds **20**–**23**, respectively, were assessed for CXCR4-antagonistic activity. Since these zinc-free compounds did not show any significant activity until 10 μM , zinc(II) atoms or conformation constrained by zinc(II) complexation might be indispensable for binding to CXCR4. As a matter of course, zinc-free analogues of **18** and **19**, compounds **24** and **25**, respectively, did not show any significant activity.

Table 4. CXCR4-Binding Activity of Compounds **2**, **5**, **7**, and **12**

| compd | IC ₅₀ (μM) ^a | compd | IC ₅₀ (μM) ^a |
|-----------|---|----------------------|---|
| 2 | 0.047 | 8 (T140) | 0.00093 |
| 5 | 0.18 | 9 (FC131) | 0.0030 |
| 7 | 0.22 | 10 (KRH-1636) | 0.034 |
| 12 | 0.42 | | |

^a IC₅₀ values are based on the inhibition of [¹²⁵I]CXCL12 binding to CXCR4 transfectants of CHO cells. All data are the mean values for at least two independent experiments.

Next, we investigated CXCR4-binding activity of the novel compounds that possess strong CXCR4-antagonistic activity (Table 4). Compounds **2**, **5**, **7**, and **12** showed potent binding activity. Especially, the potency of compound **2** is comparable to that of KRH-1636.

The best and simple compound among the present compounds (**2**) was evaluated for anti-HIV activity. Compound **2** showed significant inhibitory activity against X4-HIV-1-induced cytopathogenicity in MT-4 cells (EC₅₀ = 7.1 μM), although anti-HIV activity in cells is lower than CXCR4-antagonistic or -binding activity as usual.²³ Furthermore, the present compounds identified as CXCR4 antagonists showed no significant inhibition (<25%) at 10 μM against Ca²⁺ mobilization induced by MIP-1 α stimulation through CCR5 and at 30 μM against Ca²⁺ mobilization induced by sphingosine 1-phosphate stimulation through EDG3 (GPCR).

The present compounds, such as **2**, **5**, **7**, and **12**, have been prepared as binuclear zinc complexes for the use in several assays. The extracellular concentration of zinc is normally ~100 $\mu\text{g}/100$ mL (approximately 15 μM): 30% of the total zinc is tightly bound to the metal-binding proteins. The remaining amount (70%) is loosely bound to proteins and easily released from the corresponding proteins.³¹ Thus, the extracellular concentration of zinc is sufficiently high for the compounds to be active in vivo. Furthermore, the dipicolylamine (Dpa) unit forms a stable complex with zinc ion (log K = 7.57), indicating that the compounds can maintain an active state as the zinc complex in vivo. The affinity of the Dpa unit for Ca²⁺ and Mg²⁺ ions, which are biologically essential, is considerably low (log K < 3). Thus, it is thought that these ions might not affect the zinc complexes.

Conclusion

The current study presents a new class of nonpeptide CXCR4 antagonists with low molecular weight that have a novel scaffold: a dipicolylamine–zinc(II) complex structure. These compounds showed selective and strong CXCR4-antagonistic activity. These compounds also have basic and aromatic moieties in common with several reported CXCR4 antagonists, e.g., T140, FC131, AMD3100 and KRH-1636, suggesting that these moieties are critical for interaction with CXCR4. The present results provide useful insights for the future design of new CXCR4 antagonists in association with information from other CXCR4 antagonists for development of therapeutic strategies for CXCR4-relevant diseases. Furthermore, anthracene derivatives having two sets of zinc(II)–dipicolylamine, such as compound **14**, might be used as chemical probes to study the biology of CXCR4, as these compounds are used to sense phosphorylated peptide surfaces.

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Supporting Information Available: Additional experimental procedures of novel synthetic compounds. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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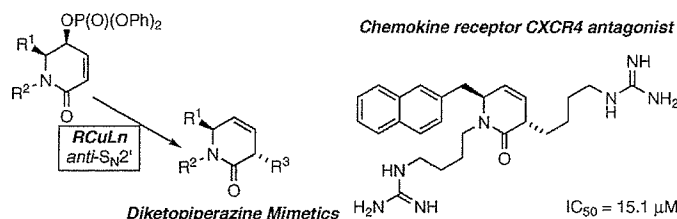
Stereoselective Synthesis of
**3,6-Disubstituted-3,6-dihydropyridin-2-ones as Potential
 Diketopiperazine Mimetics Using Organocopper-Mediated *anti*-S_N2'¹
 Reactions and Their Use in the Preparation of Low-Molecular
 CXCR4 Antagonists**

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Organocopper-mediated *anti*-S_N2' reactions of γ -phosphoryloxy- α,β -unsaturated- δ -lactams were used to prepare highly functionalized diketopiperazine mimetics. The substrate phosphates **24**, **32**, and **47** were prepared from α -amino acid-derived allylic alcohols **10** by a sequence of reactions that included ring-closing metathesis. In the reactions of phosphates with organocopper reagents, the addition of LiCl dramatically improved *anti*-S_N2' selectivity, indicating that an organocopper cluster containing lithium chloride plays an important role in the determination of regioselectivity. This reaction system was applied to the preparation of novel low molecular weight CXCR4-chemokine receptor antagonists.

Introduction

The replacement of a peptide bond in biologically active peptides with a non-hydrolyzable unit is a promising approach for peptide-lead drug discovery. (*E*)-Alkene dipeptide isosteres (EADIs) represent amide-bond mimetics that possess excellent structural homology and resistance to proteases.¹ Over the past decade, we have engaged in the development of efficient stereoselective methodology for the preparation of EADIs utilizing organocopper-mediated transformations along with their application to biologically active peptides.^{2,3} Piperazine-2,5-dione (diketopiperazine: DKP) derivatives **1** represent the smallest cyclic peptides consisting of two α -amino acid residues.

This well-organized structure is widely seen in compounds of biological or medicinal interest. Therefore, it seemed logical that the DKP nucleus could serve as a drug template with appropriately arrayed pharmacophores (Figure 1).⁴ On the basis of our studies on EADIs, we envisioned that the replacement of a DKP *cis*-amide bond with structurally similar (*Z*)-alkene units could provide DKP mimetics **2** as novel starting points for creating drug-like structures (Figure 2).

(1) For alkene dipeptide isosteres, see: (a) Christos, T. E.; Arvanitis, A.; Cain, G. A.; Johnson, A. L.; Pottorf, R. S.; Tam, S. W.; Schmidt, W. K. *Bioorg. Med. Chem. Lett.* **1993**, *3*, 1035. (b) Bohnstedt, A. C.; Prasad, J. V. N. V.; Rich, D. H. *Tetrahedron Lett.* **1993**, *34*, 5217. (c) Wai, J. S.; Bamberger, D. L.; Fisher, T. E.; Graham, S. L.; Smith, R. L.; Gibbs, J. B.; Mosser, S. D.; Oliff, A. I.; Pompliano, D. L.; Rands, E.; Kohl, N. E. *Bioorg. Med. Chem.* **1994**, *2*, 939. (d) Vasbinder, M. M.; Jarvo, E. R.; Miller, S. J. *Angew. Chem., Int. Ed.* **2001**, *40*, 2824. (e) Wipf, P.; Xiao, J. *Org. Lett.* **2005**, *7*, 103. (f) Jenkins, C. L.; Vasbinder, M. M.; Miller, S. J.; Raines, R. T. *Org. Lett.* **2005**, *7*, 2619.

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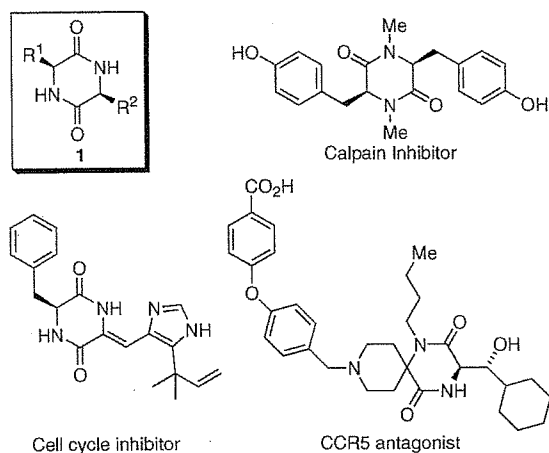


FIGURE 1. General structure of 2,5-diketopiperazine and biologically active derivatives.

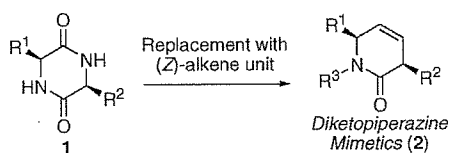
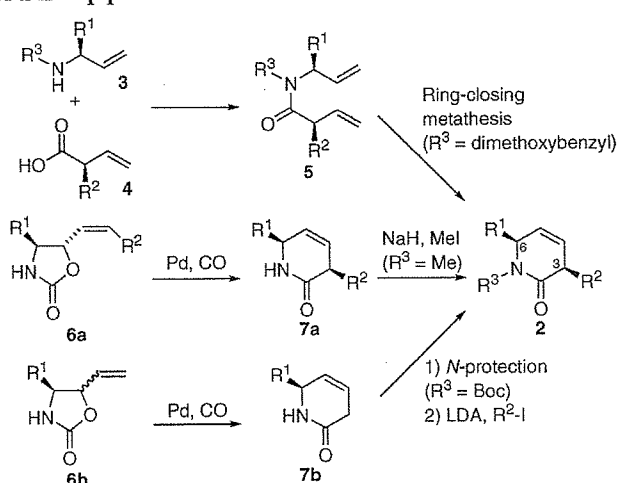


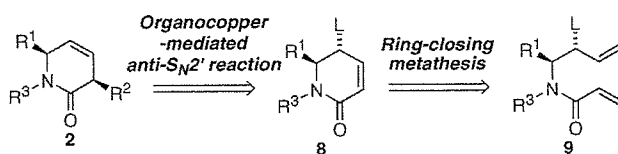
FIGURE 2. Design of diketopiperazine mimetics.

However, to our knowledge, there have been only a few reports in which 3,6-disubstituted-3,6-dihydropyridin-2-ones (DKP mimetics **2**) having side-chain functionalities have been synthesized in stereoselective fashions. These include synthetic protocols that employ the ring-closing metathesis or palladium-catalyzed carbonylation as key reactions (Scheme 1). Ring-closing metathesis of bisolefinic amide **5** with Grubbs' ruthenium alkylidene complexes yielded the desired DKP mimetics **2**, where the stereochemical outcome depends on the stereochemistry of the requisite metathesis substrate **5**, which was obtained by coupling between enantiomerically pure 1-substituted prop-2-enylamines **3** and 2-substituted but-3-enoic acids **4**.⁵ Alternatively, Knight et al. have reported the enantioselective synthesis of 3,6-dihydro-1*H*-pyridin-2-ones **7** by Pd-catalyzed decarboxylative carbonylation of 5-vinylloxazolidin-2-ones **6**,

SCHEME 1. Previous Synthetic Routes for the Preparation of Diketopiperazine Mimetics



SCHEME 2. Retrosynthetic Analysis of Diketopiperazine Mimetics Prepared by an Organocopper-Mediated *anti*- S_N2' Reaction^a



^a L = leaving group.

which were synthesized from the corresponding α -amino aldehydes.⁶ The stereochemistry at the 6-position was derived from that of the precursor α -amino aldehyde. The stereoselective introduction of substituents at the 3-position was achieved in two ways. One way involves the Pd-catalyzed reaction of 5-(alk-1-enyl)oxazolidinones **6a** to provide 3,6-disubstituted analogues **7a**,^{6a,b} however, this requires long reaction times, and the product yields were rather low. To circumvent these problems, a two-step protocol consisting of the Pd-catalyzed synthesis of the 6-substituted pyridinones **7b** followed by an enolate alkylation (**7b** to **2**) was developed.^{6c} In this methodology, the nature of both the electrophiles incoming to the enolate and the substituent on the nitrogen affect the diastereoselection at the 3-position.

Our approach for the stereoselective preparation of DKP mimetics **2** is shown in Scheme 2. We envisioned that the γ -activated- α,β -unsaturated lactams **8** could be converted into the corresponding dihydropyridinone derivatives **2** by an organocopper-mediated *anti*- S_N2' reaction. A Ru-catalyzed olefin metathesis⁷ reaction is suitable for the synthesis of key substrates

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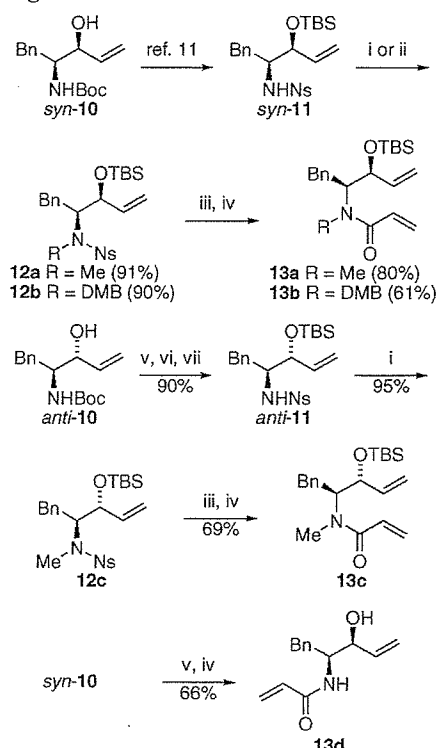
8, in terms of compatibility with various functional groups.⁸ The amide nitrogen should be appropriately protected by a group such as Me or dimethoxybenzyl (DMB) to raise the proportion of conformers with *cis*-amide geometry, which is necessary for facile olefin metathesis.⁹ We present herein the efficient conversion of γ -oxygenated- α,β -unsaturated- δ -lactams **8** to DKP mimetics **2** using organocopper-mediated *anti*-S_N2' reactions. This novel methodology was applied to the synthesis of low molecular weight CXCR4 antagonists on the basis of the DKP mimetic structure.¹⁰

Results and Discussion

Our synthesis started from the known *N*-Ns-*O*-TBS protected *syn*-1,2-amino alcohol *syn*-**11**¹¹ (Ns: 2-nitrobenzenesulfonyl; TBS: *tert*-butyldimethylsilyl) derived from allylic alcohol *syn*-**10**¹² (Scheme 3). The silyl ether *syn*-**11** was subjected to *N*-modification either with MeI, K₂CO₃, or with DMB-OH, Ph₃P, and DEAD (Mitsunobu conditions) to afford *N*-Me **12a** and *N*-DMB **12b** derivatives, respectively.¹³ After removal of the Ns group of **12** by treatment with thiolate anion under basic conditions, the resulting secondary amines were acylated with acryloyl chloride to yield the metathesis substrates **13**. The corresponding diastereomer **13c** was synthesized from *N*-Boc protected *anti*-1,2-amino alcohol *anti*-**10**¹² by a sequence of reactions identical to those used for the preparation of **13a**. Nonalkylated derivative **13d** was also synthesized.

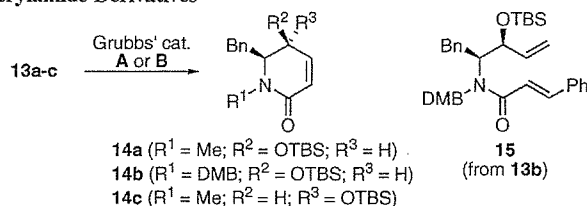
The attempted olefin metathesis of **13a–c** with Grubbs' Ru catalyst **A**¹⁴ resulted in low cyclization yields (Table 1, entries 1–4). The use of the second-generation catalyst **B**¹⁵ improved the yields (entries 5–7), although long reaction times under reflux were required. Ring-closing metathesis of **13** with catalyst **B** gave the benzylidene derivative **15** as a side product. This prompted us to postulate that the low reactivity of the substrates may be partly attributed to the presence of the bulky TBS group. Therefore, TBS-deprotected derivatives **16a–c**, obtained in high yields by the treatment of **13a–c** with TBAF (88–99%), were subjected to the metathesis reaction with catalyst **A** or **B** (Table 2). Although the reaction with catalyst **A** did not afford satisfactory results (entries 1 and 2), ring-closure with catalyst

SCHEME 3. Synthesis of Requisite Substrates for Ring-Closing Metathesis^a



^a Reagents and conditions: (i) MeI, K₂CO₃, DMF; (ii) DMB-OH, PPh₃, DEAD, THF; (iii) HSCH₂CO₂H, LiOH, DMF; (iv) CH₂=CHCOCl, Et₃N, CH₂Cl₂; (v) 4 M HCl in dioxane; (vi) Ns-Cl, 2,4,6-collidine, CHCl₃; (vii) TBSOTf, 2,6-lutidine, CH₂Cl₂.

TABLE 1. Ring-Closing Metathesis of *O*-TBS-Protected Acrylamide Derivatives



| entry | substrate | cat. (equiv) | conditions ^a | product ^{b,c} (yield, %) |
|-------|------------|--------------|-------------------------|-----------------------------------|
| 1 | 13a | A (0.6) | rt, 48 h | 14a (trace) |
| 2 | 13a | A (0.6) | reflux, 36 h | 14a (37) |
| 3 | 13b | A (0.1) | reflux, 36 h | 14b (10) |
| 4 | 13c | A (0.6) | reflux, 36 h | 14c (30) |
| 5 | 13a | B (0.6) | rt, 36 h | 14a (57) |
| 6 | 13a | B (0.6) | reflux, 48 h | 14a (61) |
| 7 | 13b | B (0.6) | reflux, 36 h | 14b (29) |

^a CH₂Cl₂ was used as solvent. ^b Isolated yield. ^c Starting materials were recovered, except for entry 6.

B proceeded smoothly at room temperature to yield the desired cyclized compounds **17a–c** in good yields (entries 3–6). Even for substrate **13d**, which lacked an *N*-alkyl substituent, the catalyst **B** afforded the cyclized product **17d** in a moderate yield (entry 7).

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