

Diversity-Oriented Synthesis of Azapeptides with Basic Amino Acid Residues: Aza-Lysine, Aza-Ornithine, and Aza-Arginine

Mariam Traoré, Ngoc-Duc Doan, and William D. Lubell*

Département de Chimie, Université de Montréal, C.P. 6128, Succursale Centre-Ville, Montréal, Québec, Canada H3C 3J7

Supporting Information

ABSTRACT: Aza-peptides with basic amino acid residues (lysine, ornithine, arginine) and derivatives were synthesized by an effective approach featuring alkylation of a hydrazone-protected aza-glycine residue with α -bromo ω -chloro propane and butane to provide the corresponding alkyl chloride side chains. Displacement of the chloride with azide and various amines gave entry to azaOrn, azaLys, and azaArg containing peptides as demonstrated by the solution and solid-phase syntheses of 29 examples, including an aza-library of Growth Hormone Releasing Peptide-6 analogs.

B asic amino acid residues, such as lysine and arginine, function in numerous biological processes including posttranslational modifications, transport across membranes, and as enzymatic cleavage sites.³ For example, acylation and methylation of lysine residues of histone proteins alter chromatin structure and gene regulation. 1,4 Cell-penetrating peptides, 5,6 such as Tat and penetratin, are composed of multiple basic residues. The protease trypsin cleaves peptides specifically at the C-terminal of lysine and arginine residues. Peptide mimics with constrained conformations and chemical modifications at lysine and arginine residues are thus desirable tools for studying and regulating such phenomena.8-14

Azapeptides employ the electronic constraints of a semicarbazide residue to rigidify peptide geometry in favor of turn conformations, with the added benefit of enhanced stability against protease degradation.¹⁵ Efforts to prepare basic azaamino acid residues have typically involved building block synthesis in solution, 16-18 which restricts analog generation. Recently, our laboratory reported the synthesis of aza-Lys peptides by a route featuring the copper-catalyzed addition of Mannich reagents to an aza-propargylglycine residue. 19 This socalled A³ reaction²⁰ was amenable to solid phase to provide rigid lysine analogs of type C (Figure 1) possessing a variety of tertiary amines on the side chain. Moreover, hydrogenation of the triple bond gave access to the Z-olefin analogs D. In principle, further reduction of the triple bond may give access to aza-lysine analogs of type B. The A³-route does however have limitations in requiring typically secondary amines, a four-carbon side chain length, and extra steps to prepare saturated analogs.

In our efforts to make constrained N-aminoimidazolidin-2-one peptides,²¹ we have recently explored the application of 1,2dibromoethane in the alkylation of both urea nitrogen of a

$$\begin{array}{c} \text{NH}_2 \\ \text{NH}_2 \\$$

Figure 1. (A) Ornithine and lysine in native peptide, and aza-lysine analogs possessing (B) saturated (C) alkyne and (D) Z-olefin side chains.

semicarbazone to prepare the heterocycle. Considering the rate of nitrogen alkylation, we perceived that the more acidic semicarbazone nitrogen could be alkylated with an α -bromo- ω -chloro alkane to furnish an alkyl chloride side chain. Subsequent displacement of the chloride using a variety of amines may then provide access to a diverse array of aza-residues with basic side chains. This strategy has now been realized using

Received: June 2, 2014 Published: June 24, 2014 Organic Letters Letter

both solution- and solid-phase methods, which have provided a wide diversity of peptide analogs possessing basic aza-residues.

Aza-lysine analogs were pursued initially by a solution-phase method (Scheme 1) featuring alkylation of benzhydrylidene-

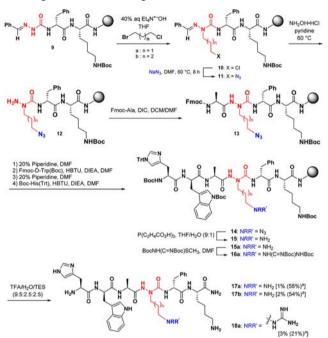
Scheme 1. Solution-Phase Synthesis of Aza-Lys and Aza-Orn Dipeptides

azaGly-Phe-OtBu (1) and benzhydrylidene-azaGly-Phe-NiPr (2).²² Employing tetraethylammonium hydroxide (120 mol %, as a 40% aqueous solution) in THF, ester 1 and amide 2 were alkylated with 1-bromo-3-chloropropane and 1-bromo-4-chlorobutane to provide the chloroalkyl semicarbazones 3-5, which were converted into aza-Orn and aza-Lys dipeptides by chloride displacement with different amines. Racemization of esters 3 and 4 (up to 15%) was however detected from the alkylation of tertbutyl ester 1 at room temperature, but was avoided by decreasing the temperature to 0 °C. On the other hand, no loss of configuration was observed using amide 2 even at rt. Chloroalkyl semicarbazones 3-5 were then converted to analogs of aza-Orn and aza-Lys 6-8 from reactions with sodium azide, primary and secondary amines, and trimethylamine in DMF at 60 °C (Scheme 1). The relatively polar final dipeptides were purified by preparative HPLC on a C18 column, except for compounds 6a and 6d, which were purified by flash chromatography on silica gel.

A method for the synthesis of azapeptides possessing basic azaamino acid residues was next developed on solid phase. Growth Hormone Releasing Peptide-6 (GHRP-6, His-D-Trp-Ala-Trp-D-Phe-Lys-NH₂), a cluster of differentiation 36 receptor (CD36) ligand,²³ was chosen for this study, because of its interesting biological activity and challenging chemical structure. In particular, [azaLys]-GHRP-6 analogues were targeted, because of the presence of two lysine residues in the related peptide EP80317 (Haic-2Me-Trp-D-Lys-Trp-D-Phe-Lys-NH₂), which has exhibited CD36 selectivity. Azapeptide analogs of GHRP-6 were thus targeted possessing respectively basic azaresidues at the 3-, 4-, and 6-positions.

Initially, the set of [azaOrn⁴]-, [azaLys⁴]-, and [azaArg⁴]-GHRP-6 analogs was pursued in order to demonstrate a solid-phase protocol for preparing aza-derivatives of the common naturally occurring basic amino acids. On rink amide resin, benzylidene-azaGly-D-Phe-Lys(Boc) 9 (Scheme 2) was synthe-

Scheme 2. Synthesis of [AzaLys⁴]-, [AzaOrn⁴]-, and [AzaArg⁴]-GHRP-6 Analogs



"Isolated yield calculated from resin loading and crude purity ascertained by LC-MS.

sized as previously described.²⁶ Alkylation of resin **9** was performed with 1-bromo-3-chloropropane and 1-bromo-4-chlorobutane, respectively, at room temperature using tetraethylammonium hydroxide (200 mol % of a 40% aqueous solution) in THF to provide the chloroalkyl resins **10a** and **10b**. Conversion was assessed to be complete by treating aliquots of resin **10** with TFA/H₂O/TES (9.5:0.25:0.25) and examination of the residue after resin filtration by LC-MS. Resins **10** were respectively treated with sodium azide in DMF to provide the corresponding azides **11a** and **11b**. The semicarbazone was removed using a 1.5 M solution of hydroxylamine hydrochloride in pyridine, and the resulting semicarbazides **12** were coupled to Fmoc-Ala by way of its symmetric anhydride using diisopropyl-carbodiimide (DIC) to provide azatetrapeptides **13**.²⁶ Elongation of the sequence was then performed using conventional Fmoc-based solid-phase peptide synthesis (SPPS).²⁷

Azides 14 were chemoselectively reduced with tris(2-carboxy)ethylphosphine (TCEP) to provide the corresponding amines 15. Resin cleavage was then performed using a cocktail of TFA/H₂O/TES (9.5:0.25:0.25) to provide respectively [azaOrn⁴]- and [azaLys⁴]-GHRP-6 (17a and 17b) in 58% and 54% crude purities. Purification of the crude azapeptides by

Organic Letters Letter

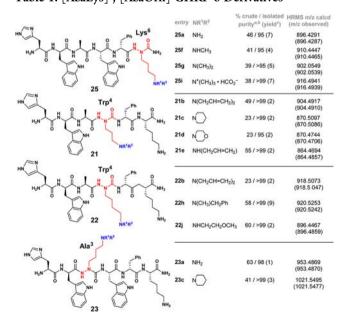
HPLC on a C18-column gave 1% and 2% overall yields of 17a and 17b. Alternatively, amine 15a was treated with *N,N*-bis(Boc)-*S*-methylisothiourea in DMF at rt for 12 h to provide [azaArg⁴]-GHRP-6 resin 16a, ³⁰ which was cleaved from the resin using the same cocktail to provide [azaArg⁴]-GHRP-6 (18a) in 21% crude purity, and 3% overall yield after purification by HPLC.

Chloroalkyl resins 10 were subsequently employed to prepare a library of diverse [azaLys]-GHRP-6 derivatives (Scheme 3).

Scheme 3. Synthesis of [AzaLys⁴]-GHRP-6 Derivatives

Employing various primary and secondary amines to displace chlorides **10a** and **10b** in DMF, substituted azaOrn and azaLys resins **19** and **20** were respectively synthesized. In the case of secondary amine analogs, the amine was subsequently protected as the corresponding *N*-Boc derivative by treating the resin with di-*tert*-butyldicarbonate in dichloromethane. The semicarbazone protecting group was removed, with the peptides elongated and cleaved from resin as previously described above. Final [azaOrn⁴] and [azaLys⁴]-GHRP-6 analogs **21** and **22** were obtained in 23–63% crude purities and isolated in 2–9% overall yields after purification by HPLC (Table 1; see Supporting Information for

Table 1. [AzaLys]-, [AzaOrn]-GHRP-6 Derivatives



^aCrude purity ascertained by LCMS analysis at 214 nm using $\rm H_2O$ (0.1% FA)/MeOH (0.1% FA) and $\rm H_2O$ (0.1% FA)/MeCN (0.1% FA) as eluents. ^bIsolated purity ascertained in same manner as crude purity. ^cIsolated yields calculated from resin loading.

details). Employing a similar synthetic strategy [azaLys³]-GHRP-6 analogues **23** were prepared from benzylidene-azaGly-Trp(Boc)-p-Phe-Lys(Boc) rink amide resin (Table 1).³¹

Finally, [azaLys⁶]-GHRP-6 analogs **25** were prepared using methyl-, dimethyl-, and trimethylamines (Scheme 4). Benzyli-

Scheme 4. Synthesis of [azaLys⁶]-GHRP-6 Analogues

dene-azaGly rink amide resin 24 was obtained by removal of the Fmoc protection from rink amide resin, and treatment of the amine resin with 4-nitrophenyl 2-(diphenylmethylidene)carbazate, which was prepared from the reaction of benzophenone hydrazone with *p*-nitrophenylchloroformate (Scheme 4). LCMS analysis of a cleaved resin aliquot of 24 showed 98% conversion to the N-(diphenylmethylidene)glycinamide product possessing the desired molecular ion. To introduce the side chains, resin 24 was alkylated with 1-bromo-4-chlorobutane and tetraethylammonium hydroxide (120 mol %, as a 40% aqueous solution) in THF, followed by chloride conversion into an amine. In the case of the methylamine analog, the amine was subsequently protected as the corresponding N-Boc derivative before peptide elongation. After removal of the semicarbazone group, azapeptides were elongated and cleaved from the resin as previously described.

In summary, 16 new GHRP-6 analogs were synthesized possessing aza-residues with different basic amine side chains (Scheme 2 and Table 1). This method has provided access to azaLys, azaOrn, and azaArg peptide sequences from a common diversity-oriented strategy featuring alkylation of an azaGly residue with an α -bromo- ω -chloroalkane followed by displacement of the resulting chloride with various amines. With this set of GHRP-6 analogs possessing basic aza-residues in hand, their impact on CD36 affinity and activity is currently being examined and will be reported in due time. Considering the power of this method for preparing azapeptides possessing diverse basic aza-residues, this approach should find significant applications in the study of various events featuring the post-translational modification and activity of lysine and arginine containing peptide structures.

ASSOCIATED CONTENT

Supporting Information

Experimental procedures and characterization data. This material is available free of charge via the Internet at http://pubs.acs.org.

Organic Letters Letter

AUTHOR INFORMATION

Corresponding Author

*E-mail: william.lubell@umontreal.ca.

Notes

The authors declare no competing financial interest.

ACKNOWLEDGMENTS

This research was supported by the Natural Sciences and Engineering Research Council of Canada (NSERC), the Canadian Institutes of Health Research (Canada-UK partnership on Antibiotic Resistance), the Ministère du développement économique de l'innovation et de l'exportation du Quebec (#878-2012, Traitement de la dégénerescence maculaire), and Amorchem.

REFERENCES

- (1) Cole, P. A. Nat. Chem. Biol. 2008, 4, 590-597.
- (2) Mitchell, D. A.; Farh, L.; Marshall, T. K.; Deschenes, R. J. J. Biol. Chem. 1994, 269, 21540–21546.
- (3) Sun, A.-Q.; Luo, Y.; Backos, D. S.; Xu, S.; Balasubramaniyan, N.; Reigan, P.; Suchy, F. *J. Mol. Pharmacol.* **2013**, 83, 1078–1086.
- (4) Allfrey, V. G.; Faulkner, R.; Mirsky, A. E. Proc. Natl. Acad. Sci. U.S.A. 1964, 51, 786–794.
- (5) Henriques, S. T.; Melo, M. N.; Castanho, M. A. R. B. *Biochem. J.* **2006**, 399, 1–7.
- (6) Madani, F.; Lindberg, S.; Langel, U.; Futaki, S.; Graeslund, A. J. Biophys. **2011**, 414729–414739.
- (7) Fittler, H.; Avrutina, O.; Glotzbach, B.; Empting, M.; Kolmar, H. Org. Biomol. Chem. **2013**, *11*, 1848–1857.
- (8) Sajjadi, Z.; Lubell, W. D. J. Pept. Res. 2005, 65, 298-310.
- (9) Feng, Z.; Lubell, W. D. J. Org. Chem. 2001, 66, 1181-1185.
- (10) Ganorkar, R.; Natarajan, A.; Mamai, A.; Madalengoitia, J. S. J. Org. Chem. **2006**, 71, 5004–5007.
- (11) Goswami, R.; Moloney, M. G. Chem. Commun. 1999, 2333-2334.
- (12) Wong, M. L.; Guzei, I. A.; Kiessling, L. L. Org. Lett. 2012, 14, 1378–1381.
- (13) Huang, Z. P.; Du, J. T.; Su, X. Y.; Chen, Y. X.; Zhao, Y. F.; Li, Y. M. *Amino Acids* **2007**, *33*, 85–89.
- (14) Chi, H.; Islam, M. S.; Nsiama, T. K.; Kato, T.; Nishino, N. *Amino Acids* **2014**, *46*, 1305–1311.
- (15) Proulx, C.; Sabatino, D.; Hopewell, R.; Spiegel, J.; Garcia-Ramos, Y.; Lubell, W. D. Future Med. Chem. **2011**, 3, 1139–1164.
- (16) Balliano, G.; Milla, P.; Giordano, C.; Gallina, C.; Coletta, M.; Menegatti, E.; Rizzi, M.; Bolognesi, M.; Ascenzi, P. *Biochem. Biophys. Res. Commun.* **1996**, 225, 557–561.
- (17) Freeman, N. S.; Tal-Gan, Y.; Klein, S.; Levitzki, A.; Gilon, C. J. Org. Chem. 2011, 76, 3078–3085.
- (18) Boeglin, D.; Lubell, W. D. J. Comb. Chem. 2005, 7, 864-878.
- (19) Zhang, J.; Proulx, C.; Tomberg, A.; Lubell, W. D. Org. Lett. 2014, 16, 298–301.
- (20) Peshkov, V. A.; Pereshivko, O. P.; Van der Eycken, E. V. Chem. Soc. Rev. 2012, 41, 3790–3807.
- (21) Doan, N.-D.; Hopewell, R.; Lubell, W. D. Org. Lett. **2014**, 16, 2232–2235.
- (22) Garcia-Ramos, Y.; Lubell, W. D. J. Pept. Sci. 2013, 19, 725-729.
- (23) Proulx, C.; Picard, E.; Boeglin, D.; Pohankova, P.; Chemtob, S.; Ong, H.; Lubell, W. D. *J. Med. Chem.* **2012**, *55*, 6502–6511.
- (24) Marleau, S.; Harb, D.; Bujold, K.; Avallone, R.; Iken, K. A.; Wang, Y.; Demers, A.; Sirois, M. G.; Febbraio, M.; Silverstein, R. L.; Tremblay, A.; Ong, H. FASEB J. 2005, 19, 1869–1871.
- (25) Demers, A.; McNicoll, N.; Febbraio, M.; Servant, M.; Marleau, S.; Silverstein, R.; Ong, H. Biochem. J. 2004, 382, 417–424.
- (26) Sabatino, D.; Proulx, C.; Klocek, S.; Bourguet, C. B.; Boeglin, D.; Ong, H.; Lubell, W. D. *Org. Lett.* **2009**, *11*, 3650–3653.
- (27) Lubell, W. D.; Blankenship, J. W.; Fridkin, G.; Kaul, R. Sci. Synth. **2005**, *21*, 713–809.

(28) Saneyoshi, H.; Ochikubo, T.; Mashimo, T.; Hatano, K.; Ito, Y.; Abe, H. *Org. Lett.* **2014**, *16*, 30–33.

- (29) Faucher, A.-M.; Grand-Maitre, C. Synth. Commun. 2003, 33, 3503-3511.
- (30) Gers, T.; Kunce, D.; Markowski, P.; Izdebski, J. Synthesis 2004, 37-42.
- (31) Sabatino, D.; Proulx, C.; Pohankova, P.; Ong, H.; Lubell, W. D. J. Am. Chem. Soc. **2011**, 133, 12493–12506.