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Asymmetric Synthesis of Functionalizable Type II β -Turn-Inducing α -Amino Acid Building Blocks

Wenzheng Gao, Jiaxin Han, Sophie Greaves, and Joseph P. A. Harrity*



ABSTRACT: Peptidomimetics are emerging as a promising class of potent and selective therapeutics. Among the current approaches to these compounds, the utilization of constrained lactams is a key element in enforcing the active peptide conformation, and the development of efficient and stereocontrolled methods for generating such lactam building blocks is an important objective. Current methods typically rely on the elaboration of existing α -amino acids, and in so doing, the side chain is sacrificed during the ring-forming process. We report a new asymmetric approach to lactam-constrained α -amino acid building blocks bearing a range of polar and hydrophobic side chains. The chemistry is amenable to rapidly generating di- and tripeptides, and the potential for these lactams to stabilize type II β -turns is demonstrated in the synthesis of the melanocyte-inhibiting factor peptidomimetic.

espite mediating a plethora of vital biological processes in living organisms, peptides remain a challenging class of drug targets due to their poor cell penetration, proteolytic instability, and unfavorable pharmacokinetics. Among the approaches employed to address these shortcomings,¹ the incorporation of conformational constraints has the potential to enhance stability and, if the constraint can mimic a bioactive conformation, improve potency.² With these goals in mind, lactam-bridged peptides were introduced by Freidinger, and these have been proven to deliver an effective class of peptidomimetics as they stabilize type II β -turns.³ For example, and as shown in Figure 1a, an analogue of the luteinizing hormone-releasing hormone containing a γ -lactam as a conformational constraint showed improved agonist activity as compared to the parent hormone due to the stabilization of a bioactive conformation containing a β -turn. There are several strategies for the synthesis of Freidinger lactams, including cyclocondensation^{3,4} and ring-closing metathesis,⁵ but in general, they provide motifs that lack functionality, which limits their further elaboration downstream (Figure 1b). In addition, current approaches rely almost exclusively on the stereospecific elaboration of available α -amino acids, and de novo asymmetric routes to functionalizable constrained amino acid building blocks are almost unknown. Indeed, in many of these cases, the side chain is sacrificed during the ring-forming process. We envisaged that the allylation of azlactones using an in situ generated Pd zwitterion reported by our group⁶ and others⁷ would allow the ready assembly of lactam monomers that were appropriately armed for incorporation into peptide

motifs. As shown in Figure 1c, these compounds would be amenable to N to C homologation by lactam alkylation, peptide coupling, and tagging at the olefin moiety. We report herein our progress toward the enantioselective synthesis of these constrained amino acids and their incorporation into small peptide arrays.

We began our studies by investigating the key lactamforming transformation, and our results are shown in Figure 2a. Pleasingly, carbamate 2a and 6-substituted analogue 2b were efficiently transformed in to lactams 3 and 4a, respectively. Notably, 4a is formed at the expense of isomer 4b with excellent E/Z selectivity, an observation we attribute to a combination of minimization of steric control and allylic strain, as highlighted in I. At the outset of this work, we recognized the importance of establishing a method that would access the constrained amino acid building blocks with synthetically useful enantiomeric ratios. On the basis that the Pd-catalyzed asymmetric allylation and benzylation of azlactones are successful using the Trost ligand series,⁸ we used these ligands to screen and optimize the enantioselectivity of this transformation. As shown in Figure 2b, we identified L4 as

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Figure 1. Synthetic approaches to Freidinger lactams.



Figure 2. Development of the lactam-forming transformation and optimization of the reaction enantioselectivity.

providing the most selective catalyst system, and further optimization led to conditions that generated 3 in both high

We next set out to explore the scope of the method for generating simple lactam-constrained mimics of representative natural α -amino acids, and our results are summarized in Table 1. In general, we were able to incorporate the key 2-



Table 1. Scope of the Enantioselective Lactam Synthesis^a

^aX-ray structures show thermal ellipsoids drawn at the 50% probability level.

aminomethyl allyl fragment with enantiomeric ratios of 90:10 or better, and these could be smoothly transformed into the corresponding lactams after removal of the Boc group with TFA. The methodology showed good generality, delivering lactam mimics bearing both polar and hydrophobic side chains. The exception was 12, which underwent the asymmetric allylation step with both a poor yield and poor enantioselectivity. In addition, we were able to crystallize lactams 3 and 5, and both showed the *R* configuration at the newly generated stereogenic center. The remaining compounds were assigned by inference. In addition, we recognized that this methodology offered the opportunity to directly prepare FL_x-Gly (FL, Freidinger lactam; x, α -amino acid mimic label) dipeptides by replacing the azlactone Ph substituent at C2 with an aminomethyl group. Pleasingly, subjecting azlactones 13-15 to our optimized conditions generated FL_{Phe}-Gly dipeptides with or without a Cbz protecting group and Cbz-protected FL_{Leu}-Gly, both with high enantiomeric ratios.⁹

As shown in Figure 3, we have applied the model developed by Lloyd-Jones and Norrby¹⁰ to put forward a rationale for the observed stereochemical outcome of the allylic alkylation process. The enantioselective step is mediated by a H-bond interaction between the azlactone enolate and the ligand N-H that allows the addition to take place with the aromatic substituent oriented away from the catalyst "roof". The

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Figure 3. Proposed origin of the reaction enantioselectivity.

alternative mode of addition places both substituents at C2 and C4 in the proximity of the ligand structure.

While the substrates prepared in Table 1 verified that the basic methodology could allow access to enantiomerically enriched constrained amino acid building blocks, we were concerned that the generation of the products as benzamides would hamper their further elaboration due to the difficulties associated with hydrolyzing this group. In this context, Connon and co-workers designed a family of azlactones that offer mild hydrolysis protocols via formation of phthalimide intermediates and set out to investigate whether this chemistry could offer a useful solution to this problem.¹¹ Pleasingly, as shown in Table 2, we were able to transform azlactones **1m**–



1q into the corresponding free amine building blocks 16-20, respectively, through a simple three-step sequence, generating the products in good yield and enantiocontrol. Interestingly, the product derived from glutamic acid-based azlactone 1r underwent further cyclization when subjected to this sequence, generating functionalized spirolactam 21.

We next took the opportunity to explore the functionalization of the alkene. As shown in Figure 4, we protected the amino group in 16 and carried out a homologation to generate ^tBu-Gly-FL_{Leu}-Boc 22. Oxidative cleavage of the alkene followed by diastereoselective reduction¹² provided an alcohol unit that was readily elaborated to the corresponding propargyl ether 24, offering a means for these lactams to be easily tagged to other molecules through "click" chemistry. In addition, 22 also underwent efficient epoxidation and hydrogenation reactions, albeit with modest diastereocontrol.



Figure 4. Reaction conditions: ^aBoc₂O, Et₃N, THF/H₂O (1:2); ^b(i) LHMDS; (ii) BrCH₂CO₂^tBu; ^cRuCl₃ (30 mol %), NaIO₄, MeCN/CH₂Cl₂/H₂O (1:1:2); ^dK-selectride, THF; ^cHCCCH₂Br, ^aBu₄NSO₄, NaOH, toluene; ^fOxone, NaHCO₃, acetone/H₂O; ^gH₂/Pd/C.

Finally, to demonstrate the applicability of this chemistry to the generation of peptidomimetics, we targeted the synthesis of a constrained analogue of melanocyte-inhibiting factor (MIF-1).¹³ MIF-1 is a hypothalamic neuropeptide derived endogenously by cleavage of the hormone oxytocin. This tripeptide displays a range of bioactivities and has been studied for the treatment of Parkinson's disease as well as for its antidepressant and nootropic activities. As shown in Figure 5,



Figure 5. ^{*d*}EDCI, HOAt, ^{*i*}Pr₂NEt, Boc-L-Pro-OH, CH₂Cl₂; ^{*b*}LHMDS, ICH₂C(O)NH₂, DMF; ^{*c*}TFA. ^{*d*}Torsional angles in parentheses are those for an ideal type II β -turn. The X-ray structure shows thermal ellipsoids drawn at the 50% probability level.

we derivatized FL_{Leu} 5 toward MIF-1 analogue 28 within a short synthetic sequence. During this synthesis, we were able to grow suitable crystals of 27 for X-ray crystallographic analysis, and this compound showed the 10-membered glycinamide–proline hydrogen bond and dihedral angles that are consistent with type II β -turns.¹⁴ Notably, the H-bond interaction between the C-terminal glycinamide hydrogen and the prolyl carbonyl oxygen has been observed in MIF-1 both in solution and in the solid state,¹⁵ highlighting that these lactam mimics can deliver peptides that maintain key secondary structural features.

We developed a de novo asymmetric route to lactamconstrained α -amino acid building blocks bearing a range of polar and hydrophobic side chains. As well as enforcing type II β -turn conformations, these intermediates are armed with an exocyclic alkene that provides a handle for ligation via "click" chemistry providing a platform for a range of tailored applications. Studies in this area are ongoing and will be reported in due course.

ASSOCIATED CONTENT

Data Availability Statement

The data underlying this study are available in the published article and its Supporting Information.

3 Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acs.orglett.3c02376.

Synthesis and characterization details, ¹H and ¹³C NMR spectra, FTIR and HRMS data, HPLC traces, and X-ray crystal structure data (PDF)

Accession Codes

CCDC 2224755 and 2224761–2224762 contain the supplementary crystallographic data for this paper. These data can be obtained free of charge via www.ccdc.cam.ac.uk/data_request/ cif, or by emailing data_request@ccdc.cam.ac.uk, or by contacting The Cambridge Crystallographic Data Centre, 12 Union Road, Cambridge CB2 1EZ, UK; fax: +44 1223 336033.

AUTHOR INFORMATION

Corresponding Author

Joseph P. A. Harrity – Department of Chemistry, University of Sheffield, Sheffield S3 7HF, United Kingdom; orcid.org/ 0000-0001-5038-5699; Email: j.harrity@shef.ac.uk

Authors

- Wenzheng Gao Department of Chemistry, University of Sheffield, Sheffield S3 7HF, United Kingdom
- Jiaxin Han Department of Chemistry, University of Sheffield, Sheffield S3 7HF, United Kingdom
- Sophie Greaves Department of Chemistry, University of Sheffield, Sheffield S3 7HF, United Kingdom

Complete contact information is available at: https://pubs.acs.org/10.1021/acs.orglett.3c02376

Notes

The authors declare no competing financial interest.

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