Graphical Abstract

Development of aspartic acid ligation for peptide cyclization derived from serine/threonine ligation

Ci Xu, Jianchao Xu, Han Liu, Xuechen Li*

Department of Chemistry, State Key Laboratory of Synthetic Chemistry, The University of Hong Kong, Hong Kong SAR, China



Based on a mechanism analogous to the serine/threonine ligation, the aspartic acid ligation, which is facilitated by the γ -amino alcohol based ligation and oxidation, is developed and applied to the synthesis of cyclic peptides. The γ -hydroxyl group triggers the ring-chain tautomerization via a 6-*endo-trig* process, while the δ -hydroxyl group facilitates the oxidative cleavage of the vicinal diol to give carboxylic acid.

This part will be used for Graphical Contents.

Communication (heading)

Development of aspartic acid ligation for peptide cyclization derived from serine/threonine ligation

Ci Xu^{a, b}, Jianchao Xu^{a, b}, Han Liu^a, Xuechen Li^{a, *}

^aDepartment of Chemistry, State Key Laboratory of Synthetic Chemistry, The University of Hong Kong, Hong Kong SAR, 999077, China ^bEqual contribution

ARTICLE INFO

Article history: Received Received in revised form Accepted Available online

Keywords: Aspartic acid ligation Peptide cyclization Serine/threonine ligation Ring-chain tautomerization Acyl transfer Selective oxidation

ABSTRACT

Based on a mechanism analogous to the serine/threonine ligation, the aspartic acid ligation, which is facilitated by the γ -amino alcohol based ligation and oxidation, is developed and applied to the synthesis of cyclic peptides. The γ -hydroxyl group triggers the ring-chain tautomerization via a 6-*endo-trig* process, while the δ -hydroxyl group facilitates the oxidative cleavage of the vicinal diol to give carboxylic acid.

As an important class of molecular constructs from both natural and synthetic sources, cyclic peptides attract the attention of the researchers from synthetic chemistry and medicinal chemistry. Compared with the corresponding linear peptides, cyclic peptides are prone to show improved binding affinity to protein targets and higher stability against degradation, benefiting from the restricted conformations [1-3]. Compared with the linear peptides which can be readily and even automatically assembled via solid phase peptide synthesis (SPPS), the synthesis of cyclic peptides pose additional challenges in the head-to-tail macrolactamization of the linear precursors with protected side chains [4]. These challenges include racemization of the Cterminal amino acid residues in the linear precursors and the undesired oligomerization, which have to be overcome by extensive optimization of the coupling conditions (reagents, bases, concentrations, etc.), variation of the cyclization sites, and even incorporation of unnatural structure units [5].

As an alternative strategy to the traditional head-to-tail peptide cyclization based on coupling, the chemical peptide ligation with the feature of site specific formation of the peptide linkage between side chain unprotected fragments without racemization

* Corresponding author.

E-mail address: xuechenl@hku.hk

have been adapted to the synthesis of cyclic peptides [6]. Since the first publication of native chemical ligation by Kent et al. in 1994 [7], several peptide ligation strategies, including native chemical ligation (NCL) [8-12], pseudoproline ligation [13-14], α-ketoacid-hydroxylamine (KAHA) ligation [15-17], traceless Staudinger ligation [18], 2-formylthiophenol ligation [19] and enzyme (subtiligase [20], butelase [21-23], sortase [24]) catalyzed ligations, have been used in the synthesis of cyclic peptides and even cyclic proteins. In 2010, our group developed a novel peptide ligation, termed serine threonine ligation (STL), between the peptide C-terminal salicylaldehyde esters and peptides with N-terminal serine or threonine residues (Scheme 1a) [25-30]. This ligation works through the imine capture between the two peptide fragments, followed by the side chain nucleophile (hyroxyl group in Ser or Thr) triggered ring-chain tautomerization and subsequent O-to-N [1,5] acyl transfer. The so-obtained N,O-benzylidene acetal is acid labile, which can be easily cleaved under mild condition to give the product with native peptide linkage. This new ligation has been applied in the synthesis of linear peptide drugs [31], proteins with posttranslational modifications [32-34], as well as cyclic

peptides/depsipeptides [35] (daptomycin [36-37], teixobactin [38-39], yunnanin C analogues [40], cyclomontanin B [41]) with variable ring sizes (from highly strained cyclic tetrapeptides [42-43] to highly flexible decapeptides). As an extension of our former work, we conceived of a new ligation method for peptide cyclization, based on the analogous side chain nucleophile triggered ring-chain tautomerization and O-to-N acyl transfer. As illustrated in Scheme 1b, we designed a aspartic acid ligation, facilitated by the γ -amino alcohol based ligation and oxidation. Compared with the serine/threonine ligation, the imine capture and the O-to-N acyl transfer are in common, while the side chain nucleophile triggered ring-chain tautomerization works in a 6endo-trig manner rather than a 5-endo-trig manner. To facilitate further transformation, a vicinal diol functionality is introduced onto the N-terminal residue instead of a single hydroxyl group, which can be cleaved under mild oxidation condition to give a carboxylic acid residue. Through this process, a native aspartic acid residue will be formed at the ligation site. Herein, we would like to document how we develop this ligation and apply it to cyclic peptide synthesis.



Scheme 1. Design of the aspartic acid ligation for peptide cyclization facilitated by γ -amino alcohol based ligation and oxidation.

To test our idea, we designed a Boc protected dihydroxylated amino acid building block (abreviated as Dh) **7** for solid phase peptide synthesis, which contains the vicinal diol functionality in the side chain. As illustrated in Figure 2, the synthesis started from commercial available Boc protected L-aspartic acid monomethyl ester **1**, which was transformed into side chain thioester **2**. After the Fukuyama reduction [44], the aldehdye **3** was directly trapped by phosphonium ylide to give alkene **4**. The methyl ester in **4** was selectively hydrolyzed in the presence of *tert*-butyl ester under mild condition, then the vicinal diol functionality was installed on **5** via Osmium(VIII) catalyzed *cis*dihydroxylation [45]. An inseparable mixture of diastereomers **6** was obtained in a moderate yield, and then the diol was further protected by acetonide to give **7**. Details of this synthesis can be found in Supporting Imformation.



a) EISH, DCC, DMAP, DCM, 90%. b) PdC, EI₂SiH, DCM. c) /BuOOCCH=PPh₃, THF, 87% over 2 steps. d) LiOH (aq), THF/H₂O, 85%. e) 4-Methylmorpholine N-oxide monohydrate, K₂OsO₄-2H₂O, acetone/H₂O, 44%. f) 2.2-Dimethoxypropane, BF₃-OE₁₂, acetone, 81%. Scheme 2. Synthesis of dihydroxylated amino acid building block (*Dh*).

After obtaining the *Dh* building block **7**, we went forward to the preparation of the peptide C-terminal salicylaldehyde (SAL) esters, the linear precursors for Ser/Thr ligation. To achieve this, the Boc-SPPS based method was used [42]. As illustrated in Scheme 3, from AM resin, the 2-hyroxylcinnamoyl linker (the surrogate of SAL) was installed to give **8**, followed by the assembly of the peptide sequence. After finishing the desired sequence with the building block **7** installed onto the *N*-terminus, the resin bounded peptide **9** was subjected to the global deprotection. The SAL ester was generated from the released peptide via ozonolysis, and the product **10a-e** were obtained after preparative HPLC purification in 25-42% yields (calculated based on resin loading).



Scheme 3. Synthesis of peptide C-terminal SAL ester with N-terminal *Dh* via Boc-SPPS.



Scheme 4. HPLC monitoring of the aspartic acid ligation facilitated peptide cyclization.

With the desired peptide SAL esters **10a-e** in hand, we started to test this peptide cyclization. As an example, the peptide **10a** was dissolved in the pyridine/acetic acid buffer to form a 20 mM

solution, and the progress of the ligation was monitored by LC-MS. After screening of the pyridine/acetic acetic buffer compositions (from 9 : 1 to 1 : 6, molar ratio), the 9 : 1 buffer was found to be optimal. The HPLC traces of the reaction under the optimized condition were shown in Scheme 4. The full conversion of the peptide SAL ester **10a** was observed after 15.5 hrs, and the *N*,*O*-benzylidene acetal intermediate (broaden peak) was formed as a mixture of diastereomers (due to the mixture of *Dh* building block) via the expected 6-*endo-trig* tetrahydro-1,3-oxazine formation and *O*-to-*N* acyl transfer. After blowing off the pyridine/acetic acid by flow gas, the crude intermediate was then treated with TFA/H₂O/TIPS 95/2.5/2.5 (volumn ratio) cocktail to cleave the benzylidene acetal, and the desired cyclic peptide **11a** with *Dh* at the ligation site was formed, also as mixture of diastereomers.

To achieve the desired aspartic acid ligation, we optimized the site-selective oxidation of the diol side chain of Dh residue. After extensive screening of the reagents and conditions (see Supporting Information for details), to our delight, the formation of the aspartic acid from Dh residue was realized in one-pot manner under the treatment of the cyclic peptide 11a by the mixture of NaIO₄, NaClO₂ and H_2O_2 in MeCN/H₂O. As an example, the crude ligation products 11a (1.0 equiv) was mixed with NaIO₄ (15.0 equiv) and NaClO₂ (10.0 equiv) in CH₃CN/H₂O (1/1, v/v) at room temperature. After the addition of H₂O₂ (50 wt.% in H₂O, 20.0 equiv), gas was evolved and a white precipitate formed in the stirred yellow solution. The oxidation reaction was completed within 10 min, as indicated by LC-MS monitoring. The product 12a was obtained in 18% yield over 3 steps (calculated based on the purified SAL ester 10a) after HPLC purification and lyophilization.



Scheme 5. Substrate scope of the aspartic acid ligation facilitated peptide cyclization.

To evaluate the scope of this new peptide cyclization method, we tested the reaction on a series of peptide SAL esters with varied sequences and chain length. As shown in Scheme 5, for substrates **10a-e**, different pyridine/acetic acid ratios (3 : 1 or 9 :

1) were found to be optimal for the ligation step. Pyridine was used as the dominant composition in general, which was in accordance with our former observations in cyclic tetrapeptide synthesis via serine/threonine ligation [42]. Val or less bulky Gly were tolerated. For the peptide **10d** without turn-inducing Pro residue, longer reaction time (up to 4 days) was need to achieve full conversion. After the acidolysis of the N,O-benzylidene acetal intermediates, the crude cyclic peptides **11a-e** were subjected to the optimized oxidation condition to give the octapeptides **12a-d** and heptapeptide **12e** in 13-26% yields after HPLC purification.

In conclusion, we developed a novel aspartic acid ligation derived from serine/threonine ligation, in which the y-amino based ligation worked through a ring-chain alcohol tautomerization via 6-endo-trig process and subsequent O-to-N [1,5] acyl transfer. The oxidative cleavage of the diol into carboxylic acid was functionality realized bv NaIO₄/NaClO₂/H₂O₂ in water containing system in one-pot manner. The method was successfully used in the synthesis of cyclic peptides with aspartic acid residue in the sequences. Nevertheless, this γ -amino alcohol based ligation was less efficient as compared to the serine/threonine ligation. Both ligation methods proceed through ring-chain tautomerization step and acyl transfer step. According to the Baldwin empirical rule, the γ -amino alcohol based ligation giving a 6-endo-trig ringchain tautomerization is more favored than the 5-endo-trig cyclization from serine/threonine ligation using β -amino alcohol. Thus, it is very likely that, during the acyl transfer step, the 5-6 fused bicyclic transition state in serine/threonine ligation is much more favored than the 6-6 fused bicyclic transition state in the γ amino alcohol based ligation. The current effort to apply for this ligation in protein chemical synthesis is ongoing in our laboratory.

Acknowledgment

This work was supported by the Research Grants Council (17309616, C6009-15G) of Hong Kong, The National Science Foundation of China (21672180, 91753101), the Area of Excellence Scheme of the University of Grants Committee of Hong Kong (AoE/P-705/16).

References

- Y.S. Ong, L. Gao, K.A. Kalesh, et al., Recent advances in synthesis and identification of cyclic peptides for bioapplications, Curr. Top. Med. Chem. 17 (2017) 2302-2318.
- [2] A.T. Bockus, C.M. McEwen, R.S. Lokey, Form and function of cyclic peptide natural products: a pharmacokinetic perspective, Curr. Top. Med. Chem. 13 (2013) 821-836.
- [3] S.H. Joo, Cyclic peptides as therapeutic agents and biochemical tools, Biomol. Ther. 20 (2012) 19-26.
- [4] C.J. White, A.K. Yudin, Contemporary strategies for peptide macrocyclization, Nat. Chem. 3 (2011) 509-524.
- [5] Z.M. Wu, S.Z. Liu, X.Z. Chen, et al., Recent progress of on-resin cyclization for the synthesis of cyclopeptidomimetics, Chin. Chem. Lett. 27 (2016) 1731-1739.
- [6] G.K.T. Nguyen, C.T.T. Wong, Making circles: recent advance in chemical and enzymatic approaches in peptide macrocyclization, J. Biochem. Chem. Sci. 2017 (2017) 1-13.
- [7] P.E. Dawson, T.W. Muir, I. Clark-Lewis, et al., Synthesis of proteins by native chemical ligation, Science 266 (1994) 776-779.
- [8] P.E. Dawson, S.B.H. Kent, Synthesis of native proteins by chemical ligation, Annu. Rev. Biochem. 69 (2000) 923-960.

- [9] G.M. Fang, Y.M. Li, F. Shen, et al., Protein chemical synthesis by ligation of peptide hydrazides, Angew. Chem. Int. Ed. 50 (2011) 7645-7649.
- [10] S. Tang, J.S. Zheng, K. Yang, et al., Synthesis of cyclic tetrapeptides via ligation of peptide hydrazides, Acta Chim. Sinica 70 (2012) 1471-1476.
- [11] J.X. Wang, G.M. Fang, Y. He, et al., Peptide o-amino anilides as cryptothioesters for protein chemical synthesis, Angew. Chem. Int. Ed. 54 (2015) 2194-2198.
- [12] K. Jin, T. Li, H.Y. Chow, et al., P-B desulfurizaton: an enabling method for protein chemical synthesis and site-specific deuteration, Angew. Chem. Int. Ed. 56 (2017) 14607-14611.
- [13] C.F. Liu, J.P. Tam, Chemical ligation approach to form a peptide bond between unprotected peptide segments. Concept and model study, J. Am. Chem. Soc. 116 (1994) 4149-4153.
- [14] P. Botti, T.D. Pallin, J.P. Tam, Cyclic peptides from linear unprotected peptide precursors through thiazolidine formation, J. Am. Chem. Soc. 118 (1996) 10018-10024.
- [15] J.W. Bode, Chemical protein synthesis with the α-ketoacidhydroxylamine ligation, Acc. Chem. Res. 50 (2017) 2104-2115.
- [16] F. Rohrbacher, G. Deniau, A. Luther, et al., Spontaneous head-to-tail cyclization of unprotected linear peptides with the KAHA ligation, Chem. Sci. 6 (2015) 4889-4896.
- [17] T. Fukuzumi, L. Ju, J.W. Bode, Chemoselective cyclization of unprotected linear peptides by α-ketoacid-hydroxylamine amide-ligation, Org. Biomol. Chem. 10 (2012) 5837-5844.
- [18] R. Kleineweischede, C.P.R. Hackenberger, Chemoselective peptide cyclization by traceless Staudinger ligation, Angew. Chem. Int. Ed. 47 (2008) 5984-5988.
- [19] C.L. Tung, C.T.T. Wong, X. Li, Peptide 2-formylthiophenol esters do not proceed through a Ser/Thr ligation pathway, but participate in a peptide aminolysis to enable peptide condensation and cyclization, Org. Biomol. Chem. 13 (2015) 6922-6926.
- [20] D.Y. Jackson, J.P. Burnier, J.A. Wells, Enzymatic cyclization of linear peptide esters using subtiligase, J. Am. Chem. Soc. 117 (1995) 819-820.
- [21] G.K.T. Nguyen, S. Wang, Y. Qiu, et al., Butelase 1 is an Asx-specific ligase enabling peptide macrocyclization and synthesis, Nat. Chem. Biol. 10 (2014) 732-738.
- [22] G.K.T. Nguyen, Y. Qiu, Y. Cao, et al., Butelase-mediated cyclization and ligation of peptides and proteins, Nat. Protoc. 11 (2016) 1977-1988.
- [23] G.K.T. Nguyen, X. Hemu, J.P. Quek, et al., Batelase-mediated macrocyclization D-amino-acid-containing peptides, Angew. Chem. Int. Ed. 55 (2016) 12802-12806.
- [24] Z.M. Wu, S.Z. Liu, X.Z. Cheng, et al., High yield synthesis of cyclic analogues of antibacterial peptides P-113 by sortase A-mediated ligation and their conformation studies, Chin. Chem. Lett. 28 (2017) 553-557.
- [25] X. Li, H.Y. Lam, Y. Zhang, C.K. Chan, Salicylaldehyde ester-induced chemoselective peptide ligations: enabling generation of natural peptidic linkages at the serine/threonine sites, Org. Lett. 12 (2010) 1724-1727.
- [26] Y. Zhang, C. Xu, H.Y. Lam, et al., Protein chemical synthesis by serine and threonine ligation, Proc. Natl. Acad. Sci. USA 110 (2013) 6657-6662.
- [27] C.T.T. Wong, T. Li, H.Y. Lam, et al., Realizing serine/threonine ligation: scope and limitations and mechanistic implication thereof, Front. Chem. 2 (2014) 10.3389/fchem.2014.00028.
- [28] C.L. Lee, X. Li, Serine/threonine ligation for the chemical synthesis of proteins, Curr. Opin. Chem. Biol. 22 (2014) 108-114.
- [29] C.L. Lee, X. Li, Chemical synthesis of glycoproteins, Sci. China Chem. 59 (2016) 1061-1064.

- [30] H. Liu, X. Li, Development and application of serine/threonine ligation for synthetic protein chemistry, Org. Biomol. Chem. 12 (2014) 3768-3773.
- [31] Y. Zhang, T. Li, X. Li, Synthesis of human growth hormone-releasing hormone via three-fragment serine/threonine ligation (STL), Org. Biomol. Chem. 11 (2013) 5584-5587.
- [32] C. Xu, H.Y. Lam, Y. Zhang, et al., Convergent synthesis of MUC1 glycopeptides via serine ligation, Chem. Commun. 49 (2013) 6200-6202.
- [33] T. Li, H. Liu, X. Li, Chemical synthesis of HMGA1a proteins with posttranslational modifications via Ser/Thr ligation, Org. Lett. 18 (2016) 5944-5947.
- [34] C.L. Lee, H. Liu, C.T.T. Wong, et al., Enabling N-to-C Ser/Thr ligation for convergent protein synthesis via combing chemical ligation approaches, J. Am. Chem. Soc. 138 (2016) 10477-10484.
- [35] C.L. Lee, H.Y. Lam, X. Li, Serine/threonine ligation for natural cyclic peptide syntheses, Nat. Prod. Rep. 32 (2015) 1274-1279.
- [36] H.Y. Lam, Y. Zhang, H. Liu, et al., Total synthesis of daptomycin by cyclization via a chemoselevtive serine ligation, J. Am. Chem. Soc. 135 (2013) 6272-6279.
- [37] D. Lin, H.Y. Lam, W. Han, et al., Structure-activity relationship of daptomycin analogues with substitution at (2*S*, 3*R*) 3-methyl glutamic acid position, Bioorg. Med. Chem. Lett. 27 (2017) 456-459.
- [38] K. Jin, I.H. Sam, K.H.L. Po, et al., Total synthesis of teixobactin, Nat. Commun. 7 (2016) 10.1038/ncomms12394.
- [39] K. Jin, K.H.L. Po, S. Wang, et al., Synthesis and structure-activity relationship of teixobactin analogues via convergent serine ligation, Bioorg. Med. Chem. 25 (2017) 4990-4995.
- [40] C.T.T. Wong, H.Y. Lam, X. Li, Effective synthesis of cyclic peptide yunnanin C and analogues via Ser/Thr ligation (STL)-mediated peptide cyclization, Tetrahedron 70 (2014) 7770-7773.
- [41] C.T.T. Wong, H.Y. Lam, X. Li, Effective synthesis of kynureninecontaining peptides via on-resin ozonolysis of tryptophan residues: synthesis of cyclomontanin B, Org. Biomol. Chem. 11 (2013) 7616-7620.
- [42] C.T.T. Wong, H.Y. Lam, T. Song, et al., Synthesis of constrained headto-tail cyclic tetrapeptides by an imine-induced ring-closing/contraction strategy, Angew. Chem., Int. Ed. 52 (2013) 10212-10215.
- [43] J.F. Zhao, X.H. Zhang, Y.J. Ding, et al., Facile synthesis of peptidyl salicylaldehyde esters and its use in cyclic peptide synthesis, Org. Lett. 15 (2013) 5182-5185.
- [44] T. Fukuyama, S.C. Lin, L. Li, Facile reduction of ethyl thiol esters to aldehydes: application to a total synthesis of (+)-neothramycin A methyl ether, J. Am. Chem. Soc. 112 (1990) 7050-7051.
- [45] Y. Yoshimura, C. Ohara, T. Imahori, et al., Synthesis of both enantiomers of hydroxypipecolic acid derivatives equivalent to 5azapyranuronic acids and evaluation of their inhibitory activities against glycosidases, Bioorg. Med. Chem. 16 (2008) 8273-8286.

Supplementary material

Supplementary data with this article, including the synthetic procedures of building block **7**, characterization of related compounds, preparation of the peptide SAL esters, optimization of oxidation conditions, and HPLC traces of the aspartic acid ligation, can be found in the online version.