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Discovery of novel aspartyl ketone dipeptides as potent and selective caspase-3 inhibitors

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Abstract—The discovery of a series of potent, selective and reversible dipeptidyl caspase-3 inhibitors are reported. The iterative discovery process of using combinatorial chemistry, parallel synthesis, moleculare modelling and structural biology will be discussed. © 2003 Elsevier Ltd. All rights reserved.

Recent studies towards understanding the molecular mechanisms of apoptosis have revealed the importance of a group of cysteinyl aspartate specific proteases, the caspases, in the programmed cell death process. So far, 13 human members of caspases have been identified and these enzymes exist as dormant proenzymes which are processed to the catalytically active mature forms under certain conditions.² Caspases can be grouped into three subfamilies based on their substrate specificity³ and cellular functions. Group I (1, 4, and 5) caspases are believed to be involved primarily in inflammation. Group II (2, 3, 6, and 7) and group III (2, 6, 8, 9, and 10) caspases are essential for apoptotic cell death. Caspase-3 (casp-3 for short), in particular, has been characterized as the dominant effector caspase involved in the proteolytic cleavage of a variety of protein substrates including cytoskeletal proteins, kinases and DNA repair enzymes during apoptosis.² Casp-3 knockout murine phenotype suggested the necessity of this enzyme during brain development⁴ and recent studies have revealed its activation in many models of apoptosis.⁵ The development of potent and selective casp-3 inhibitors has thus emerged as an attractive therapeutic target.⁶

Recent reports indicated that caspase inhibitors such as z-VAD-fmk, Boc-D-fmk and Ac-DEVD-fmk were effective in animal models of ischemia injury, burns, endotoximia, sepsis and neonatal hypoxia. These inhibitors, however, are either irreversible pan-caspase inhibitors, inhibit other cysteine proteases, or have poor whole cell activity (e.g., Ac-DEVD-CHO, see Table 1) and in vivo stability. In order to assess the importance of casp-3 activation in apoptosis, potent, selective and reversible inhibitors are highly desirable. In this communication, we will describe the iterative process that led to the discovery of such inhibitors.

Reversible P_{1'} aldehyde and ketone casp-3 inhibitors incorporating DxVD, the preferred tetrapeptide motif recognized by casp-3, have been studied previously.¹² These compounds generally lack cell potency due to limited cell permeability. For example, Bz-DAVD-phenylpropyl ketone (17) was a potent inhibitor at rh-casp-3

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Table 1. Inhibitor potency against casp-1, -3, -7 and -8 and in NT2 cells

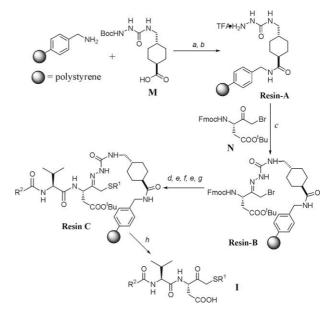
| Compd | IC ₅₀ , μM ^a | | | | |
|--------------------------|------------------------------------|--------|--------|--------|--------|
| | Casp-1 | Casp-3 | Casp-7 | Casp-8 | NT2 |
| 1 | 6.0 | 0.048 | 3.2 | 6.6 | 10 |
| 2 | 0.86 | 0.14 | 1.5 | 8.7 | 63.0 |
| 3 | 2.6 | 0.083 | 1.4 | 3.3 | 55.0 |
| 4 | 2.6 | 0.13 | 1.8 | 7.8 | 32.0 |
| 5 | 5.0 | 0.073 | 2.5 | 2.4 | 9.0 |
| 6 | 1.8 | 0.063 | >10 | 2 | 19.0 |
| 7 | 28 | 0.31 | 2.7 | 15.5 | NT^b |
| 8 | 14 | 0.21 | 3.1 | 14 | 60.0 |
| 9 | 22 | 0.75 | 8.4 | 22.5 | NT^b |
| 10 | 3.1 | 0.06 | 3.1 | 2.3 | 9.0 |
| 11 | 6.5 | 0.053 | 1.7 | 9.3 | 0.7 |
| 12 | 4.2 | 0.094 | 2.3 | 6.3 | 0.3 |
| 13 | 3.8 | 0.005 | 0.36 | 0.81 | 5.0 |
| 14 | 1.8 | 0.008 | 0.024 | 3.85 | 0.6 |
| 15 | 5.9 | 0.01 | 0.13 | 2.9 | 2.0 |
| 16 | 1.6 | 0.005 | 0.12 | 0.49 | 1.0 |
| 17 | 1.65 | 0.0018 | 0.0084 | 1.50 | 14.0 |
| Ac-DEVD-CHO ^c | 0.19 | 0.027 | 0.087 | 0.13 | > 100 |

^a Average of 2–8 titrations.

(IC₅₀, 1.8 nM) yet its potency was shifted dramatically in the whole cell assay (Table 1). Efforts to reduce the peptidic nature of these inhibitors by capping aspartic acid-phenylpropyl ketone with simple acids did not afford inhibitors with appreciable activity against recombinant human casp-3 (rh-casp-3). Preservation of the P_2 valine, however, was beneficial and resulted in inhibitors with high nano-molar potency, and thus represented a good starting point for inhibitor design. ¹³ In this communication, we will describe the development of two facile two-dimensional solid-phase combinatorial strategies that enabled the concomitant modification of both the $P_{1'}$ with heteroatom substitutions ¹⁴ and $P_2/P_3/P_4$ positions.

Central to this strategy was the direct attachment of bromomethyl ketone N¹⁵ to semicarbazide resin-A in THF in presence of acetic acid as outlined in Scheme 1.¹⁶ The semicarbazide linker M¹⁷ serves the dual purpose of protecting the ketone functionality and anchoring the reactive bromide to solid support. This strategy simplifies significantly the synthetic sequence described in the original procedure by Webb and co-workers.¹⁶ Nucleophilic displacement of the bromide with thiols in the presence of Hünig's base followed by standard Fmoc deprotection/coupling protocol for peptide synthesis afforded Resin C. The release of the final products from solid support was achieved by treatment with trifluoroacetic acid (TFA) in water (9/1, v/v) (instead of concentrated HCl).

The initial library was constructed according to the split and pool protocol using the IRORI MacroKan[®] technology. ¹⁸ 10 Carboxylic acids (a–j) and 10 thiols (A–J) were chosen as building blocks (Fig. 1). The library rapidly identified the benzylthio (thiol D) and the isobutylthio (thiol J) moieties as the preferred groups in the $P_{1'}$ position while 2,5-dimethoxyphenyl-acetate (1), 3,5-dibromo-benzoate (2) and 5-F-indole-2-carboxylate (3) were the best capping acids at P_3 – P_4 . Four of the most



Scheme 1. General strategy for preparing dipeptide of structure **I.** Reagents and conditions: (a) EDCI, HOBt, DMF; (b) TFA, DCM; (c) AcOH, THF; (d) R¹SH, *i*Pr₂NEt, DMF; (e) 20% piperidine in DMF; (f) Fmoc-V-OH, HATU, *i*Pr₂NEt, DMF; (g) R²COOH, HATU, *i*Pr₂NEt, DMF; (h) TFA, H₂O (9:1 v/v).

potent compounds were re-synthesized and their inhibition against casp-1, -3, -7, and -8 determined (Table 1). All of the inhibitors were shown to be selective, fully reversible and competitive against rh-casp-3. Compound 1, however, was the most potent among the four in the NT2 whole cell assay¹⁹ with an IC₅₀ of 10 μ M while 2, 3, and 4 were less potent.

The enhanced cell potency of 1 can be attributed to its improved cell permeability (vs tetrapeptide 17, Table 1). Using tritiated analogue ³H-1 and ³H-17 (prepared from the corresponding Br analogues), we were able to show that the intracellular concentration of 1 in NT2 cells was 6–20-fold higher than 17 (Fig. 2) (similar results were obtained in caco-2 monolayers).

Following the discovery of 1, extensive SAR studies at the $P_{1'}$ position with more than 100 thiols were carried out using the ACT-496TM automated synthesizer and the best results were obtained with lipophilic thiols such as benzylthio, substituted benzylthio and cyclohexyl-methylthio

^bNot tested.

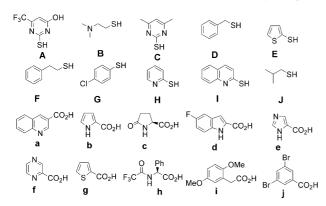


Figure 1. Building blocks for library 1 (L-1).

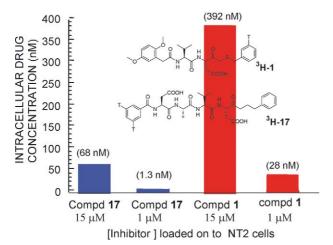
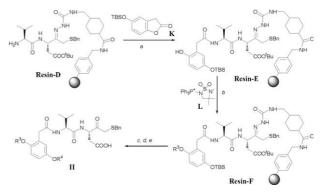


Figure 2. Cell permeability studies with tritiated compounds.

groups. In an another attempt, the 2,5-dimethoxy-phenylacetate moiety was replaced with a variety of acids (substituted phenylacetate, benzoate, phenylpropionate, naphthoate, cinnamate, hetero-aromatic, aliphatic, etc.) and no significant improvement was obtained. We instead focused on the modification of both methoxy groups. We observed that both groups were required for optimal potency. Removal of either of the methoxy group resulted in 5-fold loss of activity (7 and 8) and removal of both groups gave compound 9 which was 15-fold less active. To further address the SAR at the two OMe positions, a two-dimensional library was constructed as outlined in Scheme 2.

This library was constructed using 10 alcohols at OR³ and 50 alcohols at OR⁴. Reaction of **resin-D** with lactone **K** in the presence of DMAP in DMF at 50 °C overnight gave phenolic intermediate **resin-E**. The phenol moiety thus generated was reacted with an alcohol R³OH using reagent **L** as described by Castro et al.,²⁰ affording **resin-F**. After deprotection of the TBS group with TBAF and AcOH, the second phenolic OH was reacted with a second alcohol R⁴OH under identical conditions and the final product **II** was released from solid support upon treatment with TFA in water.

A three-point titration (0.1, 1, and 10 μ M) of this library in the NT2 whole cell assay quickly revealed that while a variety of groups were tolerated at OR³, the most potent



Scheme 2. Modification of 2,5-dimethoxyphenylacetate moiety. Reagents and conditions: (a) reagent K, DMAP, DMF, 50 °C; (b) R³OH, reagent L, toluene/DCM; (c) TBAF, AcOH, THF; (d) R⁴OH, L, toluene/DCM; (f) TFA, H₂O.

compounds in the NT2 assay contained glycolate moieties at OR^4 (11 and 12). Furthermore, these compounds were not significantly shifted (intrinsic to whole cell) in comparison to other compounds tested. We speculated that the ester was acting as the pro-drug and was hydrolyzing to the corresponding acid 13 intracellularly. This was confirmed by incubating 3H -11 (prepared similarly as described for 3H -1) in NT2 cells at 37 °C for 0.5 h. HPLC analysis revealed that the extracellular culture medium contained both ester 11 and acid 13, while the intracellular content was exclusively acid 13. Compound 13 was tested against rh-casp-3 and was found to be potent (IC_{50} : 5 nM) yet highly shifted in the NT2 assay (5 μ M). Therefore, the enhanced whole cell activity of 11 and 12 was due to their improved cell permeability.

Modelling of acid 13 into the active site of rh-casp-3 suggested that the carboxylate residue of the glycolate would be interacting with the two neutral NH bonds from residues Phe381b and Asn342 in the S₄ pocket, similarly to the P₄ Asp of substrate Ac-DEVD-CHO. This was consistent with the X-ray structure of compound 10,²¹ the bromo analogue (Fig. 3). This structure

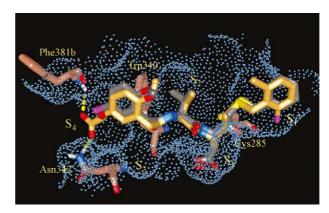


Figure 3. Compound **13** (gold) modelled into the active site of the X-ray structure of the complex between compound **10** (grey) and rh-casp-3 (pink); the two inhibitors virtually superimpose for their common structure from P₁ to P₃. In S₄, yellow dotted lines show the hydrogen bonding between the glycolate of compound **10** and the protein (Asn342 side chain and Phe381b main chain), the same hydrogen bonding present in the substrate Ac-DEVD-CHO. Above S₃, behind the plane of the figure, the edge of the Trp340 indole ring can be seen making an edge-to-face interaction with the phenyl ring for both compounds **10** and **13**.

confirmed that the $P_{1'}$, P_1 , and P_2 residues of 10 bound similarly to tetrapeptide inhibitors. The edge-to-face interaction of the phenyl group with Trp340 probably contributes to the enhanced potency of the phenylacetate moiety. The bromine atom was pointing straight to the S_4 pocket where we propose that the glycolate will be localized.

Extensive efforts were then directed towards finding a suitable neutral glycolate replacement at OR^4 . A methyl ketone was found to be an effective replacement, giving compound 15 with good intrinsic activity against rh-casp-3 (IC_{50} : 0.01 μM) and reasonable whole cell potency (IC_{50} : 2 μM). Modelling suggested that a methyl sulfone moiety would make excellent interactions with the two neutral amide NH's at the S_4 pocket of casp-3. Indeed, compound 16 was found to have excellent intrinsic and whole cell potency with IC_{50} 's of 0.005 and 1 μM , respectively. As illustrated in the Table, compound 16 was also selective against other caspases tested.

In summary, a series of potent and selective casp-3 inhibitors were discovered employing iterative combinatorial strategies. The selectivity and potency profile of compounds such as 16 makes it a useful tool for studying the importance of casp-3 activation in cell based systems.

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