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## Communications to the Editor

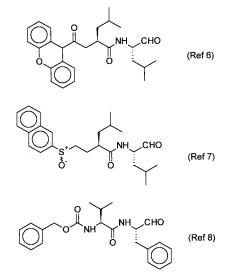
### D-Amino Acid Containing, High-Affinity Inhibitors of Recombinant Human Calpain I<sup>†</sup>

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Calpains are calcium-activated neutral proteases belonging to the papain superfamily of cysteine proteases. Two major forms of calpains have been identified: calpain I (*µ*-calpain) and calpain II (m-calpain), which require low and high micromolar Ca<sup>2+</sup> concentrations for activation, respectively. While calpain II is the predominant form in many tissues, calpain I is thought to be the predominant form activated during pathological conditions in nervous tissues.<sup>1</sup> Calpain I has been implicated in many nervous system disorders including stroke, Alzheimer's disease, motor neuron damage, and muscular dystrophy; thus, in recent years, calpain inhibition has become an important pharmacological goal.<sup>2</sup> Our involvement in inhibiting calpain emerged from our interest in finding new therapeutics<sup>3</sup> to treat stroke, one of the leading causes of mortality in the western hemisphere. An episode of stroke (focal cerebral ischemia) initiates a chain of biochemical events resulting in a delayed neurodegeneration. Calpains become activated by the increased intraneuronal calcium concentrations associated with ischemia. Acti-



#### Figure 1.

vated calpains, in turn, degrade neuronal structural proteins, contributing to the neurodegenerative process.

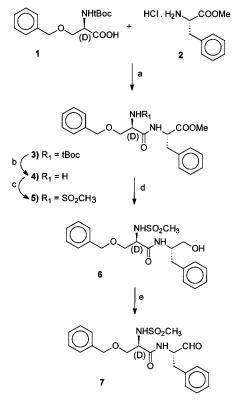
Potent peptide-based reversible (aldehyde and  $\alpha$ -ketocarbonyl)<sup>4</sup> and irreversible (halomethyl ketone, diazomethyl ketone, epoxysuccinate, and acyloxymethyl ketone)<sup>5</sup> inhibitors of calpain have been reported. In all of these inhibitors, calpain tolerated a range of amino acids at  $P_1$ . However, in potent inhibitors, the  $P_2$  amino acid was always either L-Leu or L-Val, suggesting this could be a strict structural requirement of calpain at the  $P_2$  site.<sup>4,5</sup> Recently, we disclosed potent nonpeptidic ketomethylene and carbamethylene  $(P_2-P_3)$  containing calpain I inhibitors derived from xanthene<sup>6</sup> and thionaphthalene,7 respectively (Figure 1). Our work revealed that the NH at the P<sub>2</sub> site of a potent dipeptide inhibitor can effectively be replaced by a CH<sub>2</sub>, provided an aromatic moiety is employed in the P3 region. We reasoned that the requirement of calpain I for an isobutyl group (from leucine) or an isopropyl group (from valine) might be steric in nature; either of these moieties could occupy the same pocket of the enzyme's S<sub>2</sub> subsite. In designing our target molecule, we decided to replace the P<sub>2</sub>-isopropyl group in the known potent calpain I

 $<sup>^\</sup>dagger$  Dedicated to Professor Steven M. Weinreb with admiration and best wishes.

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Scheme 1<sup>a</sup>



<sup>a</sup> Reagents: (a) BOP, 1-HOBt, NMM, DMF, 0 °C to room temperature, 2 h; (b) 90% TFA,  $CH_2Cl_2$ , room temperature, 3 h; (c)  $CH_3SO_2Cl$ , NMM,  $CH_2Cl_2$ , 0 °C to room temperature, 1 h; (d) LiBH<sub>4</sub>, THF, 0 °C to room temperature, 1 h or NaBH<sub>4</sub>, MeOH, 0 °C to room temperature, overnight; (e) Dess-Martin periodinane,  $CH_2Cl_2$ , 0 °C to room temperature, 1 h.

inhibitor, Cbz-Val-Phe-H (MDL 28170, Figure 1),<sup>8</sup> by a sulfonamido group. At the same time, as described before, we wished to incorporate an aromatic moiety in the  $P_3$  region attached by a spacer to the  $P_2$  site.

**Chemistry.** Scheme 1 depicts a representative synthesis of the target compound 7.<sup>9</sup> Commercially available Boc-D-Ser(Bn) (1) was coupled with L-Phe-OMe hydrochloride salt (2) to generate the dipeptide 3; the *t*-BocNH group in 3 was converted to amine 4 which was coupled with methanesulfonyl chloride to generate the sulfonamide 5. Reduction of the ester group in 5 produced the dipeptide alcohol 6 which underwent Dess-Martin oxidation to generate the target aldehyde 7. Alternatively, compound 6 can be prepared by coupling methanesulfonyl-D-Ser(Bn) and (*S*)-phenylalaninol; methanesulfonyl-D-Ser(Bn) can be generated from D-Ser(Bn) by treatment with methanesulfonyl chloride in the presence of aqueous NaOH.

**In Vitro Biology and Discussion.** The biological activities of the compounds were determined using recombinant human calpain I (rh calpain I), prepared as described by Meyer et al.,<sup>10</sup> with Suc-Leu-Tyr-MNA (Enzyme Systems Products, Dublin, CA) as the substrate.<sup>6</sup> Table 1 displays the inhibitory data for a series of P<sub>2</sub>-D-Ser(Bn) derived inhibitors and the reference compound Cbz-Val-Phe-H. All reaction progress curves in the presence of inhibitor were linear, consistent with rapid equilibrium binding of the compounds to calpain. As shown, P<sub>1</sub>-Phe is preferred over Abu and Leu (cf. **7** vs **8** and **9**). Interestingly, incorporation of P<sub>1</sub>-Lys(SO<sub>2</sub>-Ph) and Tyr(Bn) generated **10** and **11** (*K*<sub>i</sub> values of 2

**Table 1.**  $P_2$ -D-Ser(Bn)-Derived Inhibitors of RecombinantHuman Calpain  $I^a$ 

|     | O = O = O = O = O = O = O = O = O = O = |        |  |        |                 |                        |  |
|-----|---|--------|--|--------|-----------------|------------------------|--|
|     | compd                                   | X      | <b>R</b> <sub>1</sub>                                | $R_2$  | $R_3$           | K <sub>i</sub><br>(nM) |  |
| 7   |   | Н      | Bn   | Н      | CH <sub>3</sub> | 8                      |  |
| 8   |   | Н      | Et   | Н      | $CH_3$          | 36                     |  |
| 9   |   | Н      | <i>i</i> Bu  | Н      | CH <sub>3</sub> | 66                     |  |
| 10  |   | Н      | (CH <sub>2</sub> ) <sub>4</sub> NHSO <sub>2</sub> Ph | Н      | CH <sub>3</sub> | 2                      |  |
| 11  |   | Н      | CH <sub>2</sub> C <sub>6</sub> H <sub>4</sub> -p-OBn |        | CH <sub>3</sub> | 4                      |  |
| 12  |   | Н      | Bn   | Н      | Et              | 9                      |  |
| 13  |   | Н      | Bn   | Н      | Ph              | 22                     |  |
| 14  |   | Н      | Bn   | Н      | 2-thienyl       | 9                      |  |
| 15  |   | Н      | Bn   | $CH_3$ | CH <sub>3</sub> | 370                    |  |
| 16  |   | CONHEt | Bn   | Н      | $CH_3$          | 130                    |  |
| Cba | z-Val-Phe-H                             |        |  |        |                 | 8                      |  |

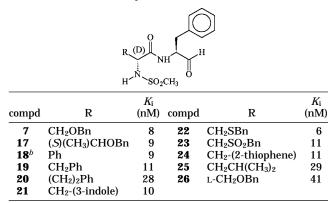
<sup>*a*</sup>  $n \ge 3$  in all cases. Assay conditions as described previously.<sup>5a</sup> The  $K_m$  for Suc-Leu-Tyr-MNA = 0.5 mM. Values for IC<sub>50</sub> were determined and converted to  $K_i$ s using the expression  $K_i = IC_{50}/(1 + S/K_m)$ , assuming a competitive mechanism of inhibition.<sup>12</sup> Replicate determinations of  $K_i$  agree within 25%.

and 4 nM, respectively), the two most potent compounds of the series; thus, it appears that the S<sub>1</sub> pocket of the enzyme tolerates large hydrophobic groups from the P1 position. It should be noted that the reference compound, Cbz-Val-Phe-H, shows a K<sub>i</sub> value of 8 nM in this assay. Variation in sulfonamide groups generated compounds 7, 12, 13, and 14. While methane- and ethanesulfonamides were preferred over benzenesulfonamide (cf. 7 and 12 vs 13), 2-thienylsulfonamide (14) was equipotent, revealing the beneficial effect of an additional heteroatom in this region. Compound 15, which contains an N-methyl methanesulfonamide, is approximately 46 times less potent than compound 7, suggesting that the NH of the sulfonamide moiety of compound 7 is involved in energetically beneficial binding. The aldehyde enzyme reactive group is approximately 16 times preferred over the  $\alpha$ -ketocarboxamide group (cf. 7 vs 16). Compound 16 ( $K_i$  130 nM) is, however, comparable to a reference L,L-dipeptide  $\alpha$ -ketocarboxamide, Cbz-Leu-Abu-CONHEt<sup>4c</sup> (K<sub>i</sub> 170 nM, in this assay).

Table 2 displays the inhibitory activity of a series of compounds with varied P<sub>2</sub>-D-amino acids (except 26, see below). D-Ser(Bn), D-Thr(Bn), D-Phgly, D-Phe, D-Trp, and D-(thiophen-2-yl)Ala (7, 17, 18, 19, 21, 24) are all well tolerated at  $P_2$ . However, D-Phe at  $P_2$  (19) is approximately 2.5 times more potent than D-Homophe at  $P_2$  (20). D-Cys(Bn) (22) and the corresponding sulfone analogue (23) at P<sub>2</sub> also maintain potency. Interestingly, incorporation of D-Leu at P<sub>2</sub> results in a ca. 4 times less potent compound (cf. **25** vs **7**), supporting our original hypothesis that the presence of an aromatic moiety in the P<sub>3</sub> region is beneficial. Finally, incorporation of L-Ser(Bn) at P2 results in a greater than 5-fold decrease in potency (cf. 26 vs 7) revealing the importance of the D configuration at P2 in this series of compounds.

**Evaluation of the Site of Binding.** It is very difficult to demonstrate competitive inhibition kinetics with calpain by the classic method of examining the effect of compounds at various concentrations of substrate. Calpain substrates cannot be varied across a

**Table 2.** Variation of  $P_2$ -D-Amino Acids: Potency againstRecombinant Human Calpain  $I^a$ 



<sup>*a*</sup>  $n \ge 3$  in all cases. Values for IC<sub>50</sub> were determined and converted to  $K_{is}$  using the expression  $K_{i} = IC_{50}/(1 + S/K_{m})$ , assuming a competitive mechanism of inhibition.<sup>12</sup> Replicate determinations of  $K_{i}$  agree within 25%. <sup>*b*</sup>Elemental analysis: C<sub>18</sub>H<sub>20</sub>N<sub>2</sub>O<sub>4</sub>S·0.75H<sub>2</sub>O; calc H, 5.79; found H, 5.30.

sufficiently large concentration range to permit determination of the mechanism of inhibition, due either to limited solubility of the substrate or to substrate inhibition. To determine whether this novel series of compounds binds at the active site of calpain I, protection against inactivation experiments were performed. In this experiment, calpain I is preincubated with the test compound, followed by the addition of E-64 (transepoxysuccinyl-L-leucylamido-(4-guanidino)butane), an irreversible inhibitor of calpain. By analogy with other well-studied cysteine proteases, E-64 is presumed to covalently label the enzyme's active site cysteine. The enzyme is then diluted into an assay mixture and its residual activity determined. The dilution factor is sufficiently large that any bound test compound will dissociate from the enzyme and the remainder will be at a concentration well below its  $IC_{50}$ . If the test compound and E-64 bind in a mutually exclusive manner, no loss of activity should be observed, while inactivation by E-64 should be apparent if the two compounds occupy distinct sites. Experiments with methanesulfonyl-D-Leu-Leu-H (27,  $K_i$  68 nM), a less potent analogue, reveal (data not shown) that this compound protects against inactivation of calpain I by E-64, consistent with the binding of both compounds to a common site on the enzyme. However, this result should not be interpreted to demonstrate that both compounds bind in exactly the same way. For example, it is possible that compound 27 could react with the active site cysteine, but might lie in the pocket on the P' side of the active site. Such a binding motif would still result in protection against inactivation by E-64, even though the two binding sites would be at least partially distinct. Elucidation of the exact nature of binding of this class of inhibitors to the enzyme thus requires determination of an X-ray crystal structure of the enzyme-inhibitor complex.

**Specificity data.** The selectivity of a subset of inhibitors was tested against several representative proteases. Table 3 lists the inhibitory data for compounds 7, **10–12**, **17**, **19**, **22**, and **24** and the reference compound, Cbz-Val-Phe-H. The closely related enzyme calpain II is inhibited by all compounds tested with  $K_i$  values which are indistinguishable (except compound

**Table 3.** Specificity Data for Compounds 7, 10–12, 17, 19, 22, and  $24^a$ 

|               |              | K <sub>i</sub> (nM         | )                           | % inh at 10 $\mu$ M   |                                  |
|---------------|--------------|----------------------------|-----------------------------|-----------------------|----------------------------------|
| compd         | calpain<br>I | calpain<br>II <sup>b</sup> | cathepsin<br>B <sup>c</sup> | thrombin <sup>d</sup> | α-chymo-<br>trypsin <sup>e</sup> |
| 7             | 8            | 5                          | 32                          | 0                     | 23                               |
| 10            | 2            | 17                         | 4                           | 0                     | 13                               |
| 11            | 4            | 5                          | 45                          | 0                     | 0                                |
| 12            | 9            | 9                          | 45                          | 9                     | 15                               |
| 17            | 9            | 7                          | 44                          | 0                     | 3                                |
| 19            | 11           | 13                         | 120                         | 0                     | 12                               |
| 22            | 6            | 7                          | 41                          | 7                     | 16                               |
| 24            | 11           | 11                         | 23                          | 0                     | 0                                |
| Cbz-Val-Phe-H | 8            | f                          | 24                          | f                     | f                                |

<sup>*a*</sup> *n* ≥ 3 in all cases. Values for IC<sub>50</sub> were determined and converted to *K*<sub>i</sub>s using the expression *K*<sub>i</sub> = IC<sub>50</sub>/(1 + S/*K*<sub>m</sub>), assuming a competitive mechanism of inhibition.<sup>12</sup> Replicate determinations of *K*<sub>i</sub> agree within 25%. <sup>*b*</sup>Porcine kidney calpain II (Calbiochem) assayed under the same conditions employed for calpain I (*K*<sub>m</sub> for Suc-Leu-Tyr-MNA = 0.4 mM). <sup>*c*</sup> Human liver cathepsin B (Athens Research and Technology), 50 mM NaOAc (pH 6.0), 1 mM EDTA, 1 mM DTT, 100 µM Z-Phe-Arg-AMC (*K*<sub>m</sub> = 300 µM), 5% DMSO. <sup>*d*</sup>Human plasma thrombin (Sigma), 50 mM Tris-Cl (pH 7.5), 10 mM CaCl<sub>2</sub>, 100 µM Z-Phe-Val-Arg-AMC (*K*<sub>m</sub> = 60 µM), 5% DMSO. <sup>*c*</sup>Bovine pancreas α-chymotrypsin (Sigma), 50 mM HEPES (pH 7.5), 0.5 M NaCl, 100 µM Suc-Ala-Ala-Pro-Phe-AMC (*K*<sub>m</sub> = 170 µM), 5% DMSO. <sup>4</sup>Not determined.

**Table 4.** Inhibitory Data for Compounds **7**, **10–12**, **17**, **19**, **22**, and **24** in Intact Cell Assay<sup>*a*</sup>

| compd | IC <sub>50</sub> (µM) | compd         | IC <sub>50</sub> (µM) |
|-------|-----------------------|---------------|-----------------------|
| 7     | 0.7                   | 19            | 0.6                   |
| 10    | 0.4                   | 22            | 0.8                   |
| 11    | 0.7                   | 24            | 0.3                   |
| 12    | 0.4                   | Cbz-Val-Phe-H | 0.3                   |
| 17    | 0.5                   |               |                       |

<sup>*a*</sup>  $n \ge 2$  in all cases.

**10**) from those obtained against calpain I. Cathepsin B is a related cysteine protease which is sensitive to inhibition by L,L-dipeptide aldehydes. While both the reference compound and the parent compound **7** show a similar preference (ca. 3-4-fold) for calpain I over cathepsin B, compounds **11** and **19** both prefer calpain I by 11-fold over cathepsin B. Table 3 also displays the inhibitory activity of these compounds against  $\alpha$ -chymotrypsin, a serine protease which is inhibited by other peptidyl P<sub>1</sub>-phenylalaninals. Excellent selectivity was observed with respect to inhibition of chymotrypsin. A second serine protease, thrombin, which displays a distinct P<sub>1</sub> specificity, also was not inhibited (Table 3).

Cellular Activity. To probe the ability of these compounds to penetrate cells and inhibit intracellular calpain I, we tested a set of compounds in an intact cell assay system. Treatment of Molt 4 cells (human leukemic T cell line) with calcium ion and an ionophore results in the elevation of intracellular calcium which, in turn, activates calpain I.<sup>11</sup> This is followed by calpain I-mediated cleavage of cytoskeletal proteins, including spectrin. Inhibition of formation of spectrin breakdown products (SBDPs) inside the cell by a compound measures its efficacy. Table 4 lists the inhibitory activities of the compounds 7, 10-12, 17, 19, 22, and 24, which showed good activity in the assay for calpain I inhibition. In general, these compounds also display good activity in this assay with IC<sub>50</sub> values in the range of 0.3–0.8  $\mu$ M. Thus, the compounds are cell-permeable and inhibit intracellular calpains.

**Conclusion.** In this paper, we described a series of novel and potent calpain inhibitors incorporating *N*-alkyl- or -arylsulfonyl-D-amino acids at  $P_2$ . To our knowledge, this is unprecedented in the design of substrate-based potent cysteine protease inhibitor. The compounds are cell-permeable and inhibit intracellular calpain I in a human cell line. This study reveals for the first time that, contrary to literature evidence, the presence of L-Leu or L-Val residue at  $P_2$  is not a preferred structural requirement for a potent calpain I inhibitor; an *N*-alkyl- or -arylsulfonyl-D-amino acid at  $P_2$  can bind with high affinity.

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