Macrocyclic Inhibitors of β -Secretase: Functional Activity in an Animal Model

Shawn J. Stachel,*,[†] Craig A. Coburn,[†]
Sethu Sankaranarayanan,[‡] Eric A. Price,[‡] Beth L. Pietrak,[‡]
Qian Huang,[‡] Janet Lineberger,[‡] Amy S. Espeseth,[‡]
Lixia Jin,[§] Joan Ellis,[§] M. Katharine Holloway,^{||}
Sanjeev Munshi,[⊥] Timothy Allison,[⊥] Daria Hazuda,[‡]
Adam J. Simon,[‡] Samuel L. Graham,[†] and Joseph P. Vacca[†]
Department of Medicinal Chemistry, Biological Chemistry, Molecular Systems and Structural Biology Merck Research Laboratories, Post Office Box 4, West Point, Pennsylvania 19486

Received July 26, 2006

Abstract: A macrocyclic inhibitor of β -secretase was designed by covalently cross-linking the P1 and P3 side chains of an isophthalamidebased inhibitor. Macrocyclization resulted in significantly improved potency and physical properties when compared to the initial lead structures. More importantly, these macrocyclic inhibitors also displayed in vivo amyloid lowering when dosed in a murine model.

Alzheimer's disease (AD^a) is a debilitating neurodegenerative disease characterized by progressive cognitive decline, inevitably leading to incapacitation and death. Although researchers have uncovered a variety of genetic mutations that result in early onset AD over the past 20 years, our current understanding of genetic predisposition accounts for <10% of AD cases.¹ The majority of AD cases are sporadic in nature, and the risk factor increases in an age-dependent manner. It is generally recognized that β -amyloid plaques and neurofibrillary tangles are the key pathological features of the disease, although the specific roles of these agents in disease progression remain debatable. However, it is well-established that the principle component of amyloid plaque is a neurotoxic peptide fragment of the amyloid precursor protein (APP), namely, $A\beta_{40-42}$.² As such, many therapeutic strategies are founded on the ability to inhibit catabolic pathways of APP that result in $A\beta$ formation.3

Proteolytic processing of APP is primarily accomplished through the activity of three discrete proteases (α -, β -, and γ -secretase).⁴ The initial cleavage event is performed by either α -secretase or β -secretase (BACE-1) releasing the APP ectodomain. The remaining membrane-bound C-terminal domains are subsequently cleaved by γ -secretase, resulting in the nonamyloidogenic P3 peptide (α -cleavage product) or the amyloidogenic A β_{40-42} (β -cleavage product). γ -Secretase inhibitors have been demonstrated to inhibit the production of $A\beta$ in vivo, however, their development as therapeutic agents remains tenuous due to the small therapeutic window that results from untoward effects on notch processing.⁵ In contrast, inhibitors of BACE-1 are predicted to be well-tolerated, as evidenced by viable knockout mice.⁶ As such, BACE-1 inhibition is considered an attractive therapeutic target for the treatment and prevention of AD.7 In this letter, we describe the

[⊥] Department of Structural Biology.

development of a series of macrocyclic active site inhibitors of BACE-1. Most noteworthy, compound **19** was shown to reduce $A\beta$ levels in a murine model following iv administration.



Figure 1. Representative potent BACE-1 inhibitors.

5-Substituted isophthalamides, as represented by structures 1 and 2 (Figure 1), emerged as early lead classes that were investigated for inhibition of BACE-1.8 Both inhibitors possessed desirable properties of both good in vitro ($IC_{50} = 15$ and 7 nM, respectively) and good cell-based potencies (IC₅₀ = 29 and 20 nM, respectively) against BACE-1. However, problems associated with their pharmacokinetic properties rendered them less than ideal candidates for advancement. A main concern was that 1 and 2 were found to be substrates of P-glycoprotein (P-gp) transport and both exhibited low apparent permeability in wild-type porcine LLC-PK1 cells. While iv administration of either 1 or 2 showed no effect on $A\beta$ levels in a murine model, intracranial administration of 1 succeeded in reducing A β levels by 50–70%.⁹ This result demonstrated that inhibition of BACE-1 can reduce $A\beta$ levels in vivo, but to be clinically relevant, the compound must be CNS penetrant. To achieve greater CNS penetration, we sought to reduce the peptidic character of our inhibitors while maintaining potency. Macrocyclization is a well-precedented technique to enhance potency by conformational preorganization, and in many cases, the physical properties of the inhibitors were also improved.¹⁰ We envisaged that, if macrocyclization were successful in enhancing potency, reduction of molecular weight and removal of some hydrogen bonding elements might be possible. Several research groups have recently reported similar macrocyclization strategies for BACE inhibitors, albeit with limited success.¹¹ In our case, examination of the X-ray crystallographic data of 1¹² complexed in the active site of BACE-1 revealed the close spatial proximity of the lipophilic P1 and P3 groups and thus directed efforts toward the design of macrocycles that would link these groups in an isophthalamide-based inhibitor.

We chose **4** as a simplified core to test whether macrocyclization would provide an increase in potency due to a reduction in conformational freedom (Figure 2). We were gratified to discover that while **4** was not a high affinity ligand for BACE-1 (IC₅₀ = 2.9 μ M) it was vastly superior to the corresponding seco variant **3** (IC₅₀ > 100 μ M). It should be noted that compound **4** was designed as a model system and thus lacks several potency enhancing features paramount for high binding affinity.



Figure 2. Macrocyclization results in increased potency.



^{*} To whom correspondence should be addressed. Phone: 216-652-2273. Fax: 215-652-3971. E-mail: shawn_stachel@merck.com.

[†] Department of Medicinal Chemistry.

[‡] Department of Biological Chemistry.

[§] Department of Drug Metabolism.

Department of Molecular Systems.

^{*a*} Abbreviations: AD, Alzheimer's disease; APP, amyloid precursor protein; BACE-1, beta-site APP cleaving enzyme 1; APP-YAC, amyloid precursor protein-yeast artificial chromosome.

Scheme 1^a



^{*a*} Reaction conditions: (a) *m*-tyrosine ethyl ester BOP, DIPEA, CH₂Cl₂, 86%; (b) *tert*-butyl(2-iodoethyl)carbamate, K₂CO₃, DMF, 75%; (c) Pearlman's catalyst, MeOH, 100%; (d)TFA, CH₂Cl₂, 100%; (e) BOP, DIPEA, CH₂Cl₂, 37%; (f) LiBH₄, THF, 51%; (g) SO₃-pyridine, TEA, DMSO; (h) (*S*)-2-amino-*N*-isobutylbutanamide, NaBH₃CN, MeOH, 60%.

Based on the encouraging results of the model system, we focused on incorporating a prime side element to enhance binding. We chose to incorporate a ψ [CH₂NH]-reduced amide isostere as present in **2**, because we have previously shown that such isosteres are not only highly active pharmacophores for BACE inhibition, but also improved cell permeability in similar series.^{8b}

The synthesis of macrocycle **10** began with a coupling reaction between isophthalamide **5** and *m*-tyrosine methyl ester to produce phenol **6** (Scheme 1). Alkylation of the phenol using *tert*-butyl(2-iodoethyl)carbamate afforded the differentially protected macrocyclic precursor **7** in 75% yield. Deprotection of the benzyl ester, followed by removal of the amino BOC protecting group, afforded *seco*-acid **8** in near quantitative yield. *seco*-Acid **8** was subjected to macrocyclization mediated by BOP reagent under high dilution conditions to provide **9** in modest yield. The ester was reduced to the alcohol **4** with lithium borohydride and subsequently oxidized to the corresponding aldehyde using Parikh–Doering conditions. Reductive amination of the corresponding aldehyde with (*S*)-2-amino-*N*-isobutylbutanamide afforded macrocyclic inhibitor **10**.

Macrocycle 10, as a 1:1 mixture of diastereomers at the tyrosinyl center, was assayed for BACE-1 activity and exhibited an $IC_{50} = 32$ nM. This represents a 90-fold increase in potency due to addition of the prime-side binding domain. It should be noted that macrocycle 10 is only 4-fold less-potent than 2 and lacks the P3 phenyl substituent that has been shown to contribute upward of 90-fold in potency. Unfortunately, when 10 was evaluated in the cell-based assay, there was a significant loss in potency (IC₅₀ = 5400 nM) when compared with **2**. It was also suspected that 10 was still a severe P-gp substrate, but due to a low apparent permeability (0.5 \times 10⁻⁶ cm/s), its susceptibility could not be accurately determined. Despite the ability to concomitantly increase potency and trim the molecular weight of the inhibitor, it was apparent that to succeed in lowering A β in vivo it would be necessary to improve the cell permeability of our inhibitors.

In a previous communication, we demonstrated that the P2/ P3 amide is noncritical for BACE-1 activity.¹³ The amide



Figure 3.

functionality acts mainly to orient the P3 ligand, and the amide hydrogen bonding contacts do not significantly contribute to the binding energy. With this in mind, we posited that the conformational constraint induced by the macrocycle would allow for excision of the amide bond without abrogating potency. It was also hoped that removal of the amide bond would result in improved physical properties, namely, cell-permeability. To test this hypothesis, we synthesized the corresponding alkyl-linked 15-membered macrocycle **11**, but were disappointed by a greater than 34-fold loss in intrinsic potency, IC₅₀ = 1100 nM (Figure 3).

Recognizing that the 15-membered macrocyclic ring size for the bisamide-containing macrocycle **10** may not be optimal for the carbon analogue, we synthesized the ring-contracted 14membered macrocycle **12** for comparison. We were gratified to see that potency was recovered and **12** exhibited a potency of 90 nM when assayed against BACE-1. Even more importantly, activity for **12** in the cell-based assay (IC₅₀ = 1139 nM) as well as apparent permeability ($P_{app} = 19 \times 10^{-6}$ cm/s) were also improved when compared with **10**. With these encouraging results, we employed some potency enhancing modifications to the sulfonamide and P1' regions of the inhibitor learned from previous studies culminating in the synthesis of macrocycle **19** as a single diastereomer.¹⁴

An efficient synthesis of the alkyl macrocycle 19 was devised as depicted in Scheme 2. The synthesis began with the iodoarene 13 that we recently reported as an intermediate in alternative series of inhibitors.¹³ Allylation of **13** under Stille conditions and subsequent hydrolylsis of the methyl ester provide benzoic acid 14. Carboxylic acid 14 was coupled with (S)-m-allyl phenylalanine 15 using standard conditions to provide the macrocyclic precursor 16. Macrocyclization of 16 was accomplished through the use of second-generation Grubbs catalyst¹⁵ **20** to provide macrocycle **17** as a single regioisomer. The metathesis reaction was high yielding and resulted in exclusive formation of the *E*-isomer, as confirmed by ¹H NMR. Hydrogenation of the olefin followed by further elaboration of the macrocycle was done in a manner similar to **10**. Reduction of the ester using LiBH₄ provided alcohol 18, and Parik-Doering oxidation followed by reductive amination with Nisobutyl-L-norleucinamide afforded the desired alkyl-linked macrocycle 19.

When assayed against BACE-1 enzyme, macrocycle **19** displayed an IC₅₀ = 4 nM. This represents a greater than 8-fold increase in intrinsic activity when compared to compound **10**. Even more impressive, in the cell-based assay, **19** displayed an IC₅₀ = 76 nM, representing a 70-fold improvement over **10** in functional activity. Most gratifying, the apparent permeability of **19** compared with **10** was significantly improved ($P_{app} = 13 \times 10^{-6}$ cm/s). With the improved permeability and reduced P-gp

Scheme 2^a



^{*a*} Reaction conditions: (a) Bu₃SnAllyl, Pd(PPh₃)₄, 82%; (b) NaOH, MeOH/THF, 83%; (c) **15**, BOP, DIPEA, 88%; (d) Grubbs2, CH₂Cl₂, 50 °C, 84%; (e) Pd/C, H₂, MeOH, 93%; (f) LiBH₄, THF, 99%; (g) SO₃-pryidine, TEA, DMSO; (h) *N*-isobutyl-L-norleucinamide, NaBH₃CN, MeOH, 71%, 2 steps.



time (hours)

Figure 4. In vivo A β 40 lowering of 100 mpk iv does of **19** at 1 and 3 h.

susceptibility (BA/AB = 5.5 mdr^{1a}), we dosed macrocycle 19 at 100 mpk in the APP-YAC16 mouse model to obtain a more relevant physiological determination of compound levels in the brain versus plasma. The ratio of compound concentration between the brain and the plasma was determined to be 20-30%, and compound concentrations of 19 in the brain were measured at ~ 1100 nM. With these encouraging results, we undertook a more robust study to evaluate efficacy. APP-YAC mice were administered a 100 mpk iv bolus of 19 and euthanized after 1 and 3 h, respectively, to determine functional efficacy. Analysis of the DEA brain extracts demonstrated a robust decrease in A β (40)^{17,18} levels of 25% when compared to vehicle. To our knowledge, this is the first reported BACE-1 inhibitor to demonstrate efficacy from an iv administration.¹⁹ In the study, the A β reduction was statistically significant in the 1 h grouping compared to the vehicle (Figure 4). However, by the 3 h timepoint, reduction of A β had begun to abate and only a 10% reduction was measurable in the group.²⁰ Analysis of the concentration of 19 in the brain correlated well with reduction



Figure 5. Brain compound levels versus $A\beta$ levels.

in A β levels, and a statistically significant reduction was observed when brain concentrations of >700 nM were achieved (Figure 5).

In summary, we have developed a series of macrocyclic inhibitors of BACE by covalently cross-linking the P1 and P3 side-chains of an isophthalamide-based inhibitor. Macrocyclization resulted in significantly improved physical properties when compared to the initial lead structure. Macrocyclic inhibitors such as **19** also displayed in vivo amyloid lowering when dosed in a murine model.

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

References

- (a) Selkoe, D. J. Translating cell biology into therapeutic advances in Alzheimer's disease. *Nature* **1999**, *399A*, 23. (b) Goate, A.; Chartier-Harlin, M. C.; Mullan, M.; Brown, J.; Crawford, F.; Findani, L.; Giuffra, L.; Haynes, A.; Irving, N.; James, L.; Mant, R.; Newton, P.; Rooke, K.; Roques, P.; Talbot, C.; Pericak, V.; Roses, A.; Williamson, R.; Rossor, M.; Owen, M.; Hardy, J. Segregation of missense mutation in the amyloid presursor protein gene with familial Alzheimer's disease. *Nature* **1991**, *349*, 523–527. (c) Hardy, J. The Alzheimer family of diseases: many etiologies, one pathogenesis? *Proc. Natl. Acad. Sci. U.S.A.* **1997**, *349*, 704–706.
- (2) Sinha, S.; Lieberburg, I. Cellular mechanisms of β-amyloid production and secretion. Proc. Natl. Acad. Sci. U.S.A. 1999, 96, 11049–11053.
- (3) Beher, D.; Graham, S. Protease inhibitors as potential diseasemodifying therapeutics for Alzheimer's disease. *Expert Opin. Invest. Drugs* 2005, 14, 1385–1409.
- (4) Selkoe, D. J. Amyloid β-protein and the genetics of Alzheimer's disease. J. Biol. Chem. 1996, 271, 18295–18298.
- (5) (a) Searfoss, G. H.; Jordan, W. H.; Calligaro, D. O.; Galbreath, E. J.; Schirtzinger, L. M.; Berridge, B. R.; Gao, H.; Higgins, M. A.; May, P. C.; Ryan, T. P. Adipsin: A biomarker of gastrointestinal toxicity mediated by a functional γ-secretase inhibitor. *J. Biol. Chem.* **2003**, 278, 46107–46116. (b) Milano, J.; McKay, J.; Dagenais, C.; Foster-Brown, L.; Pognan, F.; Gadient, R.; Jacobs, R. T.; Zacco, A.; Greenberg, B.; Ciaccio, P. J. Modulation of notch processing by γ-secretase inhibitors causes intestinal goblet cell metaplasia and induction of genes known to specify gut secretory lineage differentiation. *Toxicol. Sci.* **2004**, 82, 341–358.
- (6) (a) Cai, H.; Wang, Y.; McCarthy, D.; Wen, H.; Borchelt, D. R.; Price, D. L.; Wong, P. C. BACE1 is the major beta-secretase for generation of Abeta peptides by neurons. *Nat. Neurosci.* 2001, *4*, 233–234. (b) Roberds, S. L.; Anderson, J.; Basi, G.; Bienkowski, M. J.; Branstetter, D. G.; Chen, K. S.; Freedman, S.; Frigon, N. L.; Games, D.; Hu, K.; Johnson-Wood, K.; Kappenman, K. E.; Kawabe, T.; Kola, I.; Kuehn, R.; Lee, M.; Liu, W.; Motter, R.; Nichols, N. F.; Power, M.; Robertson, D. W.; Schenk, D.; Schoor, M.; Shopp, G. M.; Shuck, M. E.; Sihna, S.; Svensson, K. A.; Tatsuno, G.; Tintrup, H.; Wijsman, J.; Wright, S.; McConlogue, L. BACE knockout mice are healthy despite lacking the primary β-secretase activity in the brain. *Hum. Mol. Genet.* 2001, *10*, 1317–1324.
- (7) Thompson, L. A.; Bronson, J. J.; Zusi, F. C. Progress in the discovery of BACE inhibitors. *Curr. Pharm. Des.* 2005, 11, 3383–3404.

- (8) (a) Stachel, S. J.; Coburn, C. A.,; Steele, T. G.; Jones, K. G.; Loutzenhiser, E. F.; Gregro, A. R.; Rajapakse, H. A.; Lai, M.-T.; Crouthamel, M.-C.; Xu, M.; Tugusheva, K.; Lineberger, J. E.; Pietrak, B. L.; Espeseth, A. S.; Shi, X.-P.; Chem-Dodson, E.; Holloway, M. K.; Munshi, S.; Simon, A. J.; Lawrence, K.; Vacca, J. P. Structure-Based Design of Potent and Selective Cell-Permeable Inhibitors of Human β-Secretase (BACE-1). J. Med. Chem. 2004, 47, 6447–6450. (b) Coburn, C. A.; Stachel, S. J.; Jones, K. G.; Steele, T. G.; Rush, D. M.; DiMuzio, J.; Pietrak, B. L.; Lai, M.-T.; Huang, Q.; Lineberger, J.; Jin, L.; Munshi, S.; Holloway, M. K.; Espeseth, A.; Simon, A. J.; Hazuda, D.; Graham, S. L.; Vacca, J. P. BACE-1 inhibition by a series of Ψ[CH₂NH] reduced amide isosteres *Bioorg. Med. Chem. Lett.* 2006, *16*, 3635–3638.
- (9) Simon, A. J.; Crouthamel, M. C.; Wu, G.; Price, E. A.; Shi, X. P.; Tugusheva, K.; Stachel, S.; Steele, T.; Coburn, C.; Ellis, J.; Jin, L.; Sankaranarayanan, S. BACE1 inhibition after ICV infusion of a potent, cell-permeable small molecule inhibitor: effects of APP metabolism. Presented at 2005 AD/PD meeting, Sorrento, Italy, 2005.
- (10) (a) Tsantrizos, Y. S.; Bolger, G.; Bonneau, P.; Cameron, D. R.; Goudreau, N.; Kukolj, G.; LaPlante, S. R.; Llinas-Brunet, M.; Nar, H.; Lamarre, D. Macrocyclic inhibitors of the NS3 protease as potential therapeutic agents of hepatitis C virus infection. *Angew. Chem., Int. Ed.* 2003, *42*, 1356–1360. (b) Hu, X.; Nguyen, K. T.; Verlinde, C. L. M. J.; Hol, W. G. J.; Pei, D. Structure-based design of a macrocyclic inhibitor for peptide deformylase. *J. Med. Chem.* 2004, *47*, 4941–4949.
- (11) (a) Rojo, I.; Martin, J. A.; Broughton, H.; Timm, D.; Erickson, J.; Yang, H. C.; McCarthy, J. Macrocyclic peptidomimetic inhibitors of β-secretase (BACE): First X-ray structure of a macrocyclic peptidomimetic–BACE complex. *Bioorg. Med. Chem. Lett.* 2006, *16*, 191–195. (b) Hannessian, S.; Yang, G.; Rondeau, J.-M.; Neumann, U.; Betschart, C.; Tintelnot-Blomley, M. Structure-based design and synthesis of macroheterocyclic peptidomimetic inhibitors of the aspartic protease b-site amyloid precursor protein cleaving enzyme (BACE) *J. Med. Chem.* 2006, *49*, 4544–4567.
- (12) The PDB file for BACE-1/inhibitor 1 complex has been deposited with the Protein Data Bank (PDB identifier 2B8L).
- (13) Stachel, S. J.; Coburn, C. A.; Steele, T. G.; Crouthamel, M.-C.; Pietrak, B. L.; Lai, M.-T.; Holloway, M. K.; Munshi, S. K.; Graham,

S. L.; Vacca, J. P. Bioorg. Med. Chem. Lett. 2006, 16, 641-644.

- (14) Brady, S. F.; Singh, S.; Crouthamel, M.; Holloway, M. K.; Coburn, C. A.; Garsky, V. M.; Bogusky, M.; Pennington. M. W.; Vacca, J. P.; Hazuda, D.; Lai, M. Rational design and synthesis of selective BACE-1 inhibitors. *Bioorg. Med. Chem. Lett.* **2004**, *14*, 601-604.
- (15) Scholl, M.; Ding, S.; Lee, C. W.; Grubbs, R. H. Synthesis and activity of a new generation ruthenium-based olefin metahesis catalyst coordinated with 1,3-dimesityl-4,5-dihydroimidazol-2-ylidene ligands. *Org. Lett.* **1999**, *1*, 953–956.
- (16) Lamb, B. T.; Sisodia, S. S.; Lawler, A. M.; Slunt, H. H.; Kitt, C. A.; Kearns, W. G.; Pearson, P. L.; Price, D. L.; Gearhart, J. D. *Nat. Genet.* **1993**, *5*, 22–29.
- (17) Best, B. T.; Jay, M. T.; Out, F.; Ma, J.; Nadin, A.; Ellis, S.; Lewis, H. D.; Pattison, C.; Reilly, M.; Harrison, T.; Sherman, M. S.; Williamson, T. L.; Atack, J. R. J. Quantitative measurement of changes in amyloid-β(40) in the rat brain and cerebrospinal fluid following treatment with the γ-secretase inhibitor LY-411575 [N²-[(2S)-2-)3,5-difluorophenyl)-2-hydroxyethanoyl-N¹-[(7S)-5-methyll-6-oxo-6,7-dihydro-5H-dibenzo[b,d]azepin-7-yl]-l-alaninamide] Pharmacol. Exp. Ther. **2005**, 313, 902–908.
- (18) Savage, M. J.; Trusko, S. P.; Howland, D. S.; Pinsker, L. R.; Mistretta, S.; Reaume, A. G.; Greenberg, B. D.; Siman, R.; Scott, R. W. Turnover of amyloid β-protein in mouse brain and acute reduction of its level by phorbol ester *J. Neurosci.* **1998**, *18*, 1743–1752.
- (19) Biological data of a closely related analogue in this class was recently disclosed at the 10th International conference for Alzheimer's disease and related disorders. Sankaranarayanan, S.; Price, E. A.; Wu, G.; Crouthamel, M. C.; Ellis, J. D.; Jin, L.; Stachel, S. J.; Graham, S. L.; Vacca, J. P.; Coburn, C. A.; Simon, A. J. Presented at the 10th International conference for Alzheimer's disease and related disorders, Madrid, Spain, July, 2006; poster #3050.
- (20) The turnover of $A\beta$ is reported to be approximately 2 h. This is in accord with the rapid recovery when compound levels fall to subefficacious levels.

JM060884I