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Tripeptidyl-peptidase II (TPP II) Inhibitory Activity of (S)-2,3-Dihydro-2-(1H-imidazol-2-yl)-1H-indoles, a Systematic SAR Evaluation. Part 2

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Abstract—We have systematically explored the structure–activity relationship (SAR) for a series of compounds **2** as inhibitors of tripeptidyl-peptidase II (TPP II), a serine protease responsible for the degradation of cholecystokinin-8 (CCK-8). This SAR evaluation of the core structure **2** suggest a fairly restrictive pharmacophore for such related structures, but has yielded a limited set of compounds (**2b**, **2c**, **2d**, **2s**, and **2t**) with potent TPP II inhibitory activity (IC₅₀ 4-11 nM). © 2003 Elsevier Ltd. All rights reserved.

As of 1998, 18% of the US population was defined as being obese. which is a serious concern since obesity has been correlated as a causative contributor to numerous diseases and disorders.² The endogenous octa-peptide cholecystokinin-8 (CCK-8), as well as synthetic CCK agonists, exhibit pro-satiety effects, suggesting that CCK or related agents might prove useful in helping to address the obesity problem.³ In 1996 Rose et al. identified Tripeptidyl-peptidase II (TPP II; EC 3.4.14.10) as an endogenous serine protease responsible for the degradation of CCK-8, and also identified butabindide, 1, as a potent inhibitor of TPP II ($IC_{50}=7$ nM).4 Subsequently, Rose and coworkers demonstrated that treatment of mice with 1 resulted in a 50% reduced food intake over a 1-h period for treated animals relative to control animals, which is consistent with the expected result if endogenous levels of CCK-8 were elevated.

Recently we reported that 1 possesses some inherent chemical instability, with a propensity to intra-molecularly cyclize to yield butyl amine and 1's respective diketopiperazine ($IC_{50} > 10 \mu M$).⁵ Based on this finding, we subsequently prepared a small set of compounds 2 in which an imidazole served as a bio-isostere equivalent for 1's labile amide moiety.

Imidazole **2c** (IC $_{50}$ =4 nM) proved equally potent to **1** as a TPP II inhibitor while being favorably void of the intra-molecular cyclization liability of **1**. Our initially reported efforts included preparing only closely related derivatives of **1**, as a primary test of our hypothesis that compounds such as **2** may prove promising. Based on the favorable results for our initial analogues **2**, we have prepared additional heterocyclic analogues to more thoroughly evaluate the TPP II inhibitory structure–activity relationships (SARs). Below is described our related biological findings, as well as associated chemical synthesis.

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To initiate our SAR work, we retro-synthetically evaluated our generic structure 2. The resulting segmentation of the parent compound at good synthetic junction points led us to trisect the structure, which we arbitrarily labeled A-C, as outlined above for 2.

Segment A variations of parent structure 2 proved generally the most simplistic and expedient to alter, relative to segments B and C. Changes explored in segment A were generally one to three steps from common intermediate noramines 5d and 5f, whose preparations and chemical characterizations we previously described in detail.⁵ As shown in Scheme 1, we reacted secondary amines 5d and/or 5f, or their respective racemic mixtures 5e and/or 5g, under standard amide coupling conditions with generated mixed anhydrides of various commercially available N-BOC protected amino acids. Resulting amino acid adducts were simply N-deprotected to yield desired products 2a-i, 2o, and 2p. Alternatively, noramine 5g was reacted under standard Schotten-Baumann conditions with acid chlorides 6a, **6b**, and **6c** to give products **2j**, **2k**, and **2m**, respectively. Anilines 21 and 2n were prepared via standard room temperature PtO₂/H₂ hydrogenation conditions from these respective nitro precursors 2k and 2m. Product 2q was generated from 2p under reductive diborane. THF conditions in refluxing THF. Full chemical characterizations (MS, ¹H NMR, TLC, LC) were completed for this set of final targets, as well as those described below for Segments B and C, and are consistent with expectations based on comparisons to related analogues previously described in detail.⁵

For segment B variations, we began by preparing a handful of 5-substituted indolines in lieu of our previously reported un-substituted indoline analogues. As shown in Scheme 1, the synthesis followed to prepare the new 5-substituted indolines 2r-v was virtually identical to our previously reported eight step route for the unsubstituted indolines, with the simple modification of starting from 5-substituted starting material indolines 4a-c. The 5-substituted indolines 4a-c were prepared following a previously reported Mg/MeOH reduction of their respective indoles 3a-c. Indolines 4a-c led to

racemic products 2r, 2s, and 2u when Gly was used in the amide coupling step of the synthetic sequence. When L-Ala was used in the amide coupling step with 5b and **5c**, mixtures of epimers were generated, which were N-BOC de-protected as mixtures with TFA. The resulting desired final products 2t and 2v were easily isolated as pure compounds from their respective epimers via preparative HPLC. The absolute configurations for 2t and 2v were assigned based on comparison of their NMRs to the unsubstituted indoline 2p. Compound 2p, and its respective epimer, were prepared via stereospecific syntheses.⁵ For a cursory look at a different scaffold in place of the indoline scaffold of Segment B, we slightly altered the indoline synthesis to generate tetrahydroquinolines. As shown in Scheme 2, we formed our imidazole ring adjacent to the aromatic quinoline and then reduced 8 to its respective tetrahydroquinoline intermediate 9. Noramine 9 was then coupled with N-BOC-L-Ala-F⁷ 10, and the resulting epimers 11a and 11b were separated by preparative HPLC. Subsequent N-BOC de-protection of these purified intermediates yielded epimers 2w and 2x. Absolute assignment of configurations for these two products was not pursued since these compounds lacked significant biological activity.

Scheme 2.

Work around segment C included substitutions on and for the imidazole heterocycle. As shown in Scheme 3, a simple iodomethane *N*-substitution reaction on imidazole intermediate 12⁵ yielded both regioisomers 13 and 14. These regioisomer intermediates were separated via preparative HPLC and their absolute stereochemistries assigned based on comparison with similar *N*-sub-

stituted imidazoles which have been well characterized.8 Intermediates 13 and 14 were then independently hydrolyzed with refluxing 6N aqueous HCl to their respective noramines, which were carried on to their final targets 2v and 2z following similar reaction conditions described above for segment A analogues. Trisubstituted imidazole analogues 2cc-2hh were prepared by an alternate synthetic route, as depicted in Scheme 4. Indoline alcohol 15⁹ was oxidized to aldehyde 16, which was subsequently subjected to Radziszewski reaction conditions¹⁰ with diketones 17a and 17b¹¹ to yield respective imidazoles 18a and 18b. The BOC groups of 18a and 18b were removed with TFA, generating noramines 19a and 19b, which were coupled with BOC-Gly-F.⁷ The resulting amides **20a** and **20b** were then treated with TFA to yield desired products 2cc and 2dd. Related analogues 2ee and 2ff were prepared following the same Scheme, after appropriate manipulation of ester 18b. This entailed hydrolyzing ester 18b with 1 N KOH in MeOH and subjecting the resulting acid 18c to standard amide coupling conditions. The resulting 4substituted amide-imidazoles 18d were then carried on through their respective intermediates 19 and 20 to generate products 2ee and 2ff. Scheme 5 outlines the preparation of benzimidazole noramine 24, the intermediate employed to prepare final targets 2gg and 2hh. Noramine 24 was subjected to coupling and deprotecting reactions similar to those described for analogues shown in Scheme 1. Also outlined in Scheme 5 is the preparation of some other heterocycles we prepared in lieu of an imidazole.

Related intermediates 26, 27, and 29 were hydrolyzed to their respective noramines, then also coupled and deprotected in a similar manner as described for the

Scheme 3.

imidazoles of Scheme 1, ultimately yielding final targets 2ii, 2jj, and 2kk, respectively.

The related TPP II biological results for the diverse set of compounds prepared are summarized in Tables 1 and 2.¹² Evaluation of the SAR for analogues **2a–2f**, where the amino acid α appendage in segment A was the sole variation, suggest that limited lipophilicity is desirable to enhance activities [H (2a; IC_{50} 36 nM) < Me (2b; IC_{50} 6 nM) \sim Et (**2c**; IC₅₀ 4 nM) \sim Pr (**2d**; IC₅₀ 4 nM)>*i*-Pr (2e; IC_{50} 20 nM)>Bn (2f; IC_{50} 2300 nM)]. Chemical manipulations of the primary amine of **2b** (IC₅₀ 6 nM) proved disastrous in terms of maintaining favorable biological activity. Explorations included a simple N-Me substitution (2h; IC₅₀>1000 nM), a Pro-derived cyclic amine (2g; IC₅₀ 1400 nM), and insertion of an extra methylene between the carbonyl and the amine (2i; $IC_{50} > 10,000$ nM). We also found that reduction of the amide carbonyl moiety of analogue 2p led to a loss of desired biological activity [2q $(IC_{50} > 10,000 \text{ nM}) < 2p \text{ (IC}_{50} 23 \text{ nM})].$ Additional unnatural amino acids derivatives were also inactive, possessing IC₅₀s > 10,000 nM (2j, 2l, 2n). Evaluations of segment B of parent structure 2 included a cursory evaluation of 5-indoline substitutions. The related indoline SAR suggest that some variation is tolerable at the 5-position, although this substitution did not significantly enhance the desired biological activities [Set 1: Gly derivatives—OMe (2u; IC₅₀ 70 nM) < H (2a; IC₅₀ 36 nM)~F (2r; IC₅₀ 46 nM) <Cl (2s; IC₅₀ 11 nM); Set 2: Ala derivatives—OMe (2v; IC₅₀ 16 nM) \leq H (2b; IC₅₀ 6 nM) \sim Cl (2t; IC₅₀ 5 nM). Also related to segment B, we substituted the indoline heterocycle with a tetrahydroquinoline and lost most desired activity [2w (IC₅₀ 400 nM) < $2\mathbf{b}$; (IC₅₀ 6 nM)]. For segment C of the parent molecule 2, we found that a simple N-Me substitution at either imidazole nitrogen yielded significantly weaker TPP II inhibitors than their parent NH analogue **20** [**2y** (IC₅₀ > 10,000 nM) < **2z** (IC₅₀ 2,100 nM) <20 (IC₅₀ 194 nM)]. Replacement of the imidazole group with alternate five-membered ring heterocycles also proved less promising biologically, as readily seen by comparing 2ii and 2kk of Table 2 with imidazole 2a (IC₅₀ 36 nM), and **2jj** of Table 2 with imidazole **2o** (IC₅₀ 194 nM). Fusing of the pendant propyl group with the imidazole in **2b** (IC₅₀ 6 nM), to give benzimidazole derivative 2gg (IC₅₀ 1400 nM), also proved unfavorable for TPP II inhibitory activity.

$$\begin{array}{c} \text{Dess-Martin} \\ \text{OH} \\ \text{Reagent} \\ \text{15} \\ \text{BOC} \\ \text{91}\% \\ \text{16} \\ \text{BOC} \\ \text{18} \\ \text{BOC} \\ \text{CR}^{\parallel} = \text{n-Pr}; \, \text{R}^{\parallel} = \text{Me} \\ \text{MR}^{\parallel} = \text{n-Pr}; \, \text{R}^{\parallel} = \text{CO}_2 \text{Et} \\ \text{CR}^{\parallel} = \text{n-Pr}; \, \text{R}^{\parallel} = \text{CO}_2 \text{H} \\ \text{MR}^{\parallel} \\ \text{R}^{\parallel} \\ \text{20} \\ \text{2dd from 20b (50\%)} \\ \text{2dd from 20b (50\%)} \\ \text{3} \\ \text{4} \\ \text{R}^{\parallel} = \text{n-Pr}; \, \text{R}^{\parallel} = \text{Me} \\ \text{3} \\ \text{8} \\ \text{16} \\ \text{17} \\ \text{18} \\ \text{18} \\ \text{18} \\ \text{18} \\ \text{19} \\ \text{10} \\ \text{10$$

Scheme 5.

Table 1. Imidazoles, benzimidazoles and related TPP II IC₅₀'s

Compd	n	X	R	R^{I}	R^{II}	R^{III}	IC ₅₀ , (nM) ^a
2a	1	Н	Gly ^{b,5}	Н	n-Pr	Н	36
2b	1	Н	L-Åla⁵	Н	n-Pr	Н	6
2c	1	Н	L-Abu ^c ,5	Н	n-Pr	Н	4
2d	1	H	L-Norvaline	Н	<i>n</i> -Pr	Н	4
2e	1	H	L-Val	Н	<i>n</i> -Pr	Н	20
2f	1	H	L-Phe	Н	<i>n</i> -Pr	Н	2300
2g	1	H	L-Pro	Н	<i>n</i> -Pr	Н	1400
2h	1	H	L-N-Me-Ala	Н	<i>n</i> -Pr	Н	> 10,000
2i	1	H	β-Ala	Н	<i>n</i> -Pr	Н	> 10,000
2j	1	H	-CO-4-imidazole ^b	Н	<i>n</i> -Pr	Н	> 10,000
2k	1	H	2-Nitro-benzoyl ^b	Н	<i>n</i> -Pr	Н	> 10,000
21	1	H	2-Amino-benzoyl ^b	Н	<i>n</i> -Pr	Н	> 10,000
2m	1	H	3-Nitro-benzoyl ^b	Н	<i>n</i> -Pr	Н	> 10,000
2n	1	H	3-Amino-benzoyl ^b	Н	<i>n</i> -Pr	Н	> 10,000
2o	1	H	Gly^b	Н	Et	Н	194
2p	1	H	L-Ala⁵	Н	Et	Н	23
2q	1	H	-CH ₂ -CH(Me)NH ₂	Н	Et	Н	> 10,000
2r	1	F	Glyb	Н	n-Pr	Н	46
2s	1	Cl	Gly^b	Н	<i>n</i> -Pr	Н	11
2t	1	Cl	L-Ala	Н	n-Pr	Н	5
2u	1	OMe	Gly ^b	Н	<i>n</i> -Pr	Н	70
2v	1	OMe	L-Ala	Н	<i>n</i> -Pr	Н	16
2w	2	H	L-Ala ^d	Н	<i>n</i> -Pr	Н	400
2x	2	H	L-Ala ^d	Н	<i>n</i> -Pr	Н	800
2y	1	Н	Gly	Me	Et	Н	> 10,000
2z	1	Н	Gly	Me	H	Et	2100
2aa	1	Н	$\mathrm{Gly^b}$	Н	n-Bu	Н	50
2bb	1	Н	Gly^b	Н	Ph	Н	> 10,000
2cc	1	Н	Gly^b	Н	n-Pr	Me	55
2dd	1	Н	Gly^b	Н	n-Pr	CO ₂ Et	210
2ee	1	Н	Gly^b	Н	n-Pr	$CONH_2$	200
2ff	1	Н	Gly^b	Н	n-Pr	CO-Gly-OMe	135
2gg	1	H	L-Ala	H		Fused Ph	1400
2hh	1	Н	L-Abu	Н		Fused Ph	450

 $^{{}^{\}rm a}{
m IC}_{50}$ values were determined using Sigma plot C/R curves.

Table 2. Additional heterocycles and related TPP II IC₅₀'s

Compd	R	Imidazole replacement	R^{I}	R^{II}	R ^{III}	IC ₅₀ , (nM) ^a
2ii (from 26)	Gly ^b	Triazole	H	n-Pr	Н	> 10,000
2jj (from 27)	Gly ^b	Thiazole	H	Et	Н	7700
2kk (from 29)	Gly ^b	Oxadiazole	H	n-Pr	Н	2200

 $^{{}^{\}rm a}{\rm IC}_{50}$ values were determined using Sigma plot C/R curves.

^bRacemic and not S stereochemistry as depicted in Markush structure 2.

^cAbu, 2-aminobutyric acid.

^dSeparated epimers, absolute stereochemistries unassigned.

^bRacemic.

We also explored the SAR for a small set of varied 2,4-di- and 2,4,5-tri-substituted imidazoles. For 2,4-di-substituted imidazoles, we found that limited aliphatic lipophilicity is desirable at the 4-imidazole position, based on comparison of a set of R^{II} mono varied analogues [Et (2σ ; IC₅₀ 194 nM) < n-Pr (2a; IC₅₀ 36 nM) $\sim n$ -Bu (2aa; IC₅₀ 50 nM)> > Ph (2d; IC₅₀ 1>10,000 nM)]. Our evaluations of 2,4,5-trisubstituted analogues suggest that some bulk, and even some polarity, is biologically tolerable for such associated analogues, although some gradations of desired activities were noticed. Again, this assessment was based on the SAR for a set of monovaried analogues, where R^{II} was held constant as an n-Pr group [R^{III} = H (2a; IC₅₀ 36 nM) \geq R^{III} = Me (2cc; IC₅₀ 55 nM) > R^{III} = CO–NH–Gly–OMe (2ff; IC₅₀ 135 nM) \geq R^{III} = CO–NH₂ (2ee; IC₅₀ 200 nM) \sim R^{III} = CO₂Et (2dd; IC₅₀ 210 nM)].

In conclusion, we have systematically evaluated the SAR of our lead TPP II inhibitors 2. We have found that for the maintenance of good TPP II inhibitory activity the latitude of structural variations on scaffold 2 is rather limited in nature, with our most potent TPP II inhibitors being identified as 2b, 2c, 2d, 2s, and 2t. The limitations noted around the SAR of 2 are consistent with molecular modeling results that we have performed within our group, and these related modeling results will be published in detail in the near future.

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