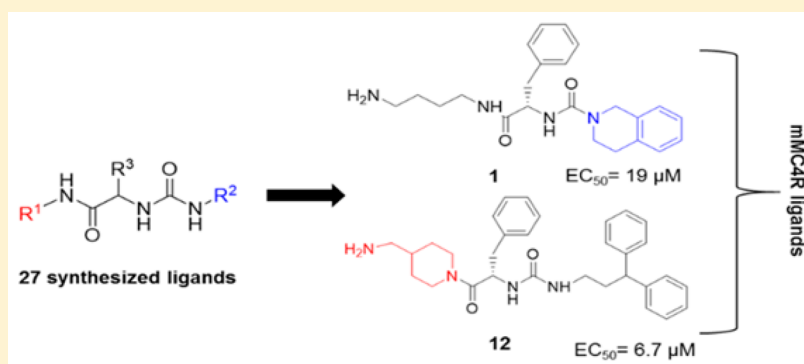


Synthesis and Structure–Activity Relationships of Substituted Urea Derivatives on Mouse Melanocortin Receptors

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S Supporting Information



ABSTRACT: The melanocortin system is involved in the regulation of several complex physiological functions. In particular, the melanocortin-3 and -4 receptors (MC3R/MC4R) have been demonstrated to regulate body weight, energy homeostasis, and feeding behavior. Synthetic and endogenous melanocortin agonists have been shown to be anorexigenic in rodent models. Herein, we report synthesis and structure–activity relationship (SAR) studies of 27 nonpeptide small molecule ligands based on an unsymmetrical substituted urea core. Three templates containing key residues from the lead compounds, showing diversity at three positions (R¹, R², R³), were designed and synthesized. The syntheses were optimized for efficient microwave-assisted chemistry that significantly reduced total syntheses time compared to a previously reported room temperature method. The pharmacological characterization of the compounds on the mouse melanocortin receptors identified compounds 1 and 12 with full agonist activity at the mMC4R, but no activity was observed at the mMC3R when tested up to 100 μM concentrations. The SAR identified compounds possessing aliphatic or saturated cyclic amines at the R¹ position, bulky aromatic groups at the R² position, and benzyl group at the R³ position resulted in mMC4R selectivity over the mMC3R. The small molecule template and SAR knowledge from this series may be helpful in further design of MC3R/MC4R selective small molecule ligands.

KEYWORDS: Melanocortin receptors, melanocortin agonist, urea ligands, small molecules, melanocortin-3 receptor, melanocortin-4 receptor

The melanocortin receptors (MCRs) belong to the family of seven transmembrane G protein-coupled receptors (GPCRs) and signal through the 3',5'-cyclic adenosine monophosphate (cAMP) signal transduction pathway. The five known melanocortin receptor subtypes (MC1R–MC5R)^{1–8} interact with the endogenous peptides to mediate several complex physiological functions including weight and energy homeostasis,⁹ feeding behavior,¹⁰ skin pigmentation,¹¹ steroidogenesis,¹² cardiovascular,¹³ and erectile function.^{14–16} The endogenous agonists for these receptors, α -, β -, and γ -melanocortin stimulating hormone (MSH) and adrenocorticotropic hormone (ACTH), are peptide hormones that are produced by post-translational processing of the pro-opiomelanocortin (POMC) gene.¹⁷ These peptides contain a conserved core sequence His⁶-Phe⁷-Arg⁸-Trp⁹ (Figure 1; α -MSH numbering) postulated to be important for melanocortin ligand–receptor recognition and activation.¹⁸ The melanocor-

tin receptors respond to all POMC-derived melanocortin peptides, albeit to varying degrees of potencies. The only exception to this is the MC2R, which only responds to ACTH but not to other MSH agonists.¹⁹

The melanocortin-3 (MC3R) and -4 receptors (MC4R) are expressed in the brain and have been demonstrated to be involved in the regulation of feeding behavior, metabolism, and energy homeostasis, as well as, erectile and cardiovascular functions.^{9,10,13,15,20–22} The role of the MC4R in feeding behavior is well documented through the use of nonselective and selective ligands and by genetic knockout studies.^{9,10,23} Activation of the MC4R via endogenous or synthetic agonists decrease food intake.¹⁰ The importance of this receptor in food

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ACTH (1-24)	YSMEHFRWGKPVGKKRRPVKVYP
α -MSH	Ac-SYSMEHFRWGKPV-NH ₂
β -MSH	AEKKDEGPYRMEHFRWGSPPKD-OH
γ -MSH	YVMGHFRWDRFG-OH

Figure 1. Sequences of the endogenous melanocortin peptides. The core His, Phe, Arg, and Trp residues amino acids are conserved in all melanocortin peptides, and this is highlighted in the sequences.

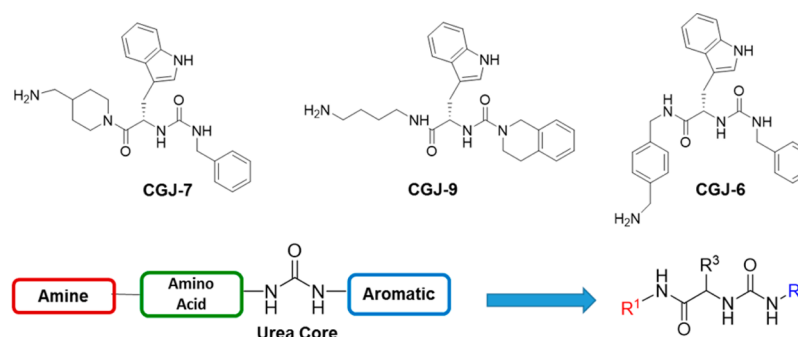


Figure 2. Example of common features of the substituted urea derivatives and structures of reported compounds CGJ-7, CGJ-9, and CGJ-6.³⁴

intake and energy homeostasis is also highlighted by naturally occurring mutations identified in morbidly obese individuals²⁴ that result in partial or total dysfunction of the MC4Rs *in vitro*. It has been reported that single nucleotide polymorphisms (SNPs) in the MC4R may be responsible for morbid obesity in ~6% children and adults.²⁴ The clinical symptoms include lack of satiety, extreme hyperphagia, decrease energy expenditure, increased linear growth, and hyperinsulinemia. The MC3R is postulated to be involved in the regulation of energy and fat homeostasis. However, the underlying mechanism of the MC3R action has not been fully elucidated.^{20–22} Therefore, discovery of receptor specific ligands for the MC3R and MC4R are vital for the advancement in the understanding of their overlapping physiological roles and to develop potential treatment for eating disorders.

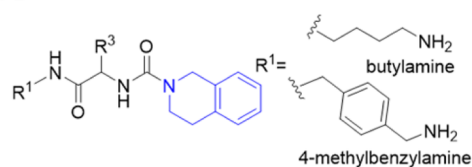
The endogenous α -MSH is a 13 amino acid long peptide agonist for the melanocortin receptors (MC1R, MC3–5Rs). Structure–activity relationship (SAR) studies of the α -MSH sequence identified an analog containing the Nle and DPhe amino acids substituted at positions 4 and 7, respectively [(Nle⁴, DPhe⁷) α -MSH, NDP-MSH].²⁵ This compound is a more potent MCR agonist than α -MSH and has emerged as a valuable biochemical tool in receptor mediated studies. However, endogenous and synthetic peptidic MSH ligands are susceptible to enzymatic degradation *in vivo*.²⁶ The poor stability and lack of selectivity limit the use of these peptides to probe and characterize the pharmacological and physiological functions of melanocortin receptors. Rational design based on the conserved peptide core of the melanocortins has formed the basis of designing MC3R/MC4R selective small molecule ligands.

Utilization of nonpeptidic scaffolds has been explored in the past to achieve selectivity and to improve pharmacokinetic properties at the MCRs. Several classes of small molecule scaffolds have been explored for melanocortin receptors including β -turn related motifs, benzodiazepines, benzimidazole pyridazinone, cyclohexylpiperidine, and urea derived compounds.^{27–35} In particular, substituted urea derivatives are an important class of carbonyl compounds that are building blocks

of several drug candidates such as HIV protease inhibitors,³⁶ Raf kinase inhibitors,³⁷ CCK-B receptor antagonists,^{38,39} and endothelin antagonists.⁴⁰ An orally bioavailable piperazine urea based compound (a selective MC4R partial agonist) has been reported to modulate food intake in animal models without any effect on erectile activity.⁴¹ An earlier study of a “focused” library based on the core tripeptide sequence, “Phe-Trp-Lys” that included a urea linkage resulted in identification of lead compounds CGJ-7 and CGJ-9 (Figure 2).³⁴ Notably, compound CGJ-9, containing the 1,2,3,4-tetrahydroisoquinoline (TIQ) functionality displayed full agonist activity at both the mMC3R and mMC4Rs. However, compound CGJ-7, containing 4-(aminomethyl)piperidine (AMP) functionality resulted in a 4 μ M agonist at the mMC3R and was unable to stimulate the mMC4R at up to 100 μ M concentrations. The present study is based on the interesting pharmacology of these two compounds. Three substituted urea-based templates were designed herein to discover small molecules that can be used to probe the molecular mechanisms of the MC3/MC4 receptors. The synthetic strategy was modified from the literature to adopt microwave technology and resulted in efficient overall synthesis.

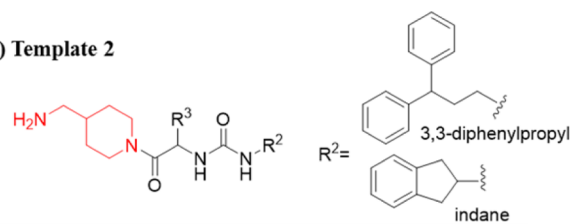
Template Design. The design strategy included a three component approach including a urea linkage as a core with an amine functionality (R¹) linked to an amino acid residue (R³) at one end and an aromatic group (R²) on the other side as delineated in Figure 2. Template 1 was designed based on the compound CGJ-9 with TIQ moiety at R² and variable amino acids (R³) and amine (R¹) functionalities (Figure 3A). Template 2 contained the fixed AMP group in R¹ position as the amine component (based on CGJ-7) and positions R² and R³ were varied (Figure 3B). Template 3 was designed as a compilation of CGJ-7 and CGJ-9, with the AMP group positioned at R¹, the TIQ moiety at R², and amino acid substitutions at R³ (Figure 3C). The amino acid substitutions at R³ were chosen based on the earlier SAR studies of melanocortin agonists that resulted in potent and/or selective pharmacology at MCRs in different templates. It is hypothesized that diversifying the three components of the urea core

A) Template 1



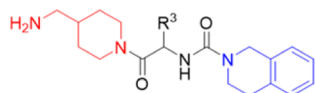
Compound	R ¹	R ³
1	butylamine	Phe
2	butylamine	(<i>p</i> I)Phe
3	butylamine	(<i>p</i> Cl)Phe
4	butylamine	His
5	butylamine	Nal(2')
6	4-methylbenzylamine	Phe
7	4-methylbenzylamine	(<i>p</i> I)Phe
8	4-methylbenzylamine	(<i>p</i> Cl)Phe
9	4-methylbenzylamine	His
10	4-methylbenzylamine	Nal(2')

B) Template 2



Compound	R ²	R ³
11	3,3-diphenylpropyl	Trp
12	3,3-diphenylpropyl	Phe
13	3,3-diphenylpropyl	(<i>p</i> I)Phe
14	3,3-diphenylpropyl	(<i>p</i> Cl)Phe
15	3,3-diphenylpropyl	His
16	3,3-diphenylpropyl	Nal(2')
17	indane	Trp
18	indane	Phe
19	indane	(<i>p</i> I)Phe
20	indane	(<i>p</i> Cl)Phe
21	indane	His

C) Template 3



Compound	R ³
22	Trp
23	Phe
24	(<i>p</i> I)Phe
25	(<i>p</i> Cl)Phe
26	His
27	Nal(2')

Figure 3. Illustration of the templates and list of compounds in each template in this study. (A) Template 1, (B) Template 2, and (C) Template 3.

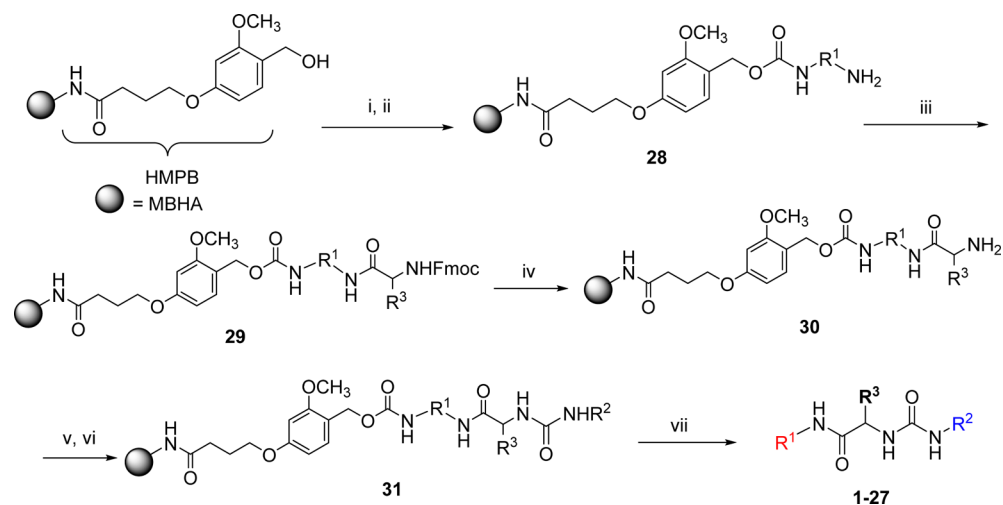
may result in more potent and/or selective melanocortin receptor ligands.

RESULTS AND DISCUSSION

Herein, the synthesis and pharmacology of substituted urea derivatives in cells expressing the cloned mouse MC1, MC3, MC4 and MC5 receptors is reported. Twenty seven compounds were designed and synthesized in total, of which 10 compounds (1–10) were part of Template 1 (Figure 3A), 11 compounds (11–21) belonged to Template 2 (Figure 3B), and 6 compounds (22–27) were part of Template 3 (Figure 3C). The solid phase syntheses of the urea compounds were modified from an earlier reported method (Scheme 1).^{34,42,43} The synthesis was optimized for a manual microwave synthesizer, which dramatically reduced the reaction time to 3 h compared to the several days in the earlier method at room temperature.³⁴ The HMPB-MBHA resin was activated with *p*-nitrophenyl chloroformate first and was coupled with the diamine subunit (R¹) through a carbamate linkage under microwave irradiation. The Fmoc-protected amino acid subunit (R³) was then added to the amine using a standard microwave SPPS protocol with coupling reagents HBTU and DIEA,

followed by microwave assisted Fmoc deprotection with 20% piperidine. The aromatic part of the template (R², Figure 2) was introduced by coupling the amine subunit (R²) through a urea linkage using *p*-nitrophenyl chloroformate as a carbonylating agent. The product was cleaved from the resin by treatment with glacial acetic acid using a microwave irradiation. The synthesized compounds were purified by semipreparative RP-HPLC and analyzed for the correct molecular weights by mass spectrometry and the purity by analytical RP-HPLC in two divergent solvent systems (Supporting Information). All compounds were found to be ≥95% pure. The purified compounds were tested for functional activity using a β -galactosidase reporter gene bioassay⁴⁴ on HEK cells stably expressing the mouse melanocortin receptors (mMC1R, mMC3–5Rs). The MC2R only responds to ACTH and therefore, it was not included in this study.

Template 1. The agonist EC₅₀ values of compounds 1–10 (Template 1) at the mouse melanocortin receptors are summarized in Table 1, along with the known agonist NDP-MSH and leads CGJ-7 and CGJ-9. Compound CGJ-9 was the lead compound for the Template 1 having Trp at the R³ position, TIQ at the R² position and butyl amine at the R¹ position and possessed low micromolar activity at all MCRs

Scheme 1. General Scheme for Microwave-Assisted Urea Based Library Synthesis^a

^aReagents and conditions: (i) *p*-nitrophenyl chloroformate (18.7 equiv), DIEA (18.7 equiv), mW (35 °C, 30 W, 5 min); (ii) diamine, R¹ (12.5 equiv), DIEA (12.5 equiv), DMF, mW (55 °C, 50 W, 10 min); (iii) Fmoc-amino acid, R³, HBTU, DIEA, mW (75 °C, 30 W, 5 min); (iv) 20% piperidine in DMF, mW (75 °C, 30 W, 4 min); (v) *p*-nitrophenyl chloroformate (1.4 equiv), DIEA (1.4 equiv), mW (35 °C, 30 W, 5 min); (vi) amine, R² (1.4 equiv), DIEA (1.4 equiv), DMF, mW (55 °C, 50 W, 10 min); (vii) glacial acetic acid, mW (38 °C, 20 W, 30 min).

Table 1. Pharmacology of Template 1 Urea Compounds at the Mouse Melanocortin Receptors^a

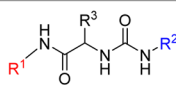
Compounds				mMC1R	mMC3R	mMC4R	mMC5R
	R ¹	R ²	R ³				
NDP-MSH				0.05 ± 0.01 nM	0.19 ± 0.04 nM	0.06 ± 0.01 nM	0.20 ± 0.05 nM
CGJ-9*	butylamine	TIQ	Trp	7.9 ± 1.0	1.7 ± 1.0	3.4 ± 1.9	4.9 ± 4.8
CGJ-7*	4-AMP	benzyl	Trp	5.2 ± 2.7	4.4 ± 2.8	> 100	8.2 ± 7.9
1	butylamine	TIQ	Phe	3.8 ± 0.3	> 100	19 ± 2.0	22 ± 4.1
2	butylamine	TIQ	(<i>p</i> I)Phe	25 ± 2.8	37 ± 19	17 ± 4.3	43 ± 11
3	butylamine	TIQ	(<i>p</i> Cl)Phe	> 100	> 100	> 100	> 100
4	butylamine	TIQ	His	50% @ 100 μM	> 100	60% @ 100 μM	80% @ 100 μM
5	butylamine	TIQ	Nal(2')	> 100	> 100	> 100	> 100
6	4-MBA	TIQ	Phe	60% @ 100 μM	> 100	> 100	70% @ 100 μM
7	4-MBA	TIQ	(<i>p</i> I)Phe	> 100	> 100	> 100	> 100
8	4-MBA	TIQ	(<i>p</i> Cl)Phe	> 100	> 100	> 100	> 100
9	4-MBA	TIQ	His	> 100	> 100	> 100	40% @ 100 μM
10	4-MBA	TIQ	Nal(2')	> 100	> 100	> 100	> 100

^aThe indicated errors represent the standard error of the mean determined from at least three independent experiments. *The values for compounds have previously been reported and included here for comparison.³⁴ >100 indicates that the compound was examined but lacked agonist activity at up to 100 μM concentrations. A percentage (%) indicates the percent maximal stimulatory response observed at 100 μM concentrations observed, but not enough stimulation was observed to determine an EC₅₀ value.

tested.³⁴ Compounds 1–10 had modifications at the R¹ and R³ positions compared to CGJ-9, while R² was fixed as TIQ. TIQ residue has previously been involved in the design of compounds that possessed MC4R agonist pharmacology. The potent MC4R compound first reported by Merck contained a TIQ moiety.³³ More recently, several peptides containing a TIQ residue have been identified to rescue receptor function

and signaling in genetically mutated MC4 receptor.⁴⁵ Compounds 1–5 had the R¹ position substituted with butyl amine and compounds 6–10 contained 4-methylbenzylamine (MBA) group at this position. The R³ position of compounds 1–5 contained the amino acid residues Phe, (*p*I)Phe, (*p*Cl)Phe, His, and Nal(2') residues, respectively. These amino acids have been previously incorporated into the potent melanocortin

Table 2. Pharmacology of Template 2 Urea Compounds at the Mouse Melanocortin Receptors^a

Compounds				mMC1R	mMC3R	mMC4R	mMC5R
	R ¹	R ²	R ³				
NDP-MSH				0.05 ± 0.01 nM	0.19 ± 0.04 nM	0.06 ± 0.01 nM	0.20 ± 0.05 nM
CGJ-9*	butylamine	TIQ	Trp	7.9 ± 1.0	1.7 ± 1.0	3.4 ± 1.9	4.9 ± 4.8
CGJ-7*	4-AMP	benzyl	Trp	5.2 ± 2.7	4.4 ± 2.8	> 100	8.2 ± 7.9
11	4-AMP	3,3-DPP	Trp	14 ± 0.8	40% @ 100 μM	> 100	> 100
12	4-AMP	3,3-DPP	Phe	6.9 ± 2.0	> 100	6.7 ± 3.8	7.7 ± 1.8
13	4-AMP	3,3-DPP	(<i>pI</i>)Phe	> 100	> 100	> 100	> 100
14	4-AMP	3,3-DPP	(<i>pCl</i>)Phe	> 100	> 100	> 100	> 100
15	4-AMP	3,3-DPP	His	> 100	> 100	> 100	> 100
16	4-AMP	3,3-DPP	Nal(2')	> 100	> 100	> 100	> 100
17	4-AMP	indane	Trp	14.2 ± 5.1	27.2 ± 10.2	9.1 ± 3.2	16.9 ± 7.3
18	4-AMP	indane	Phe	80% @ 100 μM	> 100	60% @ 100 μM	70% @ 100 μM
19	4-AMP	indane	(<i>pI</i>)Phe	30% @ 100 μM	> 100	60% @ 100 μM	70% @ 100 μM
20	4-AMP	indane	(<i>pCl</i>)Phe	30% @ 100 μM	> 100	> 100	> 100
21	4-AMP	indane	His	> 100	> 100	> 100	> 100

^aThe indicated errors represent the standard error of the mean determined from at least three independent experiments. *The values for compounds have previously been reported and included here for comparison.³⁴ >100 indicates that the compound was examined but lacked agonist activity at up to 100 μM concentrations. A percentage (%) indicates the percent maximal stimulatory response observed at 100 μM concentrations observed, but not enough stimulation was observed to determine an EC₅₀ value.

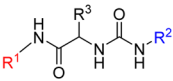
tetrapeptide agonist sequence Ac-His-DPhe-Arg-Trp-NH₂ and were shown to differentiate functional activity between the mMC3R and mMC4R.^{46–48} Compound 1 with Phe substituent at the R³, was equipotent at the mMC1R (within 3-fold experimental error), 6- and 4-fold less potent at the mMC4R and mMC5R, respectively compared to the CGJ-9. Interestingly, replacement of the Trp residue with Phe abolished ability of compound 1 to stimulate the mMC3R up to 100 μM concentrations. Similar observations were reported in other templates, where perturbation of Trp in the postulated pharmacophore region resulted in reduced, or no agonist activity at the MC3R.^{49,50} Substitution of the (*pI*)Phe residue at the R³ position (compound 2) resulted in a full agonist at all four MCRs with 3-, 22-, 5-, and 9- fold decreased potency at the mMC1R, mMC3–5R, respectively as compared to CGJ-9. The (*pCl*)Phe residue substitution at the R³ group (compound 3) abolished agonist activity at all MCRs. The placement of polar and hydrophilic residue His at this position (compound 4) resulted in only 50%, 60%, and 80% of the maximum cAMP response (relative to the control NDP-MSH) at 100 μM concentrations at the mMC1R, mMC4R, and mMC5R, respectively, but showed no activity at the mMC3R. The bulky aromatic group Nal(2') substitution of Trp (compound 5) in compound CGJ-9 resulted in no agonist activity at MCRs.

Compounds 6–10 possess the 4-methylbenzylamine group at the R¹ position and retained the TIQ moiety at the R² position. SAR of ligands 6–10 was selected to examine the effect of an aromatic ring in the place of aliphatic amine chain. The aromatic cyclic ring substitution at the R¹ position was not tolerated well in the template (Table 1). Compound 6 with Phe residue at the R³ position resulted in some partial stimulatory activity at the mMC1R (60%) and mMC5R (70%), but showed

no activity at the mMC3R and mMC4R. When the halogen groups were added to the phenyl ring in compounds 7 and 8, a complete loss of activity was observed at all four receptors tested at 100 μM concentrations. The His residue containing compound 9 resulted in 40% agonist activity at the mMC5R, but no activity at the mMC1R, mMC3R, and mMC4R resulted. The substitution of the bulky aromatic group Nal(2') (compound 10) showed no activity at all four receptors at up to 100 μM concentrations.

Template 2. Compounds 11–21 had modifications made at the R² position, while R¹ was held constant as 4-(aminomethyl)piperidiny (AMP) in the lead compound CGJ-7. Compounds 11–16 contained the bulkier 3,3-diphenylpropyl (DPP) group at the R² position as compared to the benzyl group in CGJ-7, and the R³ position was substituted with Trp, Phe, (*pI*)Phe, (*pCl*)Phe, His, and Nal(2'), respectively. The agonist EC₅₀ values for Template 2 urea derivatives are provided in Table 2. Compound 11, chemically similar to the lead compound CGJ-7 except at the R² position, showed equipotent activity (within experimental error) at the mMC1R, some agonist activity at the mMC3R compared to CGJ-7, but no activity at the mMC4R and mMC5R. Substitution with the Phe residue (12) resulted in equipotent agonist activity at the mMC1R and mMC5R, as compared to the lead compound. Interestingly, compound 12 possessed full agonist activity at the mMC4R (6.7 ± 3.8 μM) and no activity at the mMC3R, which was the opposite pharmacology with respect to CGJ-7. Replacement of the indole ring (compound 11) to the phenyl ring (compound 12) resulted in same potency at the mMC1R, restored activity at the mMC4R and mMC5R, but abolished activity at the mMC3R. All other amino acid substitutions (compounds 13–16) did not show agonist

Table 3. Pharmacology of Template 3 Urea Compounds at the Mouse Melanocortin Receptors^a

Compound				mMC1R	mMC3R	mMC4R	mMC5R
	R ¹	R ²	R ³				
NDP-MSH				0.05 ± 0.01 nM	0.19 ± 0.04 nM	0.06 ± 0.01 nM	0.20 ± 0.05 nM
CGJ-9*	butylamine	TIQ	Trp	7.9 ± 1.0	1.7 ± 1.0	3.4 ± 1.9	4.9 ± 4.8
CGJ-7*	4-AMP	benzyl	Trp	5.2 ± 2.7	4.4 ± 2.8	> 100	8.2 ± 7.9
22	4-AMP	TIQ	Trp	40% @ 100 μM	> 100	30% @ 100 μM	> 100
23	4-AMP	TIQ	Phe	> 100	> 100	> 100	> 100
24	4-AMP	TIQ	(<i>p</i> I)Phe	> 100	> 100	> 100	> 100
25	4-AMP	TIQ	(<i>p</i> Cl)Phe	> 100	> 100	> 100	> 100
26	4-AMP	TIQ	His	6.4 ± 1.1	14 ± 0.4	7.7 ± 1.7	12 ± 1.3
27	4-AMP	TIQ	Nal(2')	> 100	> 100	> 100	> 100

^aThe indicated errors represent the standard error of the mean determined from at least three independent experiments. *The values for compounds have previously been reported and included here for comparison.³⁴ >100 indicates that the compound was examined but lacked agonist activity at up to 100 μM concentrations. A percentage (%) indicates the percent maximal stimulatory response observed at 100 μM concentrations observed, but not enough stimulation was observed to determine an EC₅₀ value.

activity at any of the four MCRs when tested at up to 100 μM concentrations.

Compounds 17–21 contained the Indane group at the R² position were designed to determine whether constraining the aromatic group at this position would change the activity of the template. Compound 17, with a Trp residue at the R³ position, resulted in full micromolar agonist activity at all four receptors. Diminished activity was observed when Phe was substituted at the R³ position (18), with 80% (mMC1R), 60% (mMC4R), and 70% (mMC5R) of the maximum cAMP response at 100 μM concentrations resulted. No stimulatory activity was observed at the mMC3R at 100 μM concentrations. When the Phe ring was substituted by a *p*-iodo substitution (compound 19), some agonist activity was retained at the mouse MC1, MC4, and MC5 receptors while no activity was observed at the mMC3R. The *p*-chloro substitution at the Phe ring, compound 20, lost activity at the mMC3–5Rs and possessed slight agonist activity at the mMC1R. Compound 21 having the His residue at the R³ position, lacked agonist activity at any of the studied receptors.

Template 3. Compounds in Template 3 had fixed R¹ (4-AMP group) and R² (TIQ) positions that were present in lead compounds CGJ-7 and CGJ-9. The R³ position was substituted by the Trp, Phe, (*p*I)Phe, (*p*Cl)Phe, His and Nal(2') residues in compounds 22–27, respectively. Table 3 summarizes the agonist activity for this template at the melanocortin receptors. Compound 26, with a His substitution, resulted in a full μM agonist at all four studied melanocortin receptors and showed equipotency (within 3-fold experimental error) at the mMC1R and mMC5R compared to both CGJ-7 and CGJ-9. Compound 22, with Trp at the R³ position, showed 40% mMC1R and 30% mMC4R agonist activity as compared to the maximum cAMP response at 100 μM concentrations. All other substitutions in compounds 23–25 and 27 resulted in a lack of agonist activity at the MCRs tested.

Central melanocortin signaling has been implicated to regulate feeding behavior and energy homeostasis. There has

been intensive research on identifying small molecules based on peptidic and nonpeptidic template to selectively target centrally located melanocortin receptors. Nonpeptidic small molecules may be favorable as potential drugs because of increased metabolic stability, oral bioavailability and ability to cross blood-brain barrier (BBB). The first small molecule template for melanocortins was identified in a β-turn-based small molecules library screening by Haskell-Luevano and colleagues.²⁷ Following this, a potent and selective small molecule MC4R agonist molecule was reported by Merck & Co.³³ This compound was shown to reduce food intake in rats as well as stimulate erectile activity. Moreover, it was shown that these effects of the compound were mediated by MC4R since MC3R/4R double knockout mice did not show any of the above effects.³³ Several other small molecules have been reported from the Merck Research Laboratories including a report on a piperazine urea based MC4R partial agonist with nM potency at the human MC4R, but without an effect on erectile activity in the rodents.⁴¹ A focused urea-based nonpeptide agonist library has previously been reported by our lab.³⁴ Compound CGJ-7 was identified in that study as selective for the mMC3R (4.4 μM) as compared to the mMC4R, which showed no agonist activity up to 100 μM. Another compound CGJ-9 with TIQ moiety at the R¹ position and aliphatic amine at the R² position resulted in a μM agonist at all four melanocortin receptors. Both of the compounds had Trp at the R³ position and a urea core in the backbone. The urea core presents an advantage due to its similarity to an amide and it can also form hydrogen bonds to form secondary structures. The formation of intramolecular hydrogen bonds between urea and amide moieties in peptides has also been reported.⁵¹ The introduction of an additional heteroatom in the peptidyl backbone by a urea derivative may also influence ligand–receptor interactions.

The present study was undertaken to take advantage of previously published SAR and utilize a three component diversity approach to identify selective MC3R/MC4R receptor

small molecules. In the current study, five compounds (**1**, **2**, **12**, **17**, **26**) resulted in full agonist activity and another five compounds (**4**, **11**, **18**, **19**, **22**) showed a percent of maximal agonist activity either at the mMC3R, mMC4R or both. In general, aliphatic and cyclic amine groups at the R¹ position resulted in active compounds, while aromatic rings were not tolerated well and resulted mostly in inactive compounds. The R² position was modified with constrained (TIQ, Indane) and flexible (3,3-diphenyl propyl) aromatic rings and gave varied results on the melanocortin receptor subtypes. The R³ position was substituted either with nonpolar hydrophobic amino acids Trp, Phe, (*p*I)Phe, (*p*Cl)Phe, Nal(2'), or His residue, which contained a polar imidazole side chain. When the Trp residue at the R³ position in lead compound CGJ-9 was replaced with Phe residue along with the bulkier aromatic group (compound **1**), loss of activity at the mMC3R resulted. This observation support the hypothesis that an indole ring is important for maintaining agonist activity at the mMC3R in this urea-based template and may be harnessed to generate selectivity between centrally expressed MC3 and MC4 receptors. Similar reports of Trp substitution by Ala and Pro in the cyclic peptide MTII resulted in only a 2% cAMP stimulation at the hMC3R.^{52,53} In a shorter tetrapeptide sequence Ac-His-DPhe-Arg-Ala-NH₂, Trp to Ala substitution resulted in total loss of activity.⁴⁷ However, compound **1** retained full agonist activity at the mMC1R, mMC4R, and mMC5R and showed only 5-fold selectivity for the mMC1R and was equipotent at the mMC5R versus the mMC4R. The replacement of Trp with *p*-iodo substituted Phe residue (compound **2**) resulted in full agonist activity at all four MCRs, albeit with decreased potency compared to the lead compound CGJ-9. However, no activity was observed for *p*-chloro substitution (compound **3**) at this position. The difference between (*p*I)Phe and (*p*Cl)Phe pharmacology could be due to the difference in lipophilicity of these two compounds. The lipophilicity of halogens increases as F < Cl < Br < I. This may affect the interactions of these two compounds in the putative hydrophobic binding pocket of melanocortin receptors. Compound **4** with an imidazole ring showed some stimulatory activity at the mMC1R, mMC4R, and mMC5R but not enough to determine an EC₅₀ value. The lack of activity at the mMC3R was similar to the tetrapeptide Trp to His substitution SAR.⁴⁷ In the previous urea-based study, compound CGJ-6 (Figure 2) with 4-methylbenzylamine group at the R¹ position resulted in the most potent compound of that series with high nM activity at the mMC4R.³⁴ Unfortunately, when an aromatic 4-methylbenzyl group was introduced at the R¹ position in Template 1, only compound **6** showed a very slight agonist activity at the mMC1R and mMC5R and compound **9** was only able to stimulate the mMC5R to 40% maximum stimulation. No other compound from the Template 1 series was found active at the MCRs. The steric hindrance caused by the bulky aromatic ring at this position may be the reason for the nonactive compounds in this series. This may also suggest that the mMC1R and mMC5R putative binding pockets may be more flexible to accommodate bulkier side chain than the mMC3R and mMC4R.

Compounds **11**–**21** had a cyclic piperidine ring at the R¹ position, and the R² position was varied to aromatic rings that were attached either to a flexible propyl linker in 3,3-diphenylpropyl or a constrained cyclic frame in Indane. Compound **11** with the R² modified to 3,3-diphenylpropyl and Trp residue at the R³ position showed selectivity for the

mMC1R. When Trp residue was replaced with Phe, resulting in compound **12**, equipotent activity at the mMC1R and mMC5R resulted, as compared to the lead compound CGJ-7, but no activity at the mMC3R was observed. Compound **12** showed 6.7 μM full agonist activity at the mMC4R and was selective for the centrally located mMC4R versus the mMC3R. However, this analog was not selective when compared to the predominantly peripheral receptors mMC1R and mMC5R. When the aromatic ring was constrained by a cyclic frame as opposed to a diphenyl ring that was attached to a flexible propyl linker, compound **17** containing Trp residue showed full agonist activity at all four receptors. Compounds **18** and **19**, containing Phe and (*p*I)Phe, respectively, did not show full agonist activities and contained varied stimulatory activity at the mMC1R, mMC4R, and mMC5R. Both of these compounds were inactive at the mMC3R. Compound **18** showed more activity than the compound **19** at the mMC1R. The differences in SAR for these two groups can be attributed to the relatively smaller size of Indane and the propyl linker of the 3,3-DPP group, which allows increased rotational freedom of the aromatic rings and renders it less rigid as compared to Indane substitution.

Finally, template 3 (compounds **22**–**27**) was designed to see effects of removing the NH proton by placing a ring system directly attached to nitrogen atoms at both ends. The TIQ residue was positioned at the R¹ and 4-AMP at the R² position and were both directly attached to nitrogen atoms. Interestingly, compound **26** with the His substitution at the R³ position resulted in full micromolar agonist at all four melanocortin receptors. The Trp substitution (compound **22**) resulted in only slight agonist activity at the mMC1R and mMC4R with the Trp residue. All other compounds **23**–**25** and **27** having nonpolar hydrophobic substitutions did not show any activity at MCRs. This observation suggests that the hydrogen bond may play an important role in ligand–receptor interactions or stabilizing the ligand in bioactive conformation in this small molecule template.

In summary, we have designed, synthesized, and characterized three novel series of melanocortin agonists based on an unsymmetrical substituted urea core. The synthesis was performed on solid phase using microwave irradiation, resulting in significantly faster syntheses than previously reported. The structure–activity relationship studies based on urea scaffold presented herein provides insight into important ligand–receptor interactions for MC3R vs MC4R activity. In particular, compounds having the aliphatic or saturated cyclic group at the R¹ position, bulky aromatic group at the R² position and indole or phenyl group at the R³ position showed activity at MCRs in the small urea based molecule template. Compounds **1** and **12** showed selectivity for the centrally expressed mMC4R, as compared to the mMC3R, but no significant selectivity profile was observed at the mMC1R and mMC5R versus the mMC4R. These results may be useful in the further design of selective nonpeptide small molecule ligands for the melanocortin receptors.

METHODS

Chemistry. The amino acids *N*α 9-fluorenylmethoxycarbonyl (Fmoc)-Trp-OH, Fmoc-Phe-OH, Fmoc-(*p*Cl)Phe-OH, and Fmoc-Nal(2')-OH, the resin 4-(4-hydroxymethyl-3-methoxyphenoxy)-butyryl-MBHA (HMPB-MBHA Resin, 0.47 mequiv/g substitution), and the coupling reagent *O*-(benzotriazol-1-yl)-*N,N,N',N'*-tetramethyluronium hexafluorophosphate (HBTU) were purchased from

Peptides International (Louisville, KY). Fmoc-(*p*1)Phe-OH was purchased from Synthetech (Albany, OR). Fmoc-His-OH was purchased from Bachem (Torrance, CA). 4-Nitrophenyl chloroformate, 1,4-diaminobutane, 1,2,3,4-tetrahydroisoquinoline, 4-(aminomethyl)piperidine, 3,3-diphenylpropylamine, and 2-aminoindane, diisopropylethylamine (DIEA) were purchased from Sigma-Aldrich (St. Louis, MO). Dichloromethane (DCM), tetrahydrofuran (THF), glacial acetic acid, methanol, acetonitrile, isopropanol (*i*-PrOH), and anhydrous diethyl ether were purchased from Fisher (Fair Lawn, NJ). *N,N*-Dimethylformamide (DMF) was purchased from Burdick and Jackson (McGaw Park, IL). All chemicals were obtained from commercial suppliers and were used without further purification.

General Procedure for Compounds 1–27. The solid-phase synthesis (SPS) of the library compounds used in this study is modified from the previously published method.³⁴ Microwave synthesizer (CEM Discover SPS) was used instead of conventional SPS synthesis. Discover SPS is equipped with a fiber-optic temperature probe for controlling the microwave power delivery. Agitation is accomplished by nitrogen gas bubbling through external setup. A separate vacuum manifold is provided with the assembly for solvent drainage.

Coupling of Diamine Unit (R¹) on Solid Phase. HMPB-MBHA resin (0.1 mmol) was swelled in 25 mL polypropylene reaction vessel (CEM) with THF/DCM (1:1) for 1 h. The solvent was drained, and then *p*-nitrophenyl chloroformate (18.7 equiv in THF/DCM (1:1), 2 mmol) and DIEA (18.7 equiv in THF/DCM (1:1), 2 mmol) were added. The reaction mixture was bubbled with nitrogen for mixing and subjected to the microwave irradiation (35 °C, 30 W) for 5 min. After which, the resins were washed with 2 × THF/DCM (1:1), 2 × DCM, 1 × DMF. The appropriate diamine, R¹ subunit (12.5 eq, 1.2 mmol) was then added, along with DIEA (12.5 equiv) in DMF under microwave irradiation (55 °C, 50W) for 10 min to generate **28**. The presence of the primary amine group was tested using the Kaiser test.³⁴ The resins were washed with 3 × DMF.

Coupling of Amino Acid Unit (R³) on Solid Phase. The diamine addition was followed by coupling of amino acid (3.0 eq, 0.3 mmol) under the microwave irradiation (75 °C, 30 W) for 5 min using HBTU (3.0 equiv) and DIEA (5.1 equiv) to the terminal *N*-Fmoc protected intermediate **29**. The coupling of the His amino acid was performed at lower temperature in the microwave synthesizer (50 °C, 30 W) for 5 min. The coupling of appropriate amino acid was confirmed by a negative Kaiser test. The resins were washed with 5 × DMF and subjected to the Fmoc deprotection by 20% piperidine in DMF 1 × 2 min at room temperature and then by irradiating the sample at 75 °C, 30 W for 4 min in the CEM Discover SPS instrument to generate free amine compound **30** on the resin. The removal of Fmoc was confirmed by a positive Kaiser test.

Coupling of R² Subunit on Solid Phase. The *N*-Fmoc deprotection was followed by activation with *p*-nitrophenyl chloroformate (1.4 equiv) and DIEA (1.4 equiv) in 1:1 THF/DCM for 5 min under same conditions as described in the first step. The resins were washed with 2 × THF/DCM (1:1), 3 × DCM, and 2 × DMF, and respective amine group (R² subunit) (1.4 equiv) was then added, along with DIEA (1.4 equiv) in DMF and irradiate under microwave irradiation (55 °C, 50 W) for 10 min to get **31**. The resins were washed with 3 × DMF and 2 × THF/DCM (1:1), 2 × THF and DCM, 2 × acetic acid and *i*-PrOH, 1 × methanol, and 3 × DCM and dried under vacuum before cleavage.

Cleavage from the Solid Phase. The cleavage of compounds from resin was performed by adding glacial acetic acid (5 mL) to the resin and subjected it to microwave irradiation for 30 min at 38 °C, 20W. Following 30 min, the cleaved product was collected, the resins were washed with an additional 1 mL of glacial acetic acid. The solutions were then frozen at –78 °C and lyophilized to achieve crude compounds 1–27.

The crude compounds were purified by reversed-phase HPLC (flow rate of 5 mL/min, acetonitrile/water 0.1% TFA) using a Shimadzu chromatography system with a photodiode array detector and a semipreparative RP-HPLC C18 bonded silica column (Vydac 218TP1010, 1.0 × 25 cm). The purified compounds were >95%

pure as determined by RP-HPLC in two diverse solvent systems and had the correct molecular mass (Supporting Information). Electrospray ionization mass spectrometry (ESI-MS) was used to record spectra on an ABI 3200Q TRAP instrument at the University of Florida. The lead compounds **1** and **12** were also characterized by NMR and HRMS. NMR spectra was acquired in MeOH-*d*₄ with Varian 400 MHz spectrometer with autosampler and HRMS (ESI positive) were obtained with a Bruker BioToF II mass Spectrometer (University of Minnesota).

Functional Bioassay. HEK-293 cells stably expressing the mouse melanocortin receptors were maintained in Dulbecco's modified Eagle's medium (DMEM) with 10% fetal calf serum and transfected with 4 μg of the CRE/β-galactosidase reporter gene as previously described.⁴⁴ Briefly, 5000–15 000 post-transfection cells were plated into collagen treated 96-well plates (Nunc) and incubated overnight. At 48 h post-transfection, the cells were stimulated with 100 μL of peptide (10^{−4}–10^{−12} M) or forskolin (10^{−4} M) control in assay medium (DMEM containing 0.1 mg/mL BSA and 0.1 mM isobutylmethylxanthine) for 6 h. The assay media was aspirated and 50 μL of lysis buffer (250 mM Tris-HCl pH = 8.0 and 0.1% Triton X-100) was added. The plates were stored at –80 °C overnight. The plates containing the cell lysates were thawed the following day. Aliquots of 10 μL were taken from each well and transferred to another 96-well plate for relative protein determination. To the cell lysate plates, 40 μL of phosphate-buffered saline with 0.5% BSA was added to each well. Subsequently, 150 μL of substrate buffer (60 mM sodium phosphate, 1 mM MgCl₂, 10 mM KCl, 5 mM β-mercaptoethanol, 2 mg/mL ortho-nitrophenyl-β-galactoside [ONPG]) was added to each well and the plates were incubated at 37 °C. The sample absorbance, OD₄₀₅, was measured using a 96-well plate reader (Molecular Devices). The relative protein was determined by adding 200 μL of 1:5 dilution Bio Rad G250 protein dye/water to the 10 μL cell lysate sample taken previously, and the OD₅₉₅ was measured on a 96-well plate reader (Molecular Devices). Data points were normalized both to the relative protein content and nonreceptor dependent forskolin stimulation. Maximal efficacy was compared to that observed for the NDP-MSH control peptide tested simultaneously on each 96-well plate.

Data Analysis. The agonist EC₅₀ values represent the mean of duplicate wells performed in three or more independent experiments. The EC₅₀ value estimates, and their associated standard errors, were determined by fitting the data to a nonlinear least-squares analysis using the PRISM software program (v4.0, GraphPad Inc.). The compounds were assayed as TFA salts.

■ ASSOCIATED CONTENT

📄 Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acscemneuro.5b00273.

Analytical data table of all synthesized compounds, characterization data of compounds **1** and **12**, and NMR spectra (PDF)

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Notes

The authors declare no competing financial interest.

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A.S. and C.H.-L. designed the research. A.S., J.K., M.L.S.D. and H.H performed the experiments. A.S. and C.H.-L. analyzed the data. A.S. wrote the manuscript with the help of C.H.-L.

ABBREVIATIONS

MCR, melanocortin receptor; MC3R, melanocortin-3 receptor; MC4R, melanocortin-4 receptor; POMC, proopiomelanocortin; SAR, structure–activity relationship; TIQ, 1,2,3,4-tetrahydroisoquinoline; AMP, 4-(aminomethyl)piperidine; MBA, (4-methyl)benzylamine; DPP, diphenylpropyl; RP-HPLC, reverse-phase high-pressure liquid chromatography; MBHA, 4-methylbenzhydrylamine

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