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Synthesis and pharmacological evaluation of novel C-8 substituted tetrahydroquinolines as balancedaffinity mu/delta opioid ligands for the treatment of pain

Anthony F Nastase, Nicholas W. Griggs, Jessica P. Anand, Thomas J. Fernandez, Aubrie A Harland, Tyler J Trask, Emily M. Jutkiewicz, John R. Traynor, and Henry I. Mosberg ACS Chem. Neurosci., Just Accepted Manuscript • DOI: 10.1021/acschemneuro.8b00139 • Publication Date (Web): 20 Apr 2018 Downloaded from http://pubs.acs.org on April 22, 2018

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26	^a Department of Medicinal Chemistry, College of Pharmacy, University of Michigan, 428 Church
27	Street, Ann Arbor, MI 48109, United States
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29	^b Interdepartmental Program in Medicinal Chemistry, College of Pharmacy, University of
30	Michigan Ann Arbor MI 48109 United States
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27	^c Department of Pharmacology, Medical School, University of Michigan, Ann Arbor, MI 48109,
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34	^a Edward F Domino Research Center, University of Michigan, Ann Arbor, MI 48109
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Abstract

The use of opioids for the treatment of pain, while largely effective, is limited by detrimental side effects including analgesic tolerance, physical dependence, and euphoria, which may lead to opioid abuse. Studies have shown that compounds with a µ-opioid receptor (MOR) agonist/δ-opioid receptor (DOR) antagonist profile reduce or eliminate some of these side effects including development of tolerance and dependence. Herein we report the synthesis and pharmacological evaluation of a series of tetrahydroquinoline-based peptidomimetics with substitutions at the C-8 position. Relative to our lead peptidomimetic with no C-8 substitution, this series affords an increase in DOR affinity and provides greater balance in MOR and DOR binding affinities. Moreover, compounds with carbonyl moieties at C-8 display the desired MOR agonist/DOR antagonist profile whereas alkyl substitutions elicit modest DOR agonism. Several compounds in this series produce a robust antinociceptive effect *in vivo* and show antinociceptive activity for greater than two hours after intraperitoneal administration in mice.

Keywords: Bifunctional ligands, dependence, drug abuse, opioids, peptidomimetics, tolerance

Introduction

Opioid analgesics are among the most effective drugs for the treatment of moderate to severe pain. Complications from long-term pain management with opioids include gastrointestinal distress, opioid-induced hyperalgesia and the development of analgesic tolerance and physical dependence. Nearly all clinically prescribed opioid analgesics exert both their beneficial and undesirable effects through activation of the μ -opioid receptor (MOR). Evidence suggests that the δ -opioid receptor (DOR) plays a key role in modulating some side effects associated with opioids including analgesic tolerance and physical dependence.^{1,2} Classic studies using rodent models have demonstrated that by co-administering a DOR antagonist with morphine, or by administering chronic morphine to DOR knockout mice, the development of analgesic tolerance, physical dependence and drug-seeking behavior are attenuated.^{3–7} By simultaneously activating MOR and blocking DOR, we aim to develop safer analgesics with reduced tolerance and dependence profiles for the management of a variety of pain conditions.

Current reports and reviews on bifunctional MOR/DOR ligands, which span a wide range of scaffolds and physicochemical properties, have provided support for the development of improved therapeutics that display favorable pharmacodynamic and pharmacokinetic profiles.^{6–12} Eluxadoline (Viberzi), an orally bioavailable, peripherally-acting MOR agonist/DOR antagonist, was FDA approved in 2015 for the treatment of irritable bowel syndrome with diarrhea (IBS-D) and has shown to be effective and well-tolerated in the clinic.^{13–15} Several labs have utilized morphine-like scaffolds to develop MOR agonist/DOR antagonist small molecules, which may show promise as analgesics with reduced antinociceptive tolerance and physical dependence liabilities.^{10,16,17} An alternative approach by Portoghese and colleagues targets the proposed

MOR-DOR heterodimer with a bivalent ligand, whereby the MOR agonist pharmacophore of oxymorphone is linked with the DOR antagonist pharmacophore of naltrindole.^{18,19} Studies using these bivalent ligands in animal models not only show decreased tolerance and dependence as a function of linker length,¹⁸ but also report attenuated conditioned place preference (CPP) and reinstatement, suggesting a lower abuse liability.¹⁹ We have recently reported on two MOR agonist/DOR antagonist lead compounds, a glycosylated cyclic peptide⁷ and a peptidomimetic, **1** (Figure 1),^{20,21} that produce long-lasting and dose-dependent antinociception in mice after peripheral administration. Elaboration of the peptidomimetic series primarily through modification of the C-6 pendant and at N-1 resulted in MOR agonist/DOR antagonist analogues with a range of relative affinities for MOR and DOR.^{22–24} Further *in vivo* evaluation of three such analogs suggested that balanced MOR/DOR affinity is associated with amelioration of tolerance, dependence, and reinforcing properties.²⁵

In order to determine what role, if any, balanced MOR/DOR affinities of MOR agonist/DOR antagonist ligands plays in eliminating opioid side effects, we continue to expand the SAR of this series for the development of MOR agonist/DOR antagonist peptidomimetics with a range of relative MOR/DOR affinities. As noted above, our SAR has focused primarily on substitutions at C-6 and N-1.



Figure 1. Lead peptidomimetic 1, featuring the THQ core and numbering convention in blue. Current SAR exploration focuses on substitutions at C-8.

To further understand and optimize the SAR for our tetrahydroquinoline (THQ) core, we synthesized and evaluated a series of C-8 substitutions while keeping the pharmacophores at C-6

and N-1 constant. We had previously found that ligands with bicyclic C-6 substitutions on the unmodified THQ core preferentially bind MOR over DOR at least 10-fold,²² and N-1 substitutions afford improved balance in MOR and DOR binding affinity, but often elicit partial DOR agonism.²⁴ The C-8 modifications presented in this series probe the spatial and electronic constraints of a binding pocket previously unexplored in this series of peptidomimetics. Here we report the synthesis and evaluation of a series of THQ-based ligands featuring a diverse set of substitutions at C-8 as part of our effort to optimize the MOR agonist/DOR antagonist profile and improve upon our lead peptidomimetic.

Results/Discussion

Following the development of **1**, we have extensively evaluated the SAR of our peptidomimetic series by incorporating diverse substitutions at the C-6 position,^{21,22} also referred to as the "pendant." Prior SAR and computational docking studies have shown that a benzyl pendant at C-6 is well tolerated by a deep binding pocket in both MOR and DOR.^{20,21,23} To validate the importance of the aryl C-6 pharmacophore in receptor binding and activity, the benzyl pendant was moved to the C-8 position (Table 1, Compound 2). Compound 2 showed significantly decreased binding affinities (higher K_i) for all 3 opioid receptors (MOR, DOR and κ -opioid receptor (KOR)), and also showed decreased MOR potency (EC₅₀ = 1200 nM) and efficacy (37% stimulation compared to the standard MOR agonist DAMGO), as measured by [³⁵S]-GTP_γS binding. Furthermore, compound **2** showed no appreciable activity at DOR or KOR. We then questioned whether this reduction in MOR activity was due to the loss of the C-6 pharmacophore, or to unfavorable ligand-receptor interactions at C-8. To examine this, we combined the C-6 and C-8 benzyl substitutions, giving us compound **7a**. As shown in Table 1, the binding affinity as well as potency and efficacy of **7a** at MOR were restored (K_i = 1 nM;

 $EC_{50} = 4$ nM; 96% stimulation), while the DOR binding affinity improved 10-fold and efficacy 2-fold compared to **1**, not only validating the importance of the C-6 pharmacophore for MOR activity, but identifying a key role for the C-8 pharmacophore in modulating DOR affinity and efficacy. This moderate loss in MOR affinity and increase in DOR affinity shifted the MOR/DOR binding ratio (DOR K_i/MOR K_i) from 43 for compound **1** to a more balanced 1.6 for compound **7a** (Table 1). Consequently, this 6-,8-disubstituted THQ analogue established C-8 as a region of interest for future SAR with the principal aim of balancing MOR and DOR binding affinities.

Table 1. Effects of Benzyl Pendant Position on Binding Affinity, Potency and Efficacy



	Binding affinity K _i (nM)			DOR K _i / MOR K _i	Potency EC ₅₀ (nM)			Efficacy (% stimulation)		
Cmpd	MOR	DOR	KOR		MOR	DOR	KOR	MOR	DOR	KOR
1	0.22 (0.02)	9.4 (0.8)	68 (2)	43	1.6 (0.3)	110 (6)	540 (70)	81 (2)	16 (2)	22 (2)
2	48 (9)	360 (60)	1500 (400)	7.5	1200 (300)	dns	dns	37 (4)	dns	dns
7a	1.0 (0.1)	1.6 (0.4)	23 (5)	1.6	4 (2)	380 (84)	dns	96 (4)	42 (7)	dns

Table 1. Binding affinities (K_i) were obtained by competitive displacement of radiolabeled [³H]-diprenorphine in membrane preparations. Functional data were obtained using agonist induced stimulation of [³⁵S]-GTP γ S binding assay. Potency is represented as EC₅₀ (nM) and efficacy as percent maximal stimulation relative to standard agonist DAMGO (MOR), DPDPE (DOR), or U69,593 (KOR) at 10 μ M. All values are expressed as the mean of three separate assays performed in duplicate with standard error of the mean (SEM) in parenthesis. dns = does not stimulate (<10%).

In vitro Structure-Activity Relationships

Subsequent compounds in the C-8 series explored the steric environment and depth of the C-8 binding pocket with various alkyl substitutions, ranging from methyl to *t*-butyl (Table 2). We extended this series to include halogens (F, CF₃, Br), which largely fit the same trend as the alkyl set. The alkyl and halogenated series generally showed potent, efficacious agonism at MOR and partial agonism at DOR. Additionally, most alkyl-substituted analogues showed no KOR activation, whereas the halogenated compounds were partial agonists at KOR. In terms of binding, the smallest C-8 substitutions (**7b**, **7c**, **7g**, **7h**, **7q**) maintained high affinity for MOR and increased affinity for DOR relative to the unsubstituted lead peptidomimetic **1**, however these compounds bind MOR over DOR by at least 8:1. Conversely, larger C-8 substitutions (**7d**, **7e**, **7f**) slightly decreased MOR affinity and displayed a modest increase in DOR affinity, leading to an improved balance of opioid receptor binding affinities.

Û		Binding	∑он g affinity,	K _i (nM)	DOR Ki/ MOR Ki	Potency, EC ₅₀ (nM)		Efficacy (% stimulation)			
Cmpd	R ₂	MOR	DOR	KOR		MOR	DOR	KOR	MOR	DOR	KOR
1	T H	0.22 (0.02)	9.4 (0.8)	68 (2)	43	1.6 (0.3)	110 (6)	540 (70)	81 (2)	16 (2)	22 (2)
7b	Т	0.24 (0.08)	1.9 (0.4)	17 (0.7)	8	4.2 (1.6)	110 (24)	760 (140)	91 (1)	71 (3)	52 (2)
7c	厂	0.09 (0.04)	1.9 (0.4)	40 (5)	21	6.2 (2.9)	32 (10)	dns	74 (2)	45 (4)	dns
7d	7	0.64 (0.08)	5.9 (1.5)	98 (18)	9	23 (7)	310 (30)	dns*	90 (6)	36 (3)	dns*
7e	Ţ	0.76 (0.28)	3.6 (0.5)	34 (5)	5	17 (4)	250 (39)	dns*	85 (2)	25 (4)	dns*
7f	\downarrow	0.47 (0.18)	3.8 (0.7)	48 (7)	8	9.9 (3.6)	240 (40)	dns*	83 (5)	42 (2)	dns*
7g	Ţ	0.11 (0.01)	3.0 (0.3)	9 (1)	27	1.6 (0.2)	97 (19)	370 (8)	95 (2)	28 (3)	40 (1)
7h	CF3	0.26 (0.10)	2.2 (0.7)	29 (10)	9	1.8 (0.9)	50 (14)	580 (14)	70 (5)	42 (2)	18 (4)
7q	Br	0.23	2.4	13 (1)	10	1.2	36 (18)	310 (95)	73 (3)	69 (4)	29 (1)

Table 2. Effects of Alkyl and Halogen Substitutions on Affinity, Potency and Efficacy

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Table 2. Binding affinities (K_i) were obtained by competitive displacement of radiolabeled [³H]-diprenorphine in membrane preparations. Functional data were obtained using agonist induced stimulation of [³⁵S]-GTP γ S binding assay. Potency is represented as EC₅₀ (nM) and efficacy as percent maximal stimulation relative to standard agonist DAMGO (MOR), DPDPE (DOR), or U69,593 (KOR) at 10 μ M. All values are expressed as the mean of three separate assays (n=3) performed in duplicate unless noted otherwise, with SEM in parenthesis. * indicates n=2; dns = does not stimulate (<10% stimulation).

Expanding upon the alkyl and halogen subsets, we synthesized a series of analogues featuring aryl, carbonyl, and amine substitutions, summarized in Table 3. The phenethyl (**7l**) and ethyl ester (**7n**) C-8 substituted compounds both improved the MOR/DOR binding affinity ratio relative to **1** (DOR K_i/MOR K_i = 4.1 and 3.7, respectively), however, they were not as balanced as the benzyl C-8 substituted compound **7a** (DOR K_i/MOR K_i = 1.6). Compounds **7l** and **7n** maintained appreciable MOR efficacy (>70% stimulation), yet only modestly stimulated DOR ($\leq 15\%$ stimulation), whereas **7a** produced partial agonism at DOR (EC₅₀ = 380 nM; 42% stimulation). Evaluation of three amide-linked C-8 substituted compounds (ethyl amide **7m**,

phenyl amide 70, and benzyl amide 7p) resulted in binding affinities that favored MOR over DOR approximately 10:1. Functionally, these compounds produced the desired MOR agonist/DOR antagonist profile. Additionally, this series of carbonyl-containing compounds increased KOR affinity relative to 1 and displayed KOR antagonism. The carboxylic acidsubstituted compound (7r) was the outlier in the carbonyl series, showing a significant loss of KOR binding affinity ($K_i = 210$ nM), however affinity and activity at MOR and DOR were comparable to the other compounds in the carbonyl series. The amine substitutions examined in this series were all cyclic, tertiary amines (e.g. piperidine (7i), morpholine (7i), and piperazine (7k). These C-8 substitutions elicited minor differences at MOR, with affinities and efficacies comparable to 1, however, these compounds diverged markedly at DOR and KOR. Compound 7k displayed the weakest DOR affinity in this series by a considerable margin ($K_i = 15$ nM), while 7i and 7j both displayed high affinity and partial agonism for DOR. Furthermore, while 7i and 7k showed a large increase in KOR binding affinity ($K_i = 0.93$ and 1.9 nM, respectively) and produced partial KOR agonism, 7j showed a more modest increase in KOR affinity ($K_i = 7.3$ nM) and did not stimulate KOR.

\bigcirc		Binding	∑он g affinity,	K _i (nM)	DOR Ki/ MOR Ki	Pote	ncy, EC ₅₀	(nM)	Efficacy	ı (% stim	ulation)
Cmpd	R ₂	MOR	DOR	KOR		MOR	DOR	KOR	MOR	DOR	KOR
7a		1.0 (0.1)	1.6 (0.4)	23 (5)	2	4 (2)	380 (84)	dns	96 (4)	42 (7)	dns
71		0.37 (0.07)	1.4 (0.7)	27 (8)	4	44 (17)	33 (13)	dns*	78 (1)	15 (2)	dns*
7m		0.20 (0.06)	1.8 (0.4)	25 (4)	9	2.1 (0.3)	dns	dns	56 (5)	dns	dns
7n	\sim	1.0 (0.3)	3.7 (0.3)	47 (2)	4	4.9 (0.3)	dns	dns*	71 (3)	dns	dns*
70	©, _₽ ₹₀	0.32 (0.08)	5.4 (0.5)	29 (5)	17	1.2 (0.7)	dns	dns*	49 (4)	dns	dns*
7р		0.17 (0.03)	3.0 (0.4)	30 (2)	18	3.4 (1.4)	dns	dns	73 (5)	dns	dns
7r	но	0.47 (0.16)	2.4 (0.4)	210 (6)	5	4.4 (1.7)	dns	dns	67 (3)	dns	dns
7i		0.07 (0.03)	4.4 (1.0)	0.93 (0.18)	63	2.3 (0.2)	27 (2)	100 (33)	93 (2)	31 (4)	30 (6)
7j		0.15 (0.04)	2.3 (0.7)	7.3 (1.4)	15	1.8 (0.4)	180 (47)	dns	96 (2)	29 (5)	dns
7k		0.35 (0.18)	15 (3)	1.9 (0.5)	43	8.2 (3.5)	290 (100)	170 (67)	60 (2)	18 (1)	17 (1)

Table 3. Effects of Aryl, Carbonyl and Amino Substitutions on Affinity, Potency and Efficacy

Table 3. Binding affinities (K_i) were obtained by competitive displacement of radiolabeled [³H]-diprenorphine in membrane preparations. Functional data were obtained using agonist induced stimulation of [³⁵S]-GTP γ S binding assay. Potency is represented as EC₅₀ (nM) and efficacy as percent maximal stimulation relative to standard agonist DAMGO (MOR), DPDPE (DOR), or U69,593 (KOR) at 10 μ M. All values are expressed as the mean of three separate assays (n=3) performed in duplicate unless noted otherwise, with SEM in parenthesis. * indicates n=2; dns = does not stimulate (<10% stimulation).

In vivo antinociceptive activity. All final compounds excluding 7k and 7o were evaluated *in vivo* for antinociceptive activity using the mouse warm water tail withdrawal (WWTW) assay (Table 4). In the alkyl series, compounds 7b, c and e were fully efficacious, showing dose dependent antinociception and reaching the cutoff latency of 20 seconds at 10 mg/kg after intraperitoneal (ip) administration, whereas 7d showed no significant antinociceptive effect at the same dose. The *t*-butyl analogue 7f was partially active *in vivo*, with a latency of 10 seconds at 10 mg/kg. In the carbonyl series, only the ethyl ester analogue 7n also showed full efficacy.

Larger substitutions including the cyclic amines 7i and j, aryl rings 7a and l, and amides 7m, o and p, produced no antinociception at the doses tested. Additionally, the relatively small carboxylic acid 7r and halogenated analogues 7g, h and q produced no antinociceptive effect at the doses tested. Results of the *in vivo* screening are summarized in Table 4. Of the bioactive analogues 7b, c, e and n, the duration of action for 7e and n proved to be the longest at 2.5 hours. This is a modest improvement over the lead 1 (2 hours).

	C-8 substitution	Antinociceptive activity at 10 mg/kg	Duration of action		C-8 substitution	Antinociceptive activity at 10 mg/kg	Duration of action
7a	benzyl	No activity		7j	methylmorpholine	No activity	
7b	methyl	Fully efficacious	1.5 hr	7k	methylpiperazine	Did not test	
7c	ethyl	Fully efficacious	1.0 hr	71	phenethyl	No activity	
7d	<i>n</i> -propyl	No activity		7m	ethyl amide	No activity	
7e	<i>n</i> -butyl	Fully efficacious	2.5 hrs	7n	ethyl ester	Fully efficacious	2.5 hrs
7f	<i>t</i> -butyl	Partially active		70	phenyl amide	Did not test	
7g	fluoro	No activity		7р	benzyl amide	No activity	
7h	trifluoromethyl	No activity		7q	bromo	No activity	
7i	methylpiperidine	No activity		7r	carboxylic acid	No activity	

Table 4. Antinociceptive Activity of Compounds 7a-r in Mouse WWTW assay

Table 4. Results from the mouse WWTW assay after ip administration of compound **7a-r** at 10 mg/kg. Antinociceptive activities defined as fully efficacious for 20 s latency to tail withdrawal, partially active for 10 s above baseline, or no activity for no significant difference from baseline. Duration of action is defined here as the time it takes to return to baseline after a 10 mg/kg bolus injection of test compound.

All compounds reported in this series maintained a high binding affinity at MOR and demonstrated partial to full MOR agonism *in vitro* compared to the standard full agonist DAMGO. With compound **7k** as the sole exception, all compounds had improved DOR affinity relative to the lead compound **1**. Accordingly, both lipophilic and polar C-8 substitutions provided compounds with a greater balance in MOR/DOR receptor binding affinity. While most compounds displayed DOR agonist activity, those with carbonyl C-8 substitutions (**7m, n, o, p**)

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and **r**) were consistently found to lack efficacy in the GTP γ S binding assay. We have previously shown that peptidomimetics that do not produce GTP γ S binding are in fact functional antagonists, as demonstrated by a shift in the concentration response curve of a standard agonist. Using SNC80 as a standard DOR agonist, we calculated the K_e values of **7m**, **n**, **o**, **p**, and **r** to be 25.4 nM, 42.5 nM, 31.5 nM, 16.7 nM, and 15.7 nM, respectively, confirming their DOR antagonist proerties.

Based on our computational models,²⁶ we predict the C-8 substitutions to primarily interact with extracellular loop 2 and the N-terminal tail of MOR and DOR, and to a lesser extent with transmembrane helices 3, 4 and 5. Due to the flexibility in these regions of the receptor, accurately correlating computationally predicted ligand-receptor interactions with the SAR data was not feasible. Comprehensive evaluation of this series of compounds suggests that C-8 carbonyl moieties block DOR activation quite effectively and bulky alkyl and aryl groups, such as *n*-butyl and phenethyl, respectively, attenuate DOR activation relative to smaller alkyl, aryl, and halogen-containing groups. We have previously shown that a bulky C-6 pendant interacts favorably with the active-state MOR binding pocket, yet there is a steric clash between a large C-6 pendant and the analogous amino acid residues in the active-state DOR.^{21,26} We propose from our SAR analysis that the active-state binding pockets of MOR and DOR likely interact with the C-8 substitutions in a similar manner. For preferentially binding the MOR active-state pocket and the DOR inactive-state pocket with a high affinity, a bulky C-6 pendant and carbonyl moiety at the C-8 position are key pharmacophore elements that produce the desired MOR agonist/DOR antagonist profile with an improved receptor binding affinity balance.

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Moreover, in addition to balancing binding affinty at MOR and DOR and demonstrating that C-8 carbonyl substitutions produce DOR antagonism *in vitro*, this series afforded several compounds with promising *in vivo* antinociceptive activities. We have shown that four compounds (**7b**, **c**, **e**, and **n**) produced a full antinociceptive effect in the WWTW assay in mice after peripheral administration. Contrary to expectations, **7d**, which incorporated the *n*-propyl substitution, showed no effect *in vivo* whereas **7e**, having the *n*-butyl substitution, produced full antinociceptive activity. **7d** has only a single carbon difference from the bioactive **7c** and **7e**, so it is surprising that its *in vivo* effects would be drastically different. Among the carbonyl subset, only the ethyl ester (**7n**) was fully efficacious.

As shown in Table 4, most compounds in this series demonstrated no antinociceptive activity at the doses tested *in vivo*. While we cannot definitively attribute a loss of activity to any individual factor for all compounds in the series, typical physicochemical properties such as high molecular weight, lipophilicity, polar surface area, and hydrogen bond partners are likely to inhibit membrane permeability and access to the CNS. Accordingly, our reported *in vivo* SAR was constrained to small C-8 modifications. Small alkyl substitutions (**7b-f**) showed the best correlation with *in vivo* antinociceptive activity. However, the relatively low molecular weight fluoro-substituted compounds (**7g** and **h**) showed no activity *in vivo* at the doses tested, indicating pharmacokinetic obstacles besides molecular weight. Another pair of relatively low molecular weight analogues with differing *in vivo* effects, **7m** and **n**, suggest new parameters affecting bioavailability. Although **7m** and **n** are comparable in size, **7m** features an hydrogen bond donating amide moiety, whereas **7n** bears a more lipophilic, hydrogen bond accepting ester functionality. The added hydrogen bond donating amide may affect specific interactions with

proteins that impact CNS access (e.g. active transporters, efflux proteins, metabolizing enzymes), or nonspecific parameters, including polarity, and by extension, passive membrane permeability. Unsurprisingly the carboxylic acid-substituted compound (**7r**) showed no *in vivo* activity despite its relatively low molecular weight, likely due to the poor blood-brain barrier permeability of carboxylic acid moieties. Lastly, the relatively large aryl rings (**7a** and **l**), aryl amides (**7o** and **p**), and amine heterocycles (**7i** and **k**) evoked no response *in vivo* at the doses tested, likely due to a combination of unfavorable physicochemical parameters.

In summary, we have shown that modifications at the C-8 position of the THQ scaffold of our peptidomimetic series maintain high MOR affinity while improving DOR affinity, thus achieving our goal of developing more balanced receptor binding affinity compounds. Additionally, we have identified the C-8 carbonyl moiety as a key pharmacophore element for blocking DOR activation, thereby achieving the pharmacologically favorable MOR agonist/DOR antagonist profile. Although it is not known that a 1:1 MOR/DOR binding affinity ratio is optimal, preliminary *in vivo* data for a set of related compounds in our peptidomimetic series show the greatest reduction in tolerance and dependence when MOR/DOR affinity is relatively balanced.²⁵ Confirmation of this trend will require evaluation of tolerance and dependence liabilities of additional analogs with a range of MOR and DOR affinities. The compounds reported to have *in vivo* activity here (**7b, c, e** and **n**) encompass a range of MOR/DOR affinity ratios (7.9, 21, 4.7, and 3.7, respectively) and therefore should provide valuable data supporting or refuting this trend. In conclusion, these bioavailable peptidomimetics serve as promising leads in the search for future MOR agonist/DOR antagonist analgesics.

<u>Methods</u>

Synthesis

All reagents and solvents were obtained commercially and were used without further purification. Intermediates were purified by flash chromatography using a Biotage Isolera One instrument. Most purification methods utilized a hexanes/ethyl acetate solvent system, with a linear gradient between and 100% ethyl acetate. though dichloromethane/methanol/triethylamine solvent systems were utilized for some polar, aminecontaining intermedites. Purification of final compounds was performed using a Waters semipreparative HPLC with a Vydac protein and peptide C18 reverse phase column, using a linear gradient of 0% solvent B (0.1% TFA in acetonitrile) in solvent A (0.1% TFA in water) to 100% solvent B in solvent A at a rate 1% per minute, monitoring UV absorbance at 230 nm. The purity of final compounds was assessed using a Waters Alliance 2690 analytical HPLC instrument with a Vydac protein and peptide C18 reverse phase column. A linear gradient (gradient A) of 0% solvent B in solvent A to 70% solvent B in solvent A in 70 min, measuring UV absorbance at 230 nm was used to determine purity. All final compounds used for testing were >95% pure, as determined by analytical HPLC. ¹H NMR and ¹³C NMR data were obtained on a 500 MHz Varian spectrometer using CDCl₃ or CD₃OD solvents. The identities of final compounds were verified by mass spectrometry using an Agilent 6130 LC-MS mass spectrometer in positive ion mode.

Compounds presented in this series were synthesized as described in Schemes 1 and 2, starting with a commercially available, appropriately substituted aniline. In Scheme 1, anilines **1a-h** were first coupled with 3-bromopropionyl chloride (**i**), yielding intermediates **2a-h**. These



intermediates then underwent (ii) a base-catalyzed β-lactam formation (**3a-h**). This was followed by a triflic acid-catalyzed Fries rearrangement (iii) to produce the THQ core (**4c**, **4g**). In order to achieve the desired aryl bromide substitution pattern for further functionalization, compounds denoted as following Scheme 1a were treated with N-bromosuccinimide to install a bromine at either C-8 (**4a**) or C-6 (**4b-h**). Compounds **4i-k**, shown in Scheme 1b, were derived from **3b**. After cyclization and aromatic bromination, the THQ amine was trifluoroacetyl-protected, then the methyl group at C-8 underwent benzylic bromination with N-bromosuccinimide and benzoyl

peroxide. The benzylic bromide was then substituted under basic conditions with potassium carbonate and the specified amine (along with partial loss of the trifluoroacetyl protecting group), leaving the C-8 substituted aryl bromides **4i-k**. Complete TFA loss observed in step **vi**.





 To convert intermediates 4 to 5, the aryl bromide was functionalized via either Suzuki coupling (step vi, Scheme 1) to give 5a-l, or carbonylation (step vii, Scheme 1c) to give 5n and 4a' followed by amide coupling (step viii, Scheme 1d) for 5m, o, and p.



In Scheme 2, intermediates **5a-p** (and **4a**) were carried forward through a reductive amination step that utilized Ti(OEt)₄ and a chiral Ellman sulfinamide, followed by NaBH₄ to yield the desired *R* stereochemistry at C-4. During this step, the methyl ester of **5n** converted to an ethyl ester, likely due to nucleophilic attack by ethoxide ions liberated from the titanium complex. Addition of concentrated HCl (**x**) cleaved the sulfinamide, leaving a primary amine HCl salt. This amine then underwent amide coupling with *N-,O-*diBoc 2',6'-dimethyl-L-tyrosine (diBocDmt), followed by Boc deprotection with trifluoroacetic acid, yielding final compounds **7a-q**. Compound **7r** was produced by hydrolysis (**xi**) of the ester from **6n**, prior to Boc deprotection. All final compounds were purified by semi-preparative reverse-phase HPLC.

In Vitro Pharmacology

Cell Lines and Membrane Preparations. All tissue culture reagents were purchased from Gibco Life Sciences (Grand Island, NY, U.S.). C6-rat glioma cells stably transfected with a rat MOR (C6-MOR) or rat DOR (C6-DOR) and Chinese hamster ovary (CHO) cells stably expressing a human KOR (CHO-KOR) were used for all *in vitro* assays. Cells were grown to confluence at 37 °C in 5% CO₂ in Dulbecco's modified Eagle medium (DMEM) containing 10% fetal bovine serum and 5% penicillin/streptomycin. Membranes were prepared by washing confluent cells three times with ice cold phosphate buffered saline (0.9% NaCl, 0.61 mM Na₂HPO₄, 0.38 mM KH₂PO₄, pH 7.4). Cells were detached from the plates by incubation in warm harvesting buffer (20 mM HEPES, 150 mM NaCl, 0.68 mM EDTA, pH 7.4) and pelleted by centrifugation at 1600 rpm for 3 min. The cell pellet was suspended in ice-cold 50 mM Tris-HCl buffer, pH 7.4, and homogenized with a Tissue Tearor (Biospec Products, Inc., Bartlesville, OK, U.S.) for 20 s. The homogenate was centrifuged at 15,000 rpm for 20 min at 4°C. The pellet

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was rehomogenized in 50 mM Tris-HCl with a Tissue Tearor for 10 s, followed by recentrifugation. The final pellet was resuspended in 50 mM Tris-HCl and frozen in aliquots at 80°C. Protein concentration was determined via a BCA protein assay (Thermo Scientific Pierce, Waltham, MA, U.S.) using bovine serum albumin as the standard.

Radioligand Competition Binding Assays. Radiolabeled compounds were purchased from Perkin-Elmer (Waltham, MA, U.S.). Opioid ligand binding assays were performed by competitive displacement of 0.2 nM [³H]-diprenorphine (250 μ Ci, 1.85 TBq/mmol) by the peptidomimetic from membrane preparations containing opioid receptors as described above. The assay mixture, containing membranes (20 µg protein/tube) in 50 mM Tris-HCl buffer (pH 7.4), [³H]-diprenorphine, and various concentrations of test peptidomimetic, was incubated at room temperature on a shaker for 1 h to allow binding to reach equilibrium. For several peptidomimetics, radioligand competition binding assays were performed in high Na+ buffer (50 mM Tris-HCl, 100 mM NaCl, 5 mM MgCl₂, 1 mM EDTA, pH 7.4) and samples were incubated at room temperature for 75 min to allow binding to reach equilibrium. Samples were rapidly filtered through Whatman GF/C filters using a Brandel harvester (Brandel, Gaithersburg, MD, U.S.) and washed five times with 50 mM Tris-HCl buffer. Bound radioactivity on dried filters was determined by liquid scintillation counting, after saturation with EcoLume liquid scintillation cocktail, in a Wallac 1450 MicroBeta (Perkin-Elmer, Waltham, MA, U.S.). Nonspecific binding was determined using 10 μ M naloxone. The results presented are the mean \pm standard error (S.E.M.) from at least three separate assays performed in duplicate. K_i (nM) values were calculated using nonlinear regression analysis to fit a logistic equation to the competition data using GraphPad Prism, version 6.0c, for Mac OS X (GraphPad Software Inc., La Jolla, CA).

[³⁵S]-GTPγS Binding Assays. Agonist stimulation of [³⁵S]guanosine 5'-O-[γ- thio]triphosphate [³⁵S]-GTPγS, 1250 Ci, 46.2 TBq/mmol) binding to G protein was measured as described previously.²⁷ Briefly, membranes (10–20 µg of protein/tube) were incubated for 1 h at 25°C in GTPγS buffer (50 mM Tris-HCl, 100 mM NaCl, 5 mM MgCl₂, pH 7.4) containing 0.1 nM [³⁵S]-GTPγS, 30 µM guanosine diphosphate (GDP), and varying concentrations of test peptidomimetic. G protein activation following receptor activation with peptidomimetic was compared with 10 µM of the standard compounds [D-Ala2,N-MePhe4,Gly-ol]enkephalin (DAMGO) at MOR, D-Pen2,5- enkephalin (DPDPE) at DOR, or U69,593 at KOR. The reaction was terminated by vacuum filtration of GF/C filters that were washed 10 times with GTPγS buffer. Bound radioactivity was measured as previously described. The results are presented as the mean ± standard error (S.E.M.) from at least three separate assays performed in duplicate; potency (EC₅₀ (nM)) and percent stimulation were determined using nonlinear regression analysis with GraphPad Prism, as above.

 K_e Determination. Agonist stimulation of [³⁵S]-GTP γ S binding by the known standard agonist SNC80 at DOR was measured as described above. This was then compared to [³⁵S]-GTP γ S binding stimulated by SNC80 in the presence of test compound (1000 nM). Both conditions produced 100% stimulation relative to SNC80. The difference between the EC₅₀ of SNC80 alone and in the presence of test antagonist is the shift in concentration response. The K_e was then calculated as K_e = (concentration of compound)/ (concentration response shift – 1). The results presented are the mean from at three separate assays performed in duplicate.

In Vivo Characterization of Compounds

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Drug preparation. All compounds were administered by intraperitoneal (ip) injection in a volume of 10 mL/kg of body weight. Test compounds were dissolved in 5% DMSO (v/v) in sterile saline (0.9% NaCl w/v).

Animals. Male C57BL/6 wild type mice (Stock number 000664, Jackson Laboratory, Sacramento CA, USA) bred in-house from breeding pairs and weighing between 20-30 g at 8-16 weeks old, were used for behavioral experiments. Mice were group-housed with free access to food and water at all times. Experiments were conducted in the housing room, maintained on a 12 h light/dark cycle with lights on at 7:00 am; all experiments were conducted during the light cycle. Studies were performed in accordance with the University of Michigan Committee on the Use and Care of Animals and the Guide for the Care and Use of Laboratory Animals (National Research Council, 2011 publication).

Antinociception. Antinociceptive effects were evaluated in the mouse WWTW assay. Withdrawal latencies were determined by briefly placing a mouse into a cylindrical plastic restrainer and immersing 2-3 cm of the tail tip into a water bath maintained at 50°C. The latency to tail withdrawal or rapidly flicking the tail back and forth was recorded with a maximum cutoff time of 20 sec (50°C) to prevent tissue damage. Antinociceptive effects were determined using a cumulative dosing procedure. Each mouse received an injection of saline *ip* and then 30 min later baseline withdrawal latencies were recorded. Following baseline determinations, cumulative doses of the test compound (1, 3.2, and 10 mg/kg) were given *ip* at 30 min intervals. Thirty min after each injection, the tail withdrawal latency was measured as described above. To determine the duration of antinociceptive action, baseline latencies were determined as described above.

Thirty minutes after baseline determination, animals were given a 10 mg/kg bolus injection of test compound *ip*. Latency to tail withdrawal was then determined at 5, 15, and 30 min after injections, and every 30 min thereafter until latencies returned to baseline values.

Supporting information

Extended synthetic schemes

Molecular formula strings (CSV)

Abbreviations

MOR, μ-opioid receptor; DOR, δ-opioid receptor; KOR, κ-opioid receptor; DAMGO, [D-Ala², *N*-MePhe⁴, Gly-ol]-enkephalin; DPDPE, [D-Pen²,D-Pen⁵] enkephalin; THQ, tetrahydroquinoline; WWTW, warm water tail withdrawal; di-Boc-Dmt, N, O-Boc 2',6'dimethyl-L-tyrosine; DIPEA, N, N-diisopropylethylamine; PyBOP, benzotriazol-1yloxytripyrrolidinophosphonium hexafluorophosphate; 6-Cl HOBt, 1-hydroxy-6-chlorobenzotriazole; CPP, conditioned place preference; MIDA, N-methyliminodiacetic acid.

Corresponding Author Information

Henry Mosberg

College of Pharmacy, University of Michigan

428 Churc St., Ann Arbor, MI 48109-1065

Phone: 734-764-8117

Fax: 734-763-5595

Email: him@umich.edu

Author Contributions

A.F.N. conceived of, synthesized and performed chemical characterization of 17 of 18 novel final compounds and their related intermediates. A.A.H. synthesized and characterized compound **7g** and its preceding intermediates. A.F.N. and A.A.H. worked

under the direction of H.I.M. *In vitro* assays were performed by N.W.G. with assistance from T.J.F. and T.J.T. under the direction of J.R.T. Animal assays were performed by J.P.A. under the direction of E.M.J.

A.F.N. wrote the manuscript with support from N.W.G. and H.I.M. and feedback from J.P.A., T.J.F., E.M.J., and J.R.T.

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Conflict of Interest

The authors declare no competing financial interest.

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