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Syntheses and opioid receptor binding properties of carboxamido-substituted opioids $\overset{\scriptscriptstyle \times}{}$

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In 2001 we reported our observation that the prototypic phenolic hydroxyl group of certain opioids can be replaced by a carboxamide group and retain high affinity binding to opioid receptors.¹ When this $OH \rightarrow CONH_2$ switch was applied to the cyclazocine core structure, for example, binding affinities for cyclazocine $(1a)^2$ and 8-carboxamidocyclazocine (8-CAC, 1b) were comparable for μ , δ and κ receptors (i.e., K_i values within 2-fold – see first entry in Table 1). A similar result was seen for the $OH \rightarrow CONH_2$ switch in other 2,6-methano-3-benzazocine (a.k.a. benzomorphan) core structures (e.g., ethylketocyclazocine 2a).¹ However, when the core structure was the pentacyclic $4,5\alpha$ -epoxymorphinan derived from natural products, a divergence in SAR was seen. For the morphine (3a/3b) and naltrexone (4a/4b) pairs, binding affinity for $\mu,$ for example, was reduced by 39- and 7-fold, respectively, when the $OH \rightarrow CONH_2$ switch was applied.^{3,4} We recently reported strong evidence that in the naltrexone case, this divergent SAR is likely a consequence of the furan O stabilizing, via a strong intramolecular H-bond, a carboxamide conformation (4c) that is not the bioactive one.⁴ Therefore, 3-desoxy-3-carboxamido naltrexone **4b** and, by analogy, the morphine analogue **3b**, must pay an energy penalty to adopt the putative bioactive conformation depicted in 4d. For the 2,6-methano-3-benzazocines 8-CAC (1b) and 2b, the putative carboxamide bioactive conformation can easily be attained since

ABSTRACT

A series of 15 novel opioid derivatives were made where the prototypic phenolic-OH group of traditional opioids was replaced by a carboxamido (CONH₂) group. For 2,6-methano-3-benzazocines and morphinans similar or, in a few instances, enhanced affinity for μ , δ and κ opioid receptors was observed when the OH \rightarrow CONH₂ switch was applied. For 4,5 α -epoxymorphinans, binding affinities for the corresponding carboxamide derivatives were much lower than the OH partner consistent with our pharmacophore hypothesis concerning carboxamide bioactive conformation. The active metabolite of tramadol and its carboxamide counterpart had comparable affinities for the three receptors.

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there is no barrier created by H-bonding to a neighboring ether bridge.





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Table 1

Opioid binding data for carboxamido-substituted 2,6-methano-3-benzazocine, morphinan, 4,5α-epoxymorphinan and tramadol derivatives



Compound	$K_i - nM \pm SE^a [KBR]^b$							κ:δ ^d
	[3H]DAMGO	[µ]	[³ H]Naltrindole	[δ]	[³ H]U69,593	[κ]		
Cyclazocine core 1a:X = OH^e 1b: X = $CONH_2^f$	0.16 ± 0.01 0.31 ± 0.03	[0.5]	2.0 ± 0.22 5.2 ± 0.36	[0.4]	0.07 ± 0.01 0.06 ± 0.001	[1.2]	2 5	30 90
EKC core 2a : $X = OH^{g,h}$ 2b : $X = CONH_2^{g,h}$	0.78 ± 0.10 1.2 ± 0.12	[0.7]	3.4 ± 0.41 9.8 ± 0.50	[0.4]	0.62 ± 0.11 0.70 ± 0.08	[0.9]	1 2	4 14
Morphine core 3a : X = OH ^{g,i} 3b : X = CONH2 ^{g,i}	0.88 ± 0.14 34 ± 1.8	[0.03]	140 ± 18 1900 ± 81	[0.07]	24 ± 2.3 2000 ± 97	[0.01]	0.037 0.017	6 1
Naltrexone core 4a : X = OH ^j 4b : X = CONH ₂ ^j	0.11 ± 0.006 0.71 ± 0.058	[0.15]	60 ± 3.2 550 ± 40	[0.11]	0.19 ± 0.005 0.36	[0.02]	0.6 0.065	3320 50
Pentazocine core 5a: X = OH 5b: X = $CONH_2^k$	6.9 ± 0.64 5.2 ± 0.41	[1.3]	180 ± 3.2 220 ± 10	[0.8]	3.0 ± 0.18 5.6 ± 0.26	[0.5]	2 1	60 39
Metazocine core 6a: X = OH 6b: X = CONH2 ^k	3.8 ± 0.66 1.6 ± 0.07	[2.4]	140 ± 17 320 ± 43	[0.4]	9.9 ± 0.22 19 ± 0.54	[0.5]	0.4 0.08	14 17
Phenazocine core 7a : X = OH 7b : X = CONH ₂ ^k	0.20 ± 0.07 0.015 ± 0.0010	[13]	5.0 ± 0.88 1.2 ± 0.27	[4.2]	2.0 ± 0.13 0.55 ± 0.029	[3.6]	0.1 0.03	2.5 2
Mr2034 (2'S) core 8a: X = OH ^g 8b: X = CONH2 ^{g,k}	0.19 ± 0.01 0.052 ± 0.013	[3.7]	3.6 ± 0.40 2.0 ± 0.15	[1.8]	0.09 ± 0.01 0.089 ± 0.004	[1.0]	2 0.6	40 22

Table 1 (continued)

Compound	$K_i - nM \pm SE^a [KBR]^b$						κ:μ ^c	κ:δ ^d
	[3H]DAMGO	[μ]	[³ H]Naltrindole	[δ]	[³ H]U69,593	[κ]		
Mr2034 diastereomer (2'R) cor 9a: X = OH ^g 9b: X = CONH ₂ ^{g,k}	e 4.0 ± 0.54 2.9 ± 0.17	[1.4]	67 ± 4.3 34 ± 0.10	[2.0]	1.5 ± 0.07 2.8 ± 0.24	[0.5]	3 1	45 12
Ketocyclazocine core 10a: X = OH ^g 10b: X = CONH2 ^{g,k}	3.3 ± 0.66 1.4 ± 0.07	[2.4]	20 ± 2.7 20 ± 2.3	[1.0]	1.0 ± 0.24 1.8 ± 0.10	[0.6]	3 0.8	20 11
Butorphanol core 11a: X = OH 11b: X = CONH2 ^k	0.12 ± 0.058 0.15 ± 0.019	[0.8]	12 ± 3.8 14 ± 2.1	[0.9]	0.22 ± 0.023 0.39 ± 0.057	[0.6]	0.55 0.38	50 40
Naloxone core 12a: X = OH 12b: X = CONH2 ^k	0.66 ± 0.11 4.5 ± 0.18	[0.1]	120 ± 13 1000 ± 59	[0.1]	1.2 ± 0.072 46 ± 1.3	[0.03]	0.55 0.1	100 22
Naltrexone-6-β-ol core 13a: X = OH 13b: X = CONH2 ^k	0.19 ± 0.016 2.5 ± 0.95	[0.1]	53 ± 3.5 >10 μM	[<0.01]	0.48 ± 0.010 30 ± 0.86	[0.02]	0.4 0.08	110 —
Naltrexone-6-α-ol core 14a : X = OH 14b : X = CONH2 ^k	0.21 ± 0.09 7.1 ± 0.88	[0.03]	56±5.1 >10 μM	[<0.01]	0.56 ± 0.11 100 ± 2.7	[0.01]	0.4 0.07	100 —
Nalmefene core 15a: X = OH 15b: X = CONH2 ^k	0.44 ± 0.083 2.6 ± 0.33	[0.2]	9.3 ± 1.5 67 ± 2.6	[0.1]	0.12 ± 0.0042 2.1 ± 0.29	[0.1]	4 1	78 32
Nalorphine core 16a: X = OH 16b: X = CONH2 ^k	0.19 ± 0.01 18 ± 2.2	[0.01]	120 ± 17 >10 μM	[<.01]	0.38 ± 0.05 100 ± 7.8	[0.004]	0.5 0.2	320 —
Buprenorphine core 17a: X = OH 17b: X = CONH2 ^k	0.21 ± 0.024 0.77 ± 0.065	[0.3]	2.9 ± 0.49 54 ± 8.4	[0.1]	0.62 ± 0.073 2.1 ± 0.12	[0.3]	0.3 0.4	5 26
Nor-BNI core 18a: X = OH 18b: X = CONH ₂ ^k	2.7 ± 0.33 400 ± 32	[0.01]	24 ± 1.6 1000 ± 241	[0.02]	0.027 ± 0.008 21 ± 4.7	[0.001]	100 19	900 50
Tramadol core 19a : X = OH 19b : X = $CONH_2^k$ 19c : X = OCH_3 (tramadol)	8.6 ± 0.37 9.5 ± 0.25 1600 ± 241	[0.9]	2900 ± 66 2400 ± 200 9.4 ± 0.79	[1.2]	450 ± 43 350 ± 10 14 ± 0.44	[1.3]	0.02 0.03 114	6 7 0.7

^a Binding assays used to screen compounds are similar to those previously reported (see Refs. 26 and 27). Membrane protein from CHO cells that stably expressed one type of the human opioid receptor or (as indicated) from guinea pig brain were incubated with 12 different concentrations of the compound in the presence of either 1 nM [³H]U69,593 (µ), 0.25 nM [³H]DAMGO (δ) or 0.2 nM [³H]naltrindole (κ) in a final volume of 1 mL of 50 mM Tris–HCl, pH 7.5, at 25 °C. Incubation times of 60 min were used for [³H]U69,593 and [³H]DAMGO. Because of a slower association of [³H]naltrindole with the receptor, a 3 h incubation was used with this radioligand. Samples incubated with [³H]altrindole also contained 10 mM MgCl₂ and 0.5 mM phenylmethylsulfonyl fluoride. Nonspecific binding was measured by inclusion of 10 µM naloxone. The binding was terminated by filtering the samples through Schleicher and Schuell No. 32 glass fiber filters using a Brandel 48-well cell harvester. The filters were subsequently washed three times with 3 mL of cold 50 mM Tris–HCl, pH 7.5, and were counted in 2 mL Ecoscint A scintillation fluid. For [³H]naltrindole and [³H]U69,593 binding, the filters were soaked in 0.1% polyethyleneimine for at least 60 min before use. IC₅₀ values will be calculated by least squares fit to a logarithm-probit analysis. K_i values of unlabeled compounds were calculated from the equation K_i = (IC₅₀)/1 + *S* where *S* = (concentration of radioligand)/(K_d of radioligand) – see Ref. 28 The K_d values for [³H]DAMGO, [³H]U69,593, and [³H]DAMGO,

^b *K*BR (K_i binding ratio) = K_i (OH)/ K_i (CONH₂) for μ , δ or κ opioid receptors.

^c κ: $\mu = K_i(\mu)/K_i(\kappa)$.

^d $\kappa: \delta = K_i(\delta)/K_i(\kappa).$

^e See text for references to known phenolic-OH opioids.

^f See Ref. 29.

^g Guinea pig membranes.

^h See Ref. 1.

ⁱ See Ref. 3.

^j See Ref. 4.

^k Proton NMR, IR and MS were consistent with the assigned structures of all new compounds. C, H, and N elemental analyses were obtained for all new targets and most intermediates and were within ±0.4% of theoretical values.

Subsequent to these findings, other researchers have published studies where the OH \rightarrow CONH₂ switch was applied to a number of other well-known opioid core structures; these include the morphinan class (e.g., levorphanol, cyclorphan, MCL-101),⁵ phenylpiperidine class^{6,7} and opioid peptides.^{8,9} We now wish to report additional examples of carboxamido-substituted opioids. The objective of this study was to take 15 well-known phenolic-OH-containing opioids (structures shown in Table 1) having divergence in core structures to use as substrates for the OH \rightarrow CONH₂ switch and compare affinity and selectivity for μ , δ and κ opioid receptors

within each pair to determine if the SAR is consistent with that seen with the limited number of pairs previously reported. Intrinsic opioid-receptor mediated activity for a number of high affinity carboxamide-containing ligands compared to the corresponding phenolic-OH prototype was also determined using [^{35}S]GTP γ S binding assays.

Carboxamide targets **5b–11b**, **13b**, **14b**, **16b**, **17b** and **19b** were made directly from their corresponding known phenols using methodology previously reported (Scheme 1).^{1,3,4} Triflate ester formation was accomplished by treating the phenol with (CF₃SO₂)₂O



Scheme 1. Reagents and conditions: (i) (CF₃SO₂)₂O, pyr, CH₂Cl₂, 25 °C (Method A) or PhN(Tf)₂, Et₃N, CH₂Cl₂, 25 °C (Method B); (ii) Zn(CN)₂, Pd(PPh₃)₄, DMF, 120 °C, 20 h or under microwave radiation 150 °C for 15 min; (iii) *t*-BuOH, KOH, 82 °C (combined ii + iii = Method C); (iv) CO/NH₃/Pd(OAc)₂/DPPF in DMSO (Method D) or CO, HN(SiMe₃)₂, PdCl₂, PPh₃, DMF followed by 2 N H₂SO₄ (Method E).



Scheme 2. Reagents and conditions: (i) Ph_3PCH_3Br , $NaNH_2$, THF, 25 °C, 20 h; (ii) K_2CO_3 , H_2O_2 , DMSO, 25 °C, 3 h.

and pyridine in CH_2Cl_2 (Method A) or PhN(SO₂CF₃)₂ and triethylamine in CH_2Cl_2 (Method B). For targets **6b–11b**, **13b**, **14b** and **16b**, preparation of the corresponding carboxamide was enabled via a two-step procedure where the triflate was first converted to a nitrile using $Zn(CN)_2$, $Pd(PPh_3)_4$, in DMF followed by partial hydrolysis of the nitrile using KOH/*t*-BuOH (Method C). Alternatively, triflate esters were directly converted to the carboxamide targets **5b** and **17b** via palladium catalyzed procedures using CO/ NH₃/Pd(OAc)₂/DPPF in DMSO (Method D) or CO/HN(SiMe₃)₂/ PdCl₂/PPh₃ in DMF followed by workup with dilute sulfuric acid (Method E).^{1,3}

Using the methodology described in Scheme 1, the following known phenolic-OH-containing opioids were converted to the corresponding novel carboxamide targets as follows: pentazocine (**5a**)¹⁰ \rightarrow **5b**, Methods A (93%) and D (18%); metazocine (**6a**)¹¹ \rightarrow **6b**, Methods A (75%) and C (26% combined); phenazocine (**7a**)¹² \rightarrow **7b**, Methods B (96%) and C (54%); Mr2034 (**8a**)¹³ \rightarrow **8b**, Methods A (95%) and C (98%); Mr2034 diastereomer (**9a**)¹³ \rightarrow **9b**, Methods A (96%) and C (88%); ketocyclazocine (**10a**)¹⁴ \rightarrow **10b**, Methods A (77%) and C (88%); hutorphanol (**11a**)¹⁵ \rightarrow **11b**, Methods A (77%) and C (89%); naltrexone-6- β -ol (**13a**)¹⁶ \rightarrow **13b**, Methods B (99%) and C (65%); naltrexone-6- α -ol (**14a**)¹⁷ \rightarrow **14b**, Methods B (98%) and C (86%); nalorphine (**16a**)¹⁰ \rightarrow **16b**, Methods B (97%) and C (11%); buprenorphine (**17a**)¹⁸ \rightarrow **17b**, Methods B (95%) and E (48%); and tramadol active metabolite (**19a**)¹⁹ \rightarrow **19b**, Methods A (91%) and C (62%).

Carboxamide target **12b** corresponding to naloxone (**12a**)²⁰ was prepared from the known 6-ethylene ketal derivative²¹ of naloxone. The OH \rightarrow CONH₂ conversion was accomplished using Methods B (74%) and E (69%); deketalization using 6 N HCl in refluxing acetone for 4 h (87%) gave **12b**. As shown in Scheme 2, carboxamide target **15b** corresponding to nalmefene (**15a**)²² was prepared from the known 3-desoxy-3-cyanonaltrexone **21**²³ by first using standard Wittig olefination conditions to provide intermediate **22** in 78% yield. Conversion of **22** to target **15b** was accomplished using K₂CO₃, H₂O₂ in DMSO in 78% yield. Lastly, the carboxamide analogue **18b** of nor-BNI (**18a**)²⁴ was made (Scheme 3) in 51% yield by treating **4b** with hydrazine hydrochloride using conditions very similar to those used to make **18a**.²⁴

Target compounds were evaluated for their affinity and selectivity for μ , δ and κ opioid receptors. Membrane protein from CHO cells that stably expressed one type of the human opioid receptor was used.²⁶ In three instances where indicated, membrane protein from guinea pig brain was used. Opioid receptors from two species were used due to a change in our primary assay midway through this study, however, where data are available. absolute and relative affinities, using human or guinea pig receptors were quite similar. Binding data for all new carboxamide targets compared to their phenolic-OH counterparts are detailed in Table 1. Also included in Table 1 are columns that summarize the K_i Binding Ratio ('KBR') for each core structure against the three receptors [*K*BR = K_i (OH)/ K_i (CONH₂)]. A *K*BR that is ≥ 0.5 and ≤ 2 means the K_i values for the pair of compounds are within 2-fold and have comparable binding affinity for that receptor. A KBR of >2 indicates the carboxamide has higher affinity for the receptor than does its OH counterpart; if it is <0.5, the OH partner has higher affinity.

For 2,6-methano-3-benzazocine core structures (**5–10**), high affinity for μ and κ receptors is seen in all CONH₂ targets; lower affinity for δ is observed. The *K*BRs for μ range from 1.3 (for **5**) to 13 (for **7**) indicating the carboxamide partners have, in several instances, much higher potency than the OH counterpart. For δ and κ receptors, within each pair, both partners have comparable affinity (*K*BRs \geq 0.5 and \leq 2) except for the phenazocine core (**7**) where the carboxamide has considerably higher affinity (*K*BR = 4.2 and 3.6, respectively). For the butorphanol pair **11**, an example of the morphinan class, a very similar trend in *K*BRs (i.e., near unity) and



Scheme 3. Reagents and conditions: (i) H₂NNH₂·HCl, DMF, 150 °C, 1 h.

receptor selectivity (i.e., higher affinity for μ and κ receptors than for δ) was seen compared to published data for the morphinans levorphanol, cyclorphan and MCL-101.⁵ Comparing receptor selectivities of carboxamides **5b–11b** to their phenolic-OH counterparts **5a–11a**, the κ : μ and κ : δ selectivity ratios are similar within each pair with two exceptions. The exceptions are the metazocine (**6**) and phenazocine (**7**) examples where a significant divergence in the κ : μ selectivity is seen; the κ : μ ratios of 0.08 and 0.03, respectively, are much lower (i.e., the carboxamide has much lower affinity for κ than μ) than seen in the other core structures.

As predicted from our previous results with morphine and naltrexone,^{3,4} there is an overwhelming trend in the seven 4,5 α -epoxymorphinan pairs (**12a/b–18a/b**) studied that the carboxamide partner has lower receptor affinity than the corresponding OHcontaining opioid. *K*BRs versus all three receptors were <0.5. For μ , *K*BRs ranged from 0.3 for the buprenorphine core **17a/b** to 0.01 for the nalorphine and nor-BNI cores, **16a/b** and **18a/b**, respectively. Against the δ receptor, binding affinity for the carboxamide partner was also very low compared to the traditional OH partner and *K*BRs ranged from 0.1 (**12a/b**, **15a/b** and **17a/b**) to <0.01 for **13a/b** and **14a/b**. Carboxamide targets **12b–18b** displayed relatively low affinity for κ and *K*BRs of all pairs (**12a/b–18a/b**) for κ were low and ranged from 0.3 (buprenorphine core **17a/b**) to as low as 0.001 (nor-BNI core **18a/b**); more than half of these values were 0.03 or lower. For the seven known phenolic-OH $4,5\alpha$ -epoxymorphinans **12a–18a** studied, six have comparable affinity for μ and κ and much less affinity for δ : these mixed μ/κ compounds are 12a-17a. Comparing this receptor selectivity pattern to that of corresponding carboxamides 12b-17b, little correlation is observed since κ affinity is very low for **12b–17b** compared to affinity for μ ; the $\kappa:\mu$ [$K_i(\mu)/K_i(\kappa)$] ratios ranged from 1 for nalmefene analogue **15b** to 0.07 for the naltrexone-6-β-ol analogue **14b**. Receptor selectivity for **18b** showed a similar trend to **18a**, the well-known κ selective ligand nor-BNI,²⁴ in that both were κ -selective. However, the carboxamide partner displayed very low affinity for all receptors, including κ . The carboxamide partner **19b** of the active metabolite **19a**¹⁹ of tramadol **19c** had comparable binding affinities (KBRs = 0.9-1.3) and a similar receptor selectivity pattern to **19a** for all three receptors. Following characterization of these compounds in our laboratories, an independent study describing their syntheses and biological properties appeared.²⁵

Intrinsic opioid-receptor mediated activity for a number of high affinity carboxamide-containing ligands compared to the corresponding phenolic-OH prototype was determined using [^{35}S]GTP γS binding assays at μ and κ receptors; results are shown in Table 2. Due to the relatively poor binding affinity to δ receptors, these compounds were not evaluated for functional activity at δ . Procedures similar to those previously reported were used.²⁹ For the

Table 2

 EC_{50} and E_{max} values for the stimulation of [³⁵S]GTP γ S binding and IC₅₀ and I_{max} values for the inhibition of agonist-stimulated [³⁵S]GTP γ S binding to the human μ and κ opioid receptors^a

Mu opioid receptorNDAMGOAgonist/antagonist 55 ± 7 116 ± 4 N ^{Ib} NIIaAgonist/antagonist 10 ± 0.70 20 ± 2.4 13 ± 2.2 67 ± 3.1 IbWeak agonist/antagonist 10 ± 0.70 20 ± 2.4 15 ± 4.0 76 ± 3.6 GaAgonist/antagonist 24 ± 7.3 23 ± 1.5 25 ± 15 70 ± 2.6 GbAgonist/antagonist 21 ± 7.3 23 ± 1.5 25 ± 15 70 ± 2.6 TaAgonist/antagonist 2.3 ± 0.87 110 ± 2.9 NINI7bAgonist/antagonist 2.7 ± 0.72 31 ± 4.9 17 ± 3.3 61 ± 3.9 SbWeak agonist/antagonist 2.7 ± 0.72 31 ± 4.9 17 ± 3.3 61 ± 3.9 9aWeak agonist/antagonist 10 ± 5.0 42 ± 2.0 250 ± 36 68 ± 2.9 10aAgonist/antagonist 21 ± 3.2 39 ± 0.21 130 ± 24 63 ± 5.2 10bAgonist/antagonist 21 ± 3.2 39 ± 0.21 130 ± 24 63 ± 5.2 10bAgonist/antagonist 11 ± 0.2 39 ± 0.21 130 ± 24 63 ± 5.2 10bAgonist/antagonist 11 ± 0.2 39 ± 0.21 130 ± 24 63 ± 5.2 10bAgonist/antagonist 11 ± 0.2 39 ± 0.21 130 ± 2.4 63 ± 5.2 10bAgonist/antagonist 12 ± 1.4 23 ± 1.5 50 ± 1.5 81 ± 0.3 12bWeak agonist/antagonist 11 ± 0.2 2.5 ± 0.5 10 ± 0.5 52 ± 3.5 10b	Compound	Functional description	EC ₅₀ (nM)	E_{\max} (% maximal stimulation)	IC ₅₀ (nM)	I _{max} (% maximal inhibition)
DAMCOAgonist55 ± 7116 ± 4NIPNI1aAgonist/antagonist0 ± 1.324 ± 7.713 ± 2.267 ± 3.11bWeak agonist/antagonist10 ± 0.7020 ± 2.415 ± 4.076 ± 3.66aAgonist/antagonist40 ± 4.678 ± 4.6NININI6bAgonist/antagonist27 ± 3.179 ± 0.82NININI7aAgonist/antagonist27 ± 3.179 ± 0.82NININI8aAgonist/antagonist2.7 ± 0.7231 ± 4.917 ± 3.361 ± 3.98bWeak agonist/antagonistNA~20%S8 ± 1489 ± 3.99bAgonist/antagonist10 ± 5.042 ± 2.0250 ± 3668 ± 2.910aAgonist/antagonist21 ± 3.239 ± 0.2110 ± 0.2 ± 0.363 ± 5.210bAgonist/antagonist4.2 ± 1.423 ± 1.510 ± 2.463 ± 5.210bAgonist/antagonist11 ± 0.230 ± 5.668 ± 5.910bAgonist/antagonist1.1 ± 0.2132 ± 0.2130 ± 5.668 ± 5.910bAgonist/antagonistNA22 ± 5.8NININI17bAntagonist1.1 ± 0.2132 ± 0.2148 ± 0.5059 ± 0.2148 ± 0.5717bAgonist/antagonist1.1 ± 0.2057 ± 3.8NININI17bAgonist/antagonist1.2 ± 0.5550 ± 2.4NININI17bAgonist/antagonist1.3 ± 0.2057 ± 3.8NI <td< td=""><td>Mu opioid recepto</td><td>r</td><td></td><td></td><td></td><td></td></td<>	Mu opioid recepto	r				
iaAgonist/antagonist40 ± 1.324 ± 2.713 ± 2.267 ± 3.1ibWeak agonist/antagonist10 ± 0.7020 ± 2.415 ± 4.076 ± 3.66aAgonist/antagonist24 ± 7.323 ± 1.510 ± 2.6NINI6bAgonist/antagonist27 ± 3.179 ± 0.82NININI7bAgonist/antagonist2.7 ± 0.7231 ± 4.9NININI8aAgonist/antagonistNA~20%NININI9bAgonist/antagonist11 ± 5.239 ± 0.2130 ± 5.250 ± 5.361 ± 5.99aWeak agonist/antagonist21 ± 5.239 ± 0.21100 ± 5.0 ± 5.773 ± 3.110aAgonist/antagonist21 ± 5.239 ± 0.21100 ± 5.0 ± 5.773 ± 3.111aWeak agonist/antagonistNA22 ± 5.814 ± 3.354 ± 2.711bWeak agonist/antagonistNA21 ± 5.605 ± 5.773 ± 3.111aWeak agonist/antagonistNA11 ± 0.60.50 ± 5.773 ± 3.111bWeak agonist/antagonistNA12 ± 6.20.50 ± 5.0 ± 7.248 ± 0.8717bAntagonist31 ± 0.2057 ± 3.8NINI17aAgonist/antagonist1.3 ± 0.2057 ± 3.8NINI17aAgonist/antagonist1.3 ± 0.2057 ± 3.8NINI17aAgonist/antagonist1.3 ± 0.2057 ± 3.8NINI17aAgonist/antagonist1.3 ± 0.20NI <th< td=""><td>DAMGO</td><td>Agonist</td><td>55 ± 7</td><td>116 ± 4</td><td>NI^b</td><td>NI</td></th<>	DAMGO	Agonist	55 ± 7	116 ± 4	NI ^b	NI
1bWeak agonist/antagonist1.0 ± 0.7020 ± 2.415 ± 4.076 ± 3.66aAgonist0.0 ± 4.678 ± 4.6NINI6bAgonist24 ± 7.323 ± 1.525 ± 1.570 ± 2.67aAgonist27 ± 3.179 ± 0.82NINI7bAgonist/antagonist27 ± 0.7231 ± 4.917 ± 3.361 ± 3.98bWeak agonist/antagonist2.7 ± 0.7231 ± 4.975 ± 3.861 ± 3.98bWeak agonist/antagonistNA~20%NINI9bAgonist/antagonist10 ± 5.042 ± 2.020 ± 3.668 ± 2.910aAgonist/antagonist10 ± 5.042 ± 2.0130 ± 2.668 ± 2.910bAgonist/antagonist12 ± 1.423 ± 1.5150 ± 5.773 ± 3.111aWeak agonist/antagonistNA11 ± 2.630 ± 5.686 ± 6.917aAgonist/antagonistNA12 ± 6.20.5 ± 0.2148 ± 0.8717bMatogonist1.3 ± 0.2077 ± 1.1NININI17bAgonist/antagonist1.3 ± 0.2057 ± 3.8NININI16aAgonist/antagonist1.3 ± 0.2057 ± 3.8NININI17bAgonist/antagonist1.3 ± 0.0559 ± 2.4NININI16aAgonist/antagonist1.3 ± 0.0559 ± 2.3NININI17bAgonist/antagonist30 ± 9.951 ± 3.6NININI16bN	1a	Agonist/antagonist	4.0 ± 1.3	24 ± 2.7	13 ± 2.2	67 ± 3.1
GaAgonist40.24.6678.4.6NINI6bAgonist/antagonist27.4.3.179.4.08225.1.570.4.2.67nAgonist27.4.3.179.4.082NINI7bAgonist/antagonist5.3.4.0.87110.4.2.9NINI8aAgonist/antagonist27.4.0.7.231.4.9.917.4.3.361.4.3.98bWeak agonist/antagonistNA~20%NINI9bAgonist/antagonist101.4.5.042.4.0.25.9.4.668.4.2.910aAgonist/antagonist21.4.3.239.0.2.1130.2.4.463.4.5.210bAgonist/antagonist21.4.3.239.0.2.1130.2.4.463.4.5.211aWeak agonist/antagonistNA11.4.2.630.5.5.86.4.6.917bAntagonist0.11.1.0.0.2132.4.6.230.4.5.686.4.6.917bAntagonist0.11.1.0.0.2132.4.6.230.4.5.686.4.6.917bAntagonist13.4.0.2077.4.1.1NINI17bAntagonist1.4.0.2132.4.6.230.4.0.2.057.4.8.417bAgonist/antagonist1.3.4.0.2057.4.3.4NINI17bAgonist/antagonist1.4.0.2130.4.7.0NINI17bAgonist/antagonist1.4.0.2150.4.3.4NINI17bAgonist/antagonist1.4.0.2150.4.3.4NINI17bAgonist/antagonist1.4.0.2150.4.3.4NINI	1b	Weak agonist/antagonist	1.0 ± 0.70	20 ± 2.4	15 ± 4.0	76 ± 3.6
6bAgonisr(antagonist)24 ± 7.323 ± 1.570 ± 2.67aAgonist27 ± 3.179 ± 0.82NINI7aAgonist5.3 ± 0.87110 ± 2.9NINI8aAgonisr(antagonist2.7 ± 0.7231 ± 4.971 ± 3.361 ± 3.98bWeak agonisr(antagonistNA~20%58 ± 1.489 ± 3.99aWeak agonisr(antagonistNA~20%S1 ± 1.489 ± 3.99aMeak agonisr(antagonist101 ± 5042 ± 2.0250 ± 3668 ± 2.910aAgonisr(antagonist2.1 ± 3.239 ± 0.21130 ± 2.463 ± 5.210bAgonisr(antagonist4.2 ± 1.42.3 ± 1.514 ± 3.354 ± 2.711aWeak agonisr(antagonistNA11 ± 2.630 ± 5.686 ± 6.917aAgonisr(antagonistNA11 ± 2.60.5 ± 0.24.5 ± 0.8717bMagonisr(antagonistNA11 ± 2.60.5 ± 0.257 ± 8.417bAgonisr(antagonistNA9.0 ± 7.04.0 ± 0.8057 ± 8.417bAgonisr(antagonist1.3 ± 0.2057 ± 3.8NININI18aAgonisr1.3 ± 0.2057 ± 3.8NINININI19bAgonisr(antagonist4.4 ± 0.6559 ± 2.4NINININI16aAgonisr(antagonist4.4 ± 0.6559 ± 2.3NINININI16bAgonisr(antagonist4.4 ± 0.6559 ± 2.4NININI <t< td=""><td>6a</td><td>Agonist</td><td>40 ± 4.6</td><td>78 ± 4.6</td><td>NI</td><td>NI</td></t<>	6a	Agonist	40 ± 4.6	78 ± 4.6	NI	NI
7aAgonist27±3.179±0.82NININI7bAgonist53±0.87110±2.9NININI7bAgonist/antagonist2.7±0.7231±4.917±3.361±3.98bWeak agonist/antagonistNA~20%NINI9aWeak agonist/antagonistNA~20%NINI9bAgonist/antagonist101±5042±2.0250±3668±2.910aAgonist/antagonist11±3.239±0.1130±2.463±5.210bAgonist/antagonist4.2±1.423±1.5150±5773±3.111aWeak agonist/antagonistNA12±2.620.55±0.2148±0.8717aAgonist/antagonist0.11±0.02132±6.20.55±0.2148±0.8717bAntagonist0.11±0.02132±6.20.55±0.2148±0.8717bAntagonist13±0.2057±3.8NININI18aAgonist13±0.2057±3.8NININI19aAgonist12±0.2659±2.4NININI16aAgonist12±0.2659±2.3NININI19bAgonist/antagonist4.4±1.260±2.664±1.3749±4.419bAgonist/antagonist14±1.183±10NINI19bAgonist/antagonist0.9±9.775±3.6NININI19bAgonist/antagonist0.9±9.775±3.6NININI11a83±10 <td>6b</td> <td>Agonist/antagonist</td> <td>24 ± 7.3</td> <td>23 ± 1.5</td> <td>25 ± 15</td> <td>70 ± 2.6</td>	6b	Agonist/antagonist	24 ± 7.3	23 ± 1.5	25 ± 15	70 ± 2.6
7bAgonistS3 0.87110 ± 0.9NININI8aAgonist/antagonist2.7 ± 0.7231 ± 4.917 ± 3.361 ± 3.08bWeak agonist/antagonistNA~20%S5 ± 1.489 ± 3.99aWeak agonist/antagonistNA~20%NINI9bAgonist/antagonist10 ± 5.030 ± 0.2130 ± 2.463 ± 5.210aAgonist/antagonist2.1 ± 3.239 ± 0.21130 ± 2.463 ± 5.210bAgonist/antagonist2.1 ± 3.239 ± 0.21150 ± 5.773 ± 3.111aWeak agonist/antagonistNA2.2 ± 5.814 ± 3.354 ± 2.711bWeak agonist/antagonistNA11 ± 2.630 ± 5.686 ± 6.917aAgonist/antagonistNA11 ± 2.630 ± 5.686 ± 6.917bAgonist/antagonistNA12 ± 6.20.5 ± 0.2184 ± 0.8717bAgonist1.3 ± 0.207 ± 1.1NININI17bAgonist36 ± 5.07 ± 1.3NININI1aAgonist1.2 ± 0.655 ± 2.4NININI1aAgonist1.2 ± 0.655 ± 2.4NININI6aAgonist1.0 ± 0.35 ± 4.5NININI1aAgonist1.0 ± 0.35 ± 4.5NININI6aAgonist1.0 ± 0.35 ± 4.5NININI1aAgonist3.0 ± 9.97 ± 3.6Agonist<	7a	Agonist	27 ± 3.1	79 ± 0.82	NI	NI
8aAgonist/antagonist2.7 ± 0.7231 ± 9.91.7 ± 3.36 ± 1.4 98bWeak agonist/antagonistNA~20%NINI9aMeak agonist/antagonist101 ± 504.2 ± 0.0250 ± 3668 ± 2.910aAgonist/antagonist2.1 ± 3.239 ± 0.2130 ± 2.463 ± 5.210bAgonist/antagonist4.2 ± 1.42.3 ± 1.5150 ± 577.3 ± 3.111aWeak agonist/antagonistNA2.2 ± 5.81.4 ± 3.35.4 ± 2.711bWeak agonist/antagonist0.11 ± 0.0213.2 ± 6.20.59 ± 0.214.8 ± 0.8717bAntagonist0.11 ± 0.0213.2 ± 6.20.59 ± 0.214.8 ± 0.8717bAntagonist0.11 ± 0.0213.2 ± 6.20.59 ± 0.214.8 ± 0.8717bAntagonist0.11 ± 0.0213.2 ± 6.20.59 ± 0.214.8 ± 0.8717bAgonist/antagonist0.11 ± 0.0213.2 ± 6.20.59 ± 0.214.8 ± 0.8717bAgonist/antagonist0.11 ± 0.0213.2 ± 6.20.59 ± 0.214.8 ± 0.8717bAgonist1.3 ± 0.207.7 ± 1.1NININI16aAgonist3.6 ± 5.07.7 ± 1.3NININI16aAgonist1.20 ± 2.650 ± 2.3NININI16bNTNTNTNTNTNTNTNT7bAgonist/antagonist1.0 ± 0.1360 ± 2.0NININI7bAgonist/antagonist3.0 ± 0.7	7b	Agonist	5.3 ± 0.87	110 ± 2.9	NI	NI
8bWeak agonist/antagonistNA~20%58 ± 1489 ± 3.99aWeak agonist/antagonistNININI9bAgonist/antagonist101 ± 5042 ± 2.0S50 ± 3668 ± 2.910aAgonist/antagonist21 ± 3.239 ± 0.21130 ± 2463 ± 5.210bAgonist/antagonist4.2 ± 1.423 ± 1.5150 ± 5773 ± 3.111aWeak agonist/antagonistNA21 ± 5.2150 ± 5773 ± 3.111bWeak agonist/antagonistNA21 ± 5.20.5 ± 5.086 ± 5.917aAgonist/antagonistNA11 ± 2.630 ± 5.086 ± 5.917bAtagonistNA9.0 ± 7.04.0 ± 0.8086 ± 5.917aAgonist/antagonistNA9.0 ± 7.04.0 ± 0.8086 ± 5.917bAtagonist36 ± 5.077 ± 1.1NININI1aAgonist36 ± 5.059 ± 2.4NININI1aAgonist2.0 ± 2.559 ± 2.4NININI1aAgonist2.0 ± 2.659 ± 2.4NININI1aAgonist2.0 ± 2.559 ± 2.4NININI1aAgonist2.0 ± 2.559 ± 2.4NININI1aAgonist2.0 ± 2.559 ± 2.4NININI7aAgonist3.0 ± 0.1360 ± 2.664 ± 1.379 ± 4.45bAgonist3.0 ± 0.1360 ± 2.664 ± 1.379 ± 4.4<	8a	Agonist/antagonist	2.7 ± 0.72	31 ± 4.9	17 ± 3.3	61 ± 3.9
9aWeak agonistNA~20%NINI9bAgonist/antagonist101±5042±0.250±3668±2.910aAgonist/antagonist11±3.239±0.21130±2463±5.210bAgonist/antagonist4.2±1.423±1.5150±5773±3.111aWeak agonist/antagonistNA21±5.814±3.354±2.711bWeak agonist/antagonistNA11±2.60.59±0.2148±0.8717bAgonist/antagonist0.11±0.02132±6.20.59±0.2148±0.8717bAtgonist0.11±0.02132±6.20.59±0.2148±0.8717bAgonist36±5.077±1.1NINI18aAgonist1.3±0.2057±3.8NINI19aAgonist1.3±0.2057±3.8NININI19aAgonist1.3±0.2057±3.8NININI19aAgonist1.20±2659±2.4NININI19aAgonist1.20±2659±2.4NININI7aAgonist1.20±2659±2.4NININI7bAgonist/antagonist4.11.260±2.0NININI7bAgonist/antagonist1.0±0.1369±2.664±1.3749±4.48aAgonist0.1±0.1360±2.0NININI7bAgonist30±9.975±3.6NTNTNI9bAgonist0.9±9.975±3.6NI	8b	Weak agonist/antagonist	NA	~20%	58 ± 14	89 ± 3.9
9bAgonist/antagonist101 ± 50 42 ± 2.0 250 ± 3668 ± 2.910aAgonist/antagonist 21 ± 3.2 99 to 21130 ± 2463 ± 5.210bAgonist/antagonist 42 ± 1.4 23 ± 1.5150 ± 5773 ± 3.111aWeak agonist/antagonistNA 22 ± 5.8 14 ± 3.354 ± 2.711bWeak agonist/antagonistNA 21 ± 2.6 0.59 ± 0.21 48 ± 0.87 17aAgonist/antagonist0.11 ± 0.021 32 ± 6.2 0.59 ± 0.21 48 ± 0.87 17bAntagonist 0.11 ± 0.021 32 ± 6.2 0.59 ± 0.21 48 ± 0.87 17bAntagonist 0.11 ± 0.021 32 ± 6.2 0.59 ± 0.21 48 ± 0.87 17bAntagonist 0.11 ± 0.021 32 ± 6.2 0.59 ± 0.21 48 ± 0.87 17bAntagonist 0.11 ± 0.021 32 ± 6.2 0.59 ± 0.21 48 ± 0.87 17bAntagonist 36 ± 5.0 77 ± 11 NININI16aAgonist $12 0.20$ 57 ± 3.8 NININI16aAgonist 120 ± 26 59 ± 2.4 NININI17bAgonist/antagonist 44 ± 1.2 60 ± 2.6 640 ± 137 49 ± 4.4 7bAgonist/antagonist 10 ± 0.33 60 ± 2.6 NININI7bAgonist 10 ± 0.33 60 ± 2.6 NININI7bAgonist $10 \pm 0.33 \pm 0.17$ 84 ± 1.2 NININI7bAgoni	9a	Weak agonist	NA	~20%	NI	NI
10aAgonist/antagonist21 ± 3.2 39 ± 0.21 130 ± 24 63 ± 5.2 10bAgonist/antagonist4.2 ± 1.4 23 ± 1.5 150 ± 57 73 ± 3.1 11aWeak agonist/antagonistNA 22 ± 5.8 14 ± 3.3 54 ± 2.7 11bWeak agonist/antagonistNA 22 ± 5.8 14 ± 3.3 54 ± 2.7 11bWeak agonist/antagonist0.11 ± 0.021 32 ± 6.2 0.59 ± 0.21 48 ± 0.87 17bAntagonist0.11 ± 0.021 32 ± 6.2 0.59 ± 0.21 48 ± 0.87 Kappa opioid receptUU 0.57 ± 3.8 NINIU50.488Agonist 1.3 ± 0.20 57 ± 3.8 NININI1aAgonist 2.4 ± 0.65 59 ± 2.4 NININI1bAgonist 2.4 ± 0.65 59 ± 2.4 NININI1bAgonist 0.10 ± 2.6 50 ± 2.3 NININI1bAgonist 0.10 ± 2.6 50 ± 2.3 NININI1bAgonist 0.10 ± 2.6 64 ± 1.37 99 ± 4.4 6aAgonist 0.10 ± 0.53 50 ± 5.5 NININI1bAgonist 3.0 ± 9.9 75 ± 3.6 NININI1bAgonist 0.2 ± 9.9 57 ± 3.6 NININI1bAgonist 3.0 ± 0.7 84 ± 1.2 NININI1bAgonist 3.9 ± 0.1 75 ± 3.6 NININI1b </td <td>9b</td> <td>Agonist/antagonist</td> <td>101 ± 50</td> <td>42 ± 2.0</td> <td>250 ± 36</td> <td>68 ± 2.9</td>	9b	Agonist/antagonist	101 ± 50	42 ± 2.0	250 ± 36	68 ± 2.9
10bAgonist/antagonist 4.2 ± 1.4 23 ± 1.5 150 ± 57 73 ± 3.1 11aWeak agonist/antagonistNA 22 ± 5.8 14 ± 3.3 54 ± 2.7 11bWeak agonist/antagonistNA 11 ± 2.6 30 ± 5.6 86 ± 6.9 17aAgonist/antagonist0.11 ± 0.021 32 ± 6.2 59 ± 0.21 48 ± 0.87 17bAntagonistNA 9.0 ± 7.0 4.0 ± 0.80 57 ± 8.4 <i>Kappa opioid receptor</i> US0.488Agonist 36 ± 5.0 77 ± 11 NINI1aAgonist 36 ± 5.0 77 ± 13.8 NINI1aAgonist 2.4 ± 0.65 59 ± 2.4 NINI6aAgonist 1.20 ± 2.6 50 ± 2.3 NINI6bNTNTNTNTNTNT7aAgonist 8.1 ± 0.93 95 ± 4.5 NINI7bAgonist 1.0 ± 0.13 60 ± 2.0 NINI7bAgonist 30 ± 9.9 75 ± 3.6 NTNT7bAgonist 30 ± 9.9 75 ± 3.6 NTNT9aAgonist 3.3 ± 0.17 84 ± 1.2 NINI9bAgonist 3.3 ± 0.17 84 ± 1.2 NINI9bAgonist 3.2 ± 1.0 73 ± 8.8 NINI11bAgonist 3.6 ± 0.41 60 ± 0.97 NINI11aAgonist 3.8 ± 0.42 55 ± 4.1 NINI<	10a	Agonist/antagonist	21 ± 3.2	39 ± 0.21	130 ± 24	63 ± 5.2
11aVeak agonist/antagonistNA 22 ± 5.8 14 ± 3.3 54 ± 2.7 11bWeak agonist/antagonistNA 11 ± 2.6 30 ± 5.6 86 ± 6.9 17aAgonist/antagonistNA 9.0 ± 7.0 0.59 ± 0.21 48 ± 0.87 17bAntagonistNA 9.0 ± 7.0 4.0 ± 0.80 57 ± 8.4 Kappa opioid receptorUS0,488Agonist 36 ± 5.0 77 ± 11 NININI1aAgonist 36 ± 5.0 77 ± 13.8 NININI1bAgonist 2.4 ± 0.65 59 ± 2.4 NININI6aAgonist 120 ± 26 50 ± 2.3 NININI6bNTNTNTNTNTNT7aAgonist 8.1 ± 0.99 95 ± 4.5 NININI7bAgonist 30 ± 9.9 75 ± 3.6 NININI8bAgonist 30 ± 9.9 75 ± 3.6 NTNTNT9bAgonist 30 ± 9.9 75 ± 3.6 NTNTNT9bAgonist 30 ± 9.9 75 ± 3.6 NTNTNT9bAgonist 3.3 ± 0.17 84 ± 1.2 NININI10bAgonist 3.3 ± 0.17 84 ± 1.2 NININI111Agonist 3.8 ± 0.42 55 ± 4.1 NININI111Agonist 9.6 ± 4.1 60 ± 0.97 NININI <tr< td=""><td>10b</td><td>Agonist/antagonist</td><td>4.2 ± 1.4</td><td>23 ± 1.5</td><td>150 ± 57</td><td>73 ± 3.1</td></tr<>	10b	Agonist/antagonist	4.2 ± 1.4	23 ± 1.5	150 ± 57	73 ± 3.1
11bWeak agonist/antagonistNA 11 ± 2.6 30 ± 5.6 86 ± 6.9 17aAgonist/antagonist 0.11 ± 0.021 32 ± 6.2 0.59 ± 0.21 48 ± 0.87 17bAntagonistNA 9.0 ± 7.0 40 ± 0.80 57 ± 8.4 Kappa opioid receptUS0,488Agonist 36 ± 5.0 77 ± 11 NINI1aAgonist 36 ± 5.0 77 ± 11 NININI1aAgonist 2.4 ± 0.65 59 ± 2.4 NININI6bNTNTNTNTNTNT7aAgonist 8.1 ± 0.93 95 ± 4.5 NININI7bAgonist/antagonist 4.4 ± 1.2 60 ± 2.0 NININI7bAgonist 0 ± 9.9 7 ± 3.6 NTNTNT9aAgonist 0 ± 9.9 7 ± 3.6 NTNTNI10bAgonist 0 ± 9.9 7 ± 3.6 NTNT9aAgonist 0 ± 9.9 7 ± 3.6 <	11a	Weak agonist/antagonist	NA	22 ± 5.8	14 ± 3.3	54 ± 2.7
17aAgonist/antagonist 0.11 ± 0.021 32 ± 6.2 0.59 ± 0.21 48 ± 0.87 17bAntagonistNA 9.0 ± 7.0 4.0 ± 0.80 57 ± 8.4 Kappa opioid receptorU50,488Agonist 36 ± 5.0 77 ± 11 NININI1aAgonist 36 ± 5.0 77 ± 13 NININI1aAgonist 2.4 ± 0.65 59 ± 2.4 NININI6aAgonist 120 ± 26 50 ± 2.3 NININI6bNTNTNTNTNTNT7aAgonist/antagonist 8.1 ± 0.93 95 ± 4.5 NININI7bAgonist/antagonist 1.0 ± 0.13 60 ± 2.0 NININI7bAgonist/antagonist 1.0 ± 0.13 60 ± 2.0 NININI8bAgonist 30 ± 9.9 75 ± 3.6 NTNTNT9bAgonist 30 ± 9.9 75 ± 3.6 NTNTNT9bAgonist 3.3 ± 0.17 84 ± 1.2 NININI9bAgonist 3.3 ± 0.17 84 ± 1.2 NININI10bAgonist 3.8 ± 0.42 55 ± 4.1 NININI11bAgonist/antagonist 3.8 ± 0.42 55 ± 4.1 NINI11bAgonist/antagonist 3.8 ± 0.42 55 ± 5.1 NINI17bAntagonist/antagonistNA -1.3 ± 2.5 15 \pm 6.8	11b	Weak agonist/antagonist	NA	11 ± 2.6	30 ± 5.6	86 ± 6.9
17bAntagonistNA9.0 ± 7.04.0 ± 0.8057 ± 8.4Kappa opioid receptorU50,488Agonist36 ± 5.077 ± 1.1NINI1aAgonist13 ± 0.2057 ± 3.8NINI1bAgonist2.4 ± 0.6559 ± 2.4NINI6aAgonist120 ± 2.650 ± 2.3NINI6bNTNTNTNTNT7aAgonist8.1 ± 0.9395 ± 4.5NINI7bAgonist1.0 ± 0.1360 ± 2.6640 ± 13749 ± 4.47bAgonist1.0 ± 0.1360 ± 2.6NINI7bAgonist1.0 ± 0.1360 ± 2.6NINI7bAgonist1.0 ± 0.1360 ± 2.6NINI7bAgonist1.0 ± 0.1360 ± 2.6NINI7bAgonist1.0 ± 0.1360 ± 2.6NINI7bAgonist3.0 ± 9.975 ± 3.6NINI7bAgonist3.0 ± 9.975 ± 3.6NINI7bAgonist3.3 ± 0.1784 ± 1.2NINI7bAgonist3.3 ± 0.1784 ± 1.2NINI7bAgonist9.6 ± 4.160 ± 0.97NINI7cAgonist3.8 ± 0.425 ± 4.1NINI7cAgonist3.8 ± 0.425 ± 4.1NINI7cAgonist/antagonist0.18 ± 0.01435 ± 2.515 ± 6.82 ±	17a	Agonist/antagonist	0.11 ± 0.021	32 ± 6.2	0.59 ± 0.21	48 ± 0.87
Kappa opioid receptor Vision Agonist 36 ± 5.0 77 ± 1.1 NI NI 1a Agonist 1.3 ± 0.20 57 ± 3.8 NI NI 1b Agonist 2.4 ± 0.65 59 ± 2.4 NI NI 6a Agonist 2.4 ± 0.65 50 ± 2.3 NI NI 6b NT NT NT NT NT 7a Agonist 8.1 ± 0.93 95 ± 4.5 NI NI 7b Agonist 8.1 ± 0.93 95 ± 4.5 NI NI NI 8a Agonist 1.0 ± 0.13 60 ± 2.0 NI NI NI 8b Agonist 1.0 ± 0.13 60 ± 2.0 NI NI NI 8b Agonist 1.0 ± 0.13 60 ± 2.0 NI NI NI 9a Agonist 1.0 ± 0.13 60 ± 2.0 NI NI NI 9b Agonist 3.0 ± 9.9 75 ± 3.6 NI NI NI <tr< td=""><td>17b</td><td>Antagonist</td><td>NA</td><td>9.0 ± 7.0</td><td>4.0 ± 0.80</td><td>57 ± 8.4</td></tr<>	17b	Antagonist	NA	9.0 ± 7.0	4.0 ± 0.80	57 ± 8.4
U50,488Agonist 36 ± 5.0 77 ± 11 NININI1aAgonist 1.3 ± 0.20 57 ± 3.8 NININI1bAgonist 2.4 ± 0.65 59 ± 2.4 NININI6aAgonist 120 ± 26 50 ± 2.3 NININI6bNTNTNTNTNTNT7aAgonist 8.1 ± 0.93 95 ± 4.5 NININI7bAgonist/antagonist 4.4 ± 1.2 60 ± 2.6 640 ± 137 49 ± 4.4 8aAgonist 1.0 ± 0.13 60 ± 2.0 NINI8bAgonist 30 ± 9.9 75 ± 3.6 NTNT9aAgonist 170 ± 29 65 ± 3.3 NTNT9bAgonist 170 ± 29 65 ± 3.3 NTNI10aAgonist 3.3 ± 0.17 84 ± 1.2 NINI11bAgonist 9.6 ± 4.1 60 ± 0.97 NINI11aAgonist 2.9 ± 1.0 73 ± 8.8 NINI11bAgonist 3.8 ± 0.42 55 ± 4.1 NINI11bAgonist/antagonist 0.18 ± 0.014 35 ± 2.5 15 ± 6.8 24 ± 1.7 17bAntagonistNA -1.3 ± 2.5 140 ± 18 59 ± 5.1	Kappa opioid rece	otor				
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1bAgonist 2.4 ± 0.65 59 ± 2.4 NINI6aAgonist 120 ± 26 50 ± 2.3 NINI6bNTNTNTNTNTNT7aAgonist 8.1 ± 0.93 95 ± 4.5 NINI7bAgonist/antagonist 4.4 ± 1.2 60 ± 2.6 640 ± 137 49 ± 4.4 8aAgonist 1.0 ± 0.13 60 ± 2.0 NINI9aAgonist 30 ± 9.9 75 ± 3.6 NTNT9bAgonist 170 ± 29 65 ± 3.3 NTNT10aAgonist 3.3 ± 0.17 84 ± 1.2 NINI10bAgonist 2.9 ± 1.0 73 ± 8.8 NINI11aAgonist 2.9 ± 1.0 73 ± 8.8 NINI11bAgonist/antagonist 0.18 ± 0.014 35 ± 2.5 15 ± 6.8 24 ± 1.7 17bAntagonist 0.18 ± 0.014 35 ± 2.5 140 ± 18 59 ± 5.1	1a	Agonist	1.3 ± 0.20	57 ± 3.8	NI	NI
6a Agonist 120 ± 26 50 ± 2.3 NINI 6b NTNTNTNTNTNT 7a Agonist 8.1 ± 0.93 95 ± 4.5 NINI 7b Agonist/antagonist 4.4 ± 1.2 60 ± 2.6 640 ± 137 49 ± 4.4 8a Agonist 1.0 ± 0.13 60 ± 2.0 NINI 8b Agonist 30 ± 9.9 75 ± 3.6 NTNT 9a Agonist 170 ± 29 65 ± 3.3 NTNT 9b Agonist 170 ± 29 65 ± 3.3 NTNI 10a Agonist 3.3 ± 0.17 84 ± 1.2 NINI 10b Agonist 2.9 ± 1.0 73 ± 8.8 NINI 11a Agonist 3.8 ± 0.42 55 ± 4.1 NINI 17a Agonist/antagonist 0.18 ± 0.014 35 ± 2.5 15 ± 6.8 24 ± 1.7 17b AntagonistNA -1.3 ± 2.5 140 ± 18 59 ± 5.1	1b	Agonist	2.4 ± 0.65	59 ± 2.4	NI	NI
6bNTNTNTNTNTNTNTNT7aAgonist 8.1 ± 0.93 95 ± 4.5 NINI7bAgonist/antagonist 4.4 ± 1.2 60 ± 2.6 640 ± 137 49 ± 4.4 8aAgonist 1.0 ± 0.13 60 ± 2.0 NINI8bAgonist 30 ± 9.9 75 ± 3.6 NTNT9aAgonist 41 ± 11 83 ± 10 NTNT9bAgonist 170 ± 29 65 ± 3.3 NTNI10aAgonist 3.3 ± 0.17 84 ± 1.2 NINI10bAgonist 9.6 ± 4.1 60 ± 0.97 NINI11aAgonist 2.9 ± 1.0 73 ± 8.8 NINI11bAgonist 3.8 ± 0.42 55 ± 4.1 NINI17bAntagonist 0.18 ± 0.014 35 ± 2.5 15 ± 6.8 24 ± 1.7	6a	Agonist	120 ± 26	50 ± 2.3	NI	NI
7aAgonist 8.1 ± 0.93 95 ± 4.5 NINI7bAgonist/antagonist 4.4 ± 1.2 60 ± 2.6 640 ± 137 49 ± 4.4 8aAgonist 1.0 ± 0.13 60 ± 2.0 NINI8bAgonist 30 ± 9.9 75 ± 3.6 NTNT9aAgonist 41 ± 11 83 ± 10 NTNT9bAgonist 170 ± 29 65 ± 3.3 NTNT10aAgonist 3.3 ± 0.17 84 ± 1.2 NINI10bAgonist 9.6 ± 4.1 60 ± 0.97 NINI11aAgonist 2.9 ± 1.0 73 ± 8.8 NINI11bAgonist 3.8 ± 0.42 55 ± 4.1 NINI17aAgonist/antagonist 0.18 ± 0.014 35 ± 2.5 15 ± 6.8 24 ± 1.7 17bAntagonistNA -1.3 ± 2.5 140 ± 18 59 ± 5.1	6b	NT	NT	NT	NT	NT
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8a Agonist 1.0 ± 0.13 60 ± 2.0 NI NI 8b Agonist 30 ± 9.9 75 ± 3.6 NT NT 9a Agonist 41 ± 11 83 ± 10 NT NT 9b Agonist 170 ± 29 65 ± 3.3 NT NT 10a Agonist 3.3 ± 0.17 84 ± 1.2 NI NI 10b Agonist 9.6 ± 4.1 60 ± 0.97 NI NI NI 11a Agonist 2.9 ± 1.0 73 ± 8.8 NI NI NI 11b Agonist 3.8 ± 0.42 55 ± 4.1 NI NI 174 Agonist/antagonist 0.18 ± 0.014 35 ± 2.5 15 ± 6.8 24 ± 1.7 17b Antagonist NA -1.3 ± 2.5 140 ± 18 59 ± 5.1	7b	Agonist/antagonist	4.4 ± 1.2	60 ± 2.6	640 ± 137	49 ± 4.4
8b Agonist 30 ± 9.9 75 ± 3.6 NT NT 9a Agonist 41 ± 11 83 ± 10 NT NT 9b Agonist 170 ± 29 65 ± 3.3 NT NT 10a Agonist 3.3 ± 0.17 84 ± 1.2 NI NI 10b Agonist 9.6 ± 4.1 60 ± 0.97 NI NI 11a Agonist 2.9 ± 1.0 73 ± 8.8 NI NI 11b Agonist 3.8 ± 0.42 55 ± 4.1 NI NI 174 Agonist/antagonist 0.18 ± 0.014 35 ± 2.5 15 ± 6.8 24 ± 1.7 17b Antagonist NA -1.3 ± 2.5 140 ± 18 59 ± 5.1	8a	Agonist	1.0 ± 0.13	60 ± 2.0	NI	NI
9a Agonist 41 ± 11 83 ± 10 NT NT 9b Agonist 170 ± 29 65 ± 3.3 NT NT 10a Agonist 3.3 ± 0.17 84 ± 1.2 NI NI 10b Agonist 9.6 ± 4.1 60 ± 0.97 NI NI 11a Agonist 2.9 ± 1.0 73 ± 8.8 NI NI 11b Agonist 3.8 ± 0.42 55 ± 4.1 NI NI 17a Agonist/antagonist 0.18 ± 0.014 35 ± 2.5 15 ± 6.8 24 ± 1.7 17b Antagonist NA -1.3 ± 2.5 140 ± 18 59 ± 5.1	8b	Agonist	30 ± 9.9	75 ± 3.6	NT	NT
9b Agonist 170 ± 29 65 ± 3.3 NT NT 10a Agonist 3.3 ± 0.17 84 ± 1.2 NI NI 10b Agonist 9.6 ± 4.1 60 ± 0.97 NI NI 11a Agonist 2.9 ± 1.0 73 ± 8.8 NI NI 11b Agonist 3.8 ± 0.42 55 ± 4.1 NI NI 17a Agonist/antagonist 0.18 ± 0.014 35 ± 2.5 15 ± 6.8 24 ± 1.7 17b Antagonist NA -1.3 ± 2.5 140 ± 18 59 ± 5.1	9a	Agonist	41 ± 11	83 ± 10	NT	NT
10a Agonist 3.3 ± 0.17 84 ± 1.2 NI NI 10b Agonist 9.6 ± 4.1 60 ± 0.97 NI NI 11a Agonist 2.9 ± 1.0 73 ± 8.8 NI NI 11b Agonist 3.8 ± 0.42 55 ± 4.1 NI NI 17a Agonist/antagonist 0.18 ± 0.014 35 ± 2.5 15 ± 6.8 24 ± 1.7 17b Antagonist NA -1.3 ± 2.5 140 ± 18 59 ± 5.1	9b	Agonist	170 ± 29	65 ± 3.3	NT	NT
10b Agonist 9.6 ± 4.1 60 ± 0.97 NI NI 11a Agonist 2.9 ± 1.0 73 ± 8.8 NI NI 11b Agonist 3.8 ± 0.42 55 ± 4.1 NI NI 17a Agonist/antagonist 0.18 ± 0.014 35 ± 2.5 15 ± 6.8 24 ± 1.7 17b Antagonist NA -1.3 ± 2.5 140 ± 18 59 ± 5.1	10a	Agonist	3.3 ± 0.17	84 ± 1.2	NI	NI
11a Agonist 2.9 ± 1.0 73 ± 8.8 NI NI 11b Agonist 3.8 ± 0.42 55 ± 4.1 NI NI 17a Agonist/antagonist 0.18 ± 0.014 35 ± 2.5 15 ± 6.8 24 ± 1.7 17b Antagonist NA -1.3 ± 2.5 140 ± 18 59 ± 5.1	10b	Agonist	9.6 ± 4.1	60 ± 0.97	NI	NI
11b Agonist 3.8 ± 0.42 55 ± 4.1 NI NI 17a Agonist/antagonist 0.18 ± 0.014 35 ± 2.5 15 ± 6.8 24 ± 1.7 17b Antagonist NA -1.3 ± 2.5 140 ± 18 59 ± 5.1	11a	Agonist	2.9 ± 1.0	73 ± 8.8	NI	NI
17a Agonist/antagonist 0.18 ± 0.014 35 ± 2.5 15 ± 6.8 24 ± 1.7 17b Antagonist NA -1.3 ± 2.5 140 ± 18 59 ± 5.1	11b	Agonist	3.8 ± 0.42	55 ± 4.1	NI	NI
17b Antagonist NA -1.3 ± 2.5 140 ± 18 59 ± 5.1	17a	Agonist/antagonist	0.18 ± 0.014	35 ± 2.5	15 ± 6.8	24 ± 1.7
	17b	Antagonist	NA	-1.3 ± 2.5	140 ± 18	59 ± 5.1

NA, not applicable.

NT, not tested.

^a See Ref. 29 for experimental details. Data are mean values ± SEM from at least three separate experiments, performed in triplicate. For calculation of the E_{max} values, the basal [³⁵S]GTP γ S binding was set at 0%. For inhibition studies, 200 nM DAMGO was used as the agonist for the μ receptor, U50,488 at final concentration of 100 nM was used for the κ receptor.

^b NI, no inhibition.

benzomorphan pair cyclazocine (1a) and 8-CAC (1b), both are mixed agonists/antagonists at the μ receptor and agonists at κ . Both displayed similar potencies that correlated well with binding affinities. In the case of the **6a/6b** pair (metazocine core) against μ , the carboxamide partner **6a** is an agonist/antagonist while the OH partner is an agonist. Because the affinity of **6b** for κ receptors was relatively weak, it was not studied in the $[^{35}S]$ GTP γ S binding assay. For **7a**/**7b** having a phenazocine core, both were agonists at μ having qualitatively similar potencies. Against κ, **7b** was a mixed agonist/antagonist and 7a was an agonist. The agonist potencies of the two were similar. A divergence in functional activity at the μ receptor was noted for 8a (agonist/antagonist) and 8b (weak agonist/antagonist). At the κ receptor, both were agonists, although **8a** had much higher potency. Divergence was also seen at the µ receptor for the **9a** (weak agonist) and **9b** (agonist/antagonist) pair. Against κ , both were agonists, however, **9a** had much higher potency. For the **10a/10b**, another benzomorphan pair, both were mixed agonists/antagonists at μ and agonists at κ ; agonist potencies at κ were similar. For the butorphanol core, an example of a tetracyclic morphinan, the OH partner **11a** exhibited a weak agonist/antagonist profile at μ , whereas the carboxamide **11b** was an antagonist. At κ , both were agonists having similar potencies. At μ and κ receptors, the carboxamide derivative **17b** of buprenorphine displayed a somewhat different profile (antagonist) than buprenorphine (**17a**) (agonist/antagonist). As μ and κ antagonists, 17a was more potent than 17b. In the $[^{35}S]GTP\gamma S$ binding assay mediated by the κ opioid receptor, some compounds showed a less than maximal activation of $[^{35}S]GTP\gamma S$ binding, but did not have antagonist properties. While most partial agonists have antagonist properties, not all compounds that produce less than a maximal effect in an assay have antagonist effects. Some compounds are less efficacious than an agonist that produces a maximal effect. While many of these compounds have antagonist properties, too, making them partial agonists, there are some compounds that are less efficacious than full agonists but do not have antagonist properties. Compounds 1a, 1b, 6a, 8a and 10b are examples of compounds that have a lower efficacy in the $[^{35}S]GTP\gamma S$ binding assay mediated by the κ receptor, but these compounds do not antagonize $[^{35}S]$ GTP γ S binding that was induced by the κ agonist U50,488.

A series of novel opioids **5b–19b** have been prepared where the phenolic-OH group of traditional and well-studied opioids 5a-19a was replaced by a carboxamide (CONH₂) group. Characterization of target and known compounds in opioid receptor binding assays revealed that carboxamide targets **5b–11b** and **19b** derived from 2,6-methano-3-benzazocine (a.k.a. benzomorphans), morphinan or tramadol-based core structures have high affinity to μ and κ receptors and relatively low affinity for δ receptors. Compared to their phenolic-OH counterparts 5a-11a and 19a, comparable or enhanced affinity was seen for all three receptors. Receptor selectivity for these carboxamides was, in general, similar to the OH partners. It is interesting to note that carboxamide **7b** having the phenazocine core, has KBRs for all three receptors considerably higher than the other 2,6-methano-3-benzazocine cores and has the least degree of selectivity between μ , δ and κ . Another note of interest is the observation that when a divergence in selectivity ratios is seen, there is a trend toward higher selectivity for μ than κ .

A divergent SAR was seen for carboxamides **12b–18b** having the 4,5 α -epoxymorphinan core. Without exception, these carboxamides had lower and in many instances, much lower affinity for μ , δ and κ opioid receptors than their phenolic-OH counterparts

12a–18a. These observations were, in fact, predicted from our earlier studies and are consistent with our pharmacophore hypothesis concerning the bioactive conformation of the carboxamide group.⁴ Receptor selectivity of carboxamides **12b–18b** for μ , δ and κ receptors was, in general, similar to that seen for corresponding OH partners **12a–18a**. SAR findings that we now report are consistent with our previous OH \rightarrow CONH₂ switch studies^{1,3–5} and as well as those from other laboratories.^{5–9} For those OH/CONH₂ pairs studied in [³⁵S]GTP γ S binding assays, within each pair, similarities in their function and potency profiles were frequently observed especially at the κ receptor. At the μ receptor, more divergence was observed with a trend towards the carboxamide partner displaying less agonist activity.

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