# Caged naloxone reveals opioid signaling deactivation kinetics

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### MOL #88096

Running title: Light-triggered opioid antagonism

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Number of text pages: 25

Number of tables: 0

Number of figures: 7

Number of references: 27

Words in Abstract: 216

Words in Introduction: 253

Words in Discussion: 1376

Supplemental tables: 1

Supplemental figures: 5

**Abbreviations:** CNB, carboxynitrobenzyl; CNV, carboxynitroveratryl; CNV-NLX, carboxynitroveratryl-naloxone; CYLE, carboxynitrobenzyl-tyrosine-[Leu<sup>5</sup>]-enkephalin; DAMGO, [D-Ala<sup>2</sup>, N-MePhe<sup>4</sup>, Gly-ol]-enkephalin; DERM, dermorphin; GIRK channel, G protein-coupled K+channel; GPCR, G protein-coupled receptor; LC, locus coeruleus; LE, [Leu<sup>5</sup>]-enkephalin; ME, [Met<sup>5</sup>]-enkephalin; MOR, mu opioid receptor; NLX, naloxone

### **Abstract**

The spatiotemporal dynamics of opioid signaling in the brain remain poorly defined. Photoactivatable opioid ligands provide a means to quantitatively measure these dynamics and their underlying mechanisms in brain tissue. Although activation kinetics can be assessed using caged agonists, deactivation kinetics are obscured by slow clearance of agonist in tissue. In order to reveal deactivation kinetics of opioid signaling we developed a caged competitive antagonist that can be quickly photoreleased in sufficient concentrations to render agonist dissociation effectively irreversible. CNV-NLX, a caged analog of the competitive opioid antagonist naloxone (NLX), was readily synthesized from commercially available NLX in good yield and found to be devoid of antagonist activity at heterologously expressed opioid receptors. Photolysis in slices of rat locus coeruleus produced a rapid inhibition of the ionic currents evoked by multiple agonists of the mu opioid receptor (MOR), but not of α-adrenergic receptors, which activate the same pool of ion channels. Using the high-affinity peptide agonist dermorphin (DERM), we established conditions under which light-driven deactivation rates are independent of agonist concentration and thus intrinsic to the agonist-receptor complex. Under these conditions, some MOR agonists yielded deactivation rates that are limited by G-protein signaling, whereas others appeared limited by agonist dissociation. Therefore, the choice of agonist determines which feature of receptor signaling is unmasked by CNV-NLX photolysis.

### Introduction

Opioid receptors are a family of G protein-coupled receptors (GPCRs) that participate in analgesia, behavioral reinforcement and drug abuse. G protein-mediated signaling is highly dynamic, exemplified by the susceptibility of opioid receptors and other GPCRs to undergo desensitization, trafficking and long term tolerance with prolonged exposure to agonist (von Zastrow and Williams, 2012; von Zastrow et al., 2003; Williams et al., 2001). We previously reported photoactivatable peptide agonists for opioid receptors (Banghart and Sabatini, 2012), which enabled fundamental signaling properties such as G protein-coupled K+ (GIRK) channel activation kinetics to be measured. These are limited by the kinetics of GPCR signaling and are largely ligand independent (Ingram et al., 1997). In contrast, deactivation kinetics can be liganddependent, likely reflecting ligand-receptor dissociation rates (Ingram et al., 1997). However, neither ligand affinity nor GIRK/GPCR coupling kinetics are by necessity constant and may reflect the state of the receptor and the cell. Understanding the determinants of signaling deactivation kinetics is necessary to fully understand the molecular mechanisms that regulate G-protein mediated signaling; however, slow diffusional clearance of agonists in brain tissue, which results in rebinding of dissociated agonist during washout, prolongs receptor activation and limits the temporal resolution of such measurements. In contrast, rapid delivery of antagonist in a diffusionindependent manner should uncover intrinsic deactivation kinetics by preventing rebinding of dissociated agonist. To provide a means for measuring opioid signaling deactivation kinetics as well as a method to block opioid signaling in spatiotemporally delimited manner, we developed a photoactivatable analogue of the broad-spectrum opioid antagonist NLX.

### **Materials and Methods**

### Chemical synthesis and characterization

Commercial reagents were obtained from reputable suppliers and used as received. Naloxone hydrochloride dihydrate was obtained from Sigma-Aldrich and SEM-protected CNVbromide (SEMO-CNV-Br) was prepared by Medicilon (Shanghai, China). All solvents were purchased in septum-sealed bottles stored under an inert atmosphere. All reactions were sealed with septa through which an argon atmosphere was introduced unless otherwise noted. Reactions were conducted in round-bottomed flasks or septum-capped amber screw-cap vials containing Teflon-coated magnetic stir bars. Reactions were monitored by thin layer chromatography (TLC) on precoated TLC glass plates (silica gel 60 F254, 250 µm thickness) or by LC/MS (4.6 mm × 150 mm 5 μm C18 column; 5 μL injection; 30-95% CH<sub>3</sub>CN/H<sub>2</sub>O linear gradient with constant 0.1% v/v TFA additive; 10 min run; 1 ml/min flow; ESI; positive ion mode; UV detection at 210 nm, 254 nm, and 350 nm). TLC chromatograms were visualized by UV illumination. Flash chromatography was performed on an automated purification system using pre-packed silica gel columns. High-resolution mass spectrometry was performed by the Small Molecule Mass Spectrometry Facility at the Harvard FAS Center for Systems Biology on an Agilent 6210 Time-of-Flight LC/MS. NMR spectra were recorded on a Varian 400 MHz spectrometer. <sup>1</sup>H and <sup>13</sup>C chemical shifts (δ) were referenced to residual solvent peaks. Data for <sup>1</sup>H NMR spectra are reported as follows: chemical shift (δ ppm), multiplicity (s = singlet, d = doublet, t = triplet, q = quartet, dd = doublet of doublets, m = multiplet), coupling constant (Hz), integration. Data for <sup>13</sup>C NMR spectra are reported by chemical shift (δ ppm). UV/VIS spectra were recorded on a NanoDrop 2000 UV-Vis Spectrophotometer (Thermo Scientific). Room lights were covered with Rosculux Canary Yellow #312 film to filter out wavelengths of light that would lead to unintentional photolysis during purification and handling.

A vial was charged with naloxone hydrochloride dihydrate (110 mg, 276 nmol) and 4Å, 2.5 µm activated molecular sieves powder (50 mg) and then flushed with argon.

Dimethylformamide (DMF, 400 µI) was added and the resulting suspension was sonicated and then allowed to stir for 15 min before the addition of K<sub>2</sub>CO<sub>3</sub> (84 mg, 607 nmol). SEMO-CNV-Br (128 mg, 304 nmol) dissolved in DMF (500 µl) was added dropwise and the reaction was allowed to stir for 10 h at room temperature while covered with aluminum foil. It was subsequently diluted with ethyl acetate (20 ml) and extracted with saturated NaHCO3 (20 ml, 2x), washed with brine (20 ml, 3x) dried (MgSO4), and filtered. Silica gel (2.5 g) was added and the mixture was concentrated to dryness. Flash chromatography (dry load with silica gel, 0-3% CH<sub>3</sub>OH/CH<sub>2</sub>Cl<sub>2</sub>, linear gradient) provided 178 mg of a pale pink solid, 175 mg of which was immediately dissolved in CH<sub>2</sub>Cl<sub>2</sub> (15 ml) and covered with aluminum foil. TFA (2.5 ml) was added dropwise, which caused a precipitate to form, and the reaction was allowed to stir for 4 h at room temperature. The reaction mixture was concentrated to dryness to provide CNV-NLX as the TFA salt (119 mg, 175 nmol, 66% overall yield). To remove traces of naloxone, this material was further purified in 10-20 mg batches by reverse-phase HPLC (21.2 mm × 150 mm 5 µm Zorbax SB-C8 column (Agilent); 400 µL injections of ~ 20 mM solution in 5% CH<sub>3</sub>CN/H<sub>2</sub>O with 0.1% v/v TFA; 5–60% CH<sub>3</sub>CN/H<sub>2</sub>O linear gradient with constant 0.1% v/v TFA additive; 45 min run; 5 mL/min flow; with the UV/VIS detectors off to avoid photolysis of the purified product). The material eluting between 32-36 minutes was collected, combined, removed of solvent in vacuo, and lyophilized to dryness. Typically, excellent recoveries were obtained (>95 % by weight). Diastereomers were not resolved under these conditions. <sup>1</sup>H NMR (Methanol- $d_4$ , 400 MHz): 7.45-7.61 (m, 1H), 7.19-7.33 (m, 1H), 6.73-6.87 (m, 1H), 6.61-6.71 (m, 1H), 6.47-6.60 (m, 1H), 5.76-5.92 (m, 1H), 5.48-5.61 (m, 2H), 4.78-4.85 (m, 1H), 3.76-3.87 (m, 7H), 3.66-3.74 (m, 1H), 3.45-3.64 (m, 1H), 3.23-3.40 (m, 1H), 2.98-3.15 (m, 1H), 2.78-2.98 (m, 2H), 2.43-2.74 (m, 2H), 2.07-2.17 (m, 1H), 1.79-1.92 (m, 1H), 1.41-1.71 (m, 2H);  $^{13}$ C NMR (Methanol- $d_4$ , 100 MHz); 206.61, 206.5, 153.48, 153.19, 148.63, 148.59, 145.58, 145.55, 141.05, 140.99, 140.80, 140.51, 128.30, 128.00, 127.05, 126.58, 126.52, 125.11, 124.35, 123.73, 121.25, 120.42, 120.26, 119.26, 111.66, 110.25, 108.00, 107.97,  $107.67,\ 107.64,\ 89.75,\ 89.69,\ 89.48,\ 89.42,\ 69.87,\ 69.65,\ 63.03,\ 62.55,\ 62.50,\ 55.52,\ 55.43,$ 49.08, 46.07, 34.22, 30.80, 30.72, 27.36, 22.65; HRMS (ESI) calc'd for  $C_{29}H_{31}N_2O_{10}$  [M+H]<sup>+</sup> 567.1973, found 567.1986.

### Photolysis in solution

Several 40 µl samples of 0.5 mM CNV-NLX in phosphate buffered saline, pH 7.2, contained in a quartz cuvette, were irradiated for various periods of time with ~150 mW of light from a 0.22 NA fiber optic cable (ThorLabs) coupled to a 405 nm laser (Vortran Stradus 405-250) with periodic mixing (Figure S1). Each sample was then analyzed by RP-HPLC (Agilent) on a 4.6 mm x 150 mm Zorbax SB-C8 column (5 µm particle size) in H<sub>2</sub>O, CH<sub>3</sub>CN, 0.1% TFA at 1 ml/min. NLX and CNV-NLX were resolved by a gradient from 20% to 100% CH<sub>3</sub>CN over 10 min. Samples were monitored at 210 nm, 254 nm and 350 nm. Mass spectrometry (Agilent 6420 Triple Quadropole LC/MS) confirmed photolytic conversion of CNV-NLX to NLX (Figure S1).

### Secreted alkaline phosphatase assay

HEK293 cells stably transfected with a chimeric G<sub>s</sub>-G<sub>i</sub> protein that causes G<sub>i</sub>-coupled receptors to activate rather than inhibit adenylate cyclase (Liberles and Buck, 2006; Liberles and Buck, Personal communication) were grown in DMEM (Invitrogen) containing 5% FBS (Invitrogen) and 500 µg/ml G-418 (Invitrogen) and maintained at 37°C in an atmosphere of 5% CO<sub>2</sub>. Cells were plated in 96-well plates at 50,000 cells/well and co-transfected with the GPCR and cAMP-dependent reporter plasmid using Lipofectamine® (Invitrogen) and PLUS® reagent (Invitrogen). The transfection media was replaced with ligand-containing DMEM (200 µl/well) and cells were incubated for 36-48 hrs 37°C/5% CO<sub>2</sub>. Care was taken to avoid exposure to bright, non-filtered light. Each plate was then wrapped in plastic wrap and incubated at 68°C for 2 hrs to heat-inactivate native phosphatases. After transferring 100 µl aliquots from each well to a fresh 96-well plate, 100 µl of an aqueous buffer containing 2 M diethanolamine bicarbonate and 1.2 mM methylumbelliferone phosphate (MUP), pH 10.0, was added to each well. Plates were imaged using a Perkin Elmer Envision plate reader using optical settings for methylumbelliferone fluorescence, taking care to image all conditions to be compared on the same day for a single receptor at the same time-point after MUP addition. In every individual experiment, the NLX was evaluated alongside CNV-NLX at the same receptor and within each plate an agonist-only and

no-ligand control were included for normalization between plates. cDNAs encoding human delta (OPRD1) and human kappa opioid receptors (OPRK1) contained an N-terminal 3xHA epitope tag (Missouri S&T cDNA Resource center, <a href="www.cdna.org">www.cdna.org</a>) and the murine mu opioid receptor (OPRM1) contained an N-terminal FLAG epitope tag. Data shown reflect the average of 6 replicates run in parallel for each condition.

### **Brain slice preparation**

Horizontal brain slices containing locus coeruleus (LC) neurons were prepared as described previously (Williams and North, 1984). Briefly, rats were killed and the brain was removed, blocked and mounted in a vibratome chamber (Leica VT 1200S). Horizontal slices (230 – 240 μm) were prepared in ice-cold cutting solution containing (in mM): 126 NaCl, 2.5 KCl, 1.2 MgCl<sub>2</sub>, 2.6 CaCl<sub>2</sub>, 1.2 NaH<sub>2</sub>PO<sub>4</sub>, 11 D-glucose and 21.4 NaHCO<sub>3</sub> and 0.01 MK801 (Abcam), equilibrated with 95% O<sub>2</sub>/ 5% CO<sub>2</sub>. Slices were stored at 34°C in glass vials with oxygenated (95% O<sub>2</sub>/ 5% CO<sub>2</sub>) artificial cerebrospinal fluid (ACSF) containing (in mM): 126 NaCl, 2.5 KCl, 1.2 MgCl<sub>2</sub>, 2.6 CaCl<sub>2</sub>, 1.2 NaH<sub>2</sub>PO<sub>4</sub>, 11 D-glucose and 21.4 NaHCO<sub>3</sub>.

### Electrophysiological recordings

After an incubation period of at least 30 min, slices were hemisected and transferred to the recording chamber which was superfused with 34°C ACSF at a rate of 1.5 - 2 ml/min. Wholecell recordings were made from LC neurons with an Axopatch 1D amplifier in voltage-clamp mode ( $V_{hold} = -60$  mV). Recording pipettes (1.7 – 2.1 M $\Omega$ ) were filled with internal solution containing (in mM): 115 potassium methanesulfonate or potassium methyl sulfate, 20 KCl, 1.5 MgCl<sub>2</sub>, 5 HEPES(K), 10 BAPTA or 0.1 EGTA, 2 Mg-ATP, 0.2 Na-GTP, pH 7.4, 275-280 mOsM. Series resistance was monitored without compensation and remained < 15 M $\Omega$  for inclusion. Current was continuously recorded at 200 Hz with PowerLab (Chart version 5.4.2; AD Instruments, Colorado Springs, CO). Episodic currents were recorded at 10 kHz for 1 min using AxoGraphX (1.4.3). Drugs were applied by bath superfusion.

Uncaging was carried out with full field illumination using a 405 nm LED (Thorlabs) coupled through a 60x objective (Olympus) of either 0.9 or 1.0 NA. Light power was 10 mW at the back aperture of the objective and 1 mW at the focal point. A solution containing CNV-NLX (5 μM, in 6 ml) was re-circulated for at least 5 min before an uncaging experiment. Opioid agonists were added to the solution containing CNV-NLX prior to photolysis. In most experiments slices were exposed to only a single light flash. In all experiments a saturating concentration of the α2-adrenoceptor agonist UK14304 (3 μM) was applied and the resulting current was used as a postsynaptic control. In many experiments the amplitude of opioid induced current was presented as a fraction of the current induced by UK14304. The small molecule opioids oxycodone, codeine, morphine, etorphine and methadone were obtained from NIDA. ME, DAMGO and endomorphin-1 were obtained from Sigma. β-endorphin was obtained from Peptides International. Dermorphin was obtained from Phoenix Peptides. UK14304 and Idazoxan were obtained from Tocris.

### Data analysis

Summary data are presented as Mean  $\pm$  SEM. The time constant of decay was fit to a single exponential using axographX. The time courses covered the area from at least 5-95% of the total time course of the decline in agonist induced current. Summary data were analyzed using the indicated statistical tests and differences were considered significant when p < 0.05.

### **Results**

### Design and synthesis of caged naloxone

We previously found that appending an α-carboxy-2-nitrobenzyl (CNB) caging group to the phenol on the conserved N-terminal tyrosine of the opioid peptides dynorphin (1-8) and [Leu<sup>5</sup>]-enkephalin (LE) sufficiently reduced agonist potency to achieve strong photoactivation (*i.e.*, CYLE (1), Figure 1) (Banghart and Sabatini, 2012). The significance of this phenol in opioid ligand-receptor interactions is underscored in the morphinan family of opiate alkaloids, which contain a small molecule surrogate of the N-terminal tyrosine in the form of a tyramine moiety that includes the oxygenated A-ring (Kane et al., 2006). For example, the potent MOR agonist morphine (2) is converted to the low affinity agonist codeine (3), simply by addition of a methyl group to the 3-hydroxy moiety (Toll et al., 1998). By extension, we reasoned that the commercially available opioid antagonist NLX (4, Figure 1), which contains a similar morphinan core structure, could be readily caged at the same position.

Although NLX contains several reactive groups, the nucleophilic 3-phenol can be chemoselectively alkylated due to its low  $pK_a$  and steric accessibility (Krassnig et al., 1998). Thus, commercially available NLX was treated under mildly basic conditions with SEMO-CNV-Br (5, Figure 1) to install the  $\alpha$ -carboxy-6-nitroveratryl (CNV) caging group (Russell et al., 2010; Russell et al., 2012). After protecting group removal with trifluoroacetic acid, CNV-NLX (6) was isolated in 66% overall yield as a mixture of diastereomers.

Similar to the CNB caging group used in CYLE, the CNV moiety contains a carboxylic acid to reduce the net hydrophobicity of the caging group, which improves solubility, and may advantageously interact electrostatically with the receptor to impair binding. In contrast to CNB, CNV contains two methoxy moieties at the 4 and 5 positions on the nitroaromatic ring, which extend the absorbance spectrum into the visible wavelength range (Figure S1A) (Russell et al., 2010; Russell et al., 2012). This offers advantages over caging groups that only respond to UV light in terms of the cost of effective light sources for photolysis, reduced scattering in tissue and

lower phototoxicity (Trigo et al., 2009). LC-MS analysis confirmed that CNV-NLX cleanly converts to NLX upon illumination at 405 nm (Figure S1B-F).

### Pharmacological evaluation in heterologously expressed opioid receptors

The functional inactivity of CNV-NLX as an antagonist of opioid receptors was assayed in a heterologous expression system. HEK293 cells were transfected with MOR along with a chimeric  $G_s$ - $G_i$  alpha subunit that causes  $G_i$ -coupled receptors to activate, rather than inhibit, adenylate cyclase, and a CREB-dependent reporter plasmid that leads to the production of secreted alkaline phosphatase (SEAP) as a function of receptor activation through the cAMP pathway (Banghart and Sabatini, 2012; Liberles and Buck, 2006; Liberles and Buck, *Personal communication*). Antagonist dose-response curves were obtained by treating cells with 1  $\mu$ M LE, a concentration that produces near maximal activation, along with various concentrations of NLX or CNV-NLX. Although NLX produced dose dependent inhibition, CNV-NLX exhibited no antagonist activity (Figure 2A). Separately, agonist dose response curves were obtained for LE in the presence of 5  $\mu$ M NLX and 5  $\mu$ M CNV-NLX. Although NLX completely suppressed the LE response, CNV-NLX was without effect (Figure 2B). Similar results were obtained at delta and kappa receptors (Figure S2).

### Light-triggered antagonism in brain tissue

We next asked whether photolysis of CNV-NLX in brain tissue leads to rapid antagonism of opioid receptors and whether this antagonism is specific for opioid receptors over α-adrenergic receptors, which activate the same signaling pathway. To this end, whole-cell voltage-clamp recordings were obtained from acute slices of rat locus coeruleus (LC), which exhibit large outward GIRK currents when exposed to MOR agonists, as well as α-adrenergic receptor agonists (Fiorillo and Williams, 1996; Williams and North, 1984). High concentrations of various agonists were applied to activate GIRK channels in the presence of CNV-NLX (5 μM), followed by full-field delivery of a 5 s flash of 405 nm light (Figure 3). The currents evoked by the MOR agonists [D-Ala², N-MePhe⁴, Gly-ol]-enkephalin (DAMGO), Dermorphin (DERM), morphine,

methionine enkephalin (ME), and methadone were rapidly inhibited immediately following the light flash, followed by a gradual recovery, consistent with photo-release of NLX, the dilution of NLX and agonist rebinding over time (Figure 3A). In contrast, the current evoked by the α-adrenergic receptor agonist UK14303 was not affected by light (Figure 3A, B and Figure S3). These data confirm that photolysis of CNV-NLX can strongly suppress opioid signaling evoked by high concentrations of many common MOR agonists, and that this suppression is not due to a non-specific action on GPCRs.

To examine the ability to control the amount of released antagonist with light, a subsaturating concentration of DAMGO (100 nM) and CNV-NLX (5 μM) were applied and the duration of the light flash was varied. Sub-saturating concentrations of MOR agonists activate sustained outward currents that do not desensitize substantially over tens of minutes, which facilitates quantification of the light-driven antagonism. Under these conditions, a 100 ms flash inhibited the DAMGO-evoked current to 39.2±7.6% (n=13) of the peak, whereas a subsequent 2 s flash decreased the current to 0.7±0.1% of the peak (n=10, Figure 4A). To ensure that DAMGO did not saturate MORs, which would confound these interpretations, a saturating concentration of UK14304 (3 μM) was applied, followed by the selective α-adrenergic antagonist idazoxan (1 μM) at the end of each recording. α-adrenergic receptors and MORs activate the same pool of GIRK channels in LC neurons (Fiorillo and Williams, 1996; Williams and North, 1984). Whereas a saturating concentration of DAMGO activates a current that is about 120% that of the peak current induced by UK14304 (Fiorillo and Williams, 1996; Williams and North, 1984), 100 nM DAMGO resulted in a current that was about 70% as large, confirming that the 100 nM DAMGO evoked current was not maximal (Figure 4B).

### Kinetic analysis of opioid signaling deactivation

By pre-equilibrating tissue with the caged antagonist, photorelease results in nearly immediate delivery directly to receptor binding sites without long range diffusion through complex and dense neuropil. During experiments in which agonists were applied at several concentrations, we observed that the rate of light-driven current decay varied with agonist identity

as well as concentration for a given agonist. Time constants of current decay ranged from up to ~ 10 s at high concentrations, down to ~1 s with low concentrations of some agonists. Across experiments utilizing identical photolysis conditions, the kinetics of current decay should reflect four parameters: 1) agonist dissociation rate (koff), which is determined by its affinity (Kd) and association rate  $(k_{on})$ , as  $K_d = k_{off} / k_{on}$ ; 2) naloxone association rate, which is constant across conditions and much faster than the observed decay rates (Figure S4); 3) the rate of G-protein mediated biochemical signaling deactivation (k<sub>deact</sub>); and 4) the relative concentrations of agonist and photoreleased competitive antagonist, the latter of which is fixed by the experimental conditions (Figure 5). Studies employing rapid, application and washout of MOR agonists in acutely isolated LC neurons indicate that, with the exception of the high affinity agonist, etorphine, GIRK deactivation had a time constant of ranging from 0.7 to 1.4 s following washout (Ingram et al., 1997). Therefore the deactivation time constants observed on the scale of several seconds likely reflect competitive binding between agonist and antagonist. At high agonist concentrations, rebinding of agonist after NLX photorelease presumably slows the rate of current decay. However, with photorelease of sufficient NLX to prevent rebinding, the observed kinetics of deactivation should remain constant over a range of low agonist concentrations.

To determine if photolysis of CNV-NLX can resolve such ligand-receptor interaction kinetics in brain slices, we obtained a kinetic dose-response curve of the effects of NLX photorelease on currents evoked by the high-affinity peptide agonist DERM (Figure 6). The concentration of DERM was varied from 10 nM to 10  $\mu$ M and applied in the presence of 5  $\mu$ M CNV-NLX, which was photolyzed by a 5 s flash at the peak of the agonist-evoked outward current. Saturating MOR-mediated currents are, on average, 1.2 fold larger than  $\alpha$ -adrenergic receptor-mediated currents in the same cells (Fiorillo and Williams, 1996; Williams and North, 1984). By measuring the current evoked by each concentration of DERM in comparison to that evoked by UK14304 (3  $\mu$ M) in the same cells, DERM was found to activate MORs with an EC50 of ~10 nM, and saturating responses were reached at ~ 500 nM. As predicted, the currents evoked by low concentrations of DERM decayed faster after photolysis of CNV-NLX than those evoked by large concentrations (Figure 6A) and complete block was achieved at all but the highest

concentrations. Furthermore, the time constant of deactivation did not vary significantly at the three lowest concentrations tested (10, 30 & 100 nM; one-way ANOVA, p=0.26) and only slowed upon reaching near-saturating concentrations (500 nM – 10  $\mu$ M, Figure 6B). This likely reflects rebinding of agonist due to reversible, competitive receptor occupancy by the photoreleased NLX at high agonist concentrations, whereas at EC50 agonist concentrations and lower, the photoreleased NLX out competes agonist at the receptor thus eliminating rebinding. These results suggest that the rate of deactivation measured at low, sub-saturating concentrations of agonists can reflect the rate of agonist dissociation ( $k_{off}$ ). Furthermore, the post-photolysis current amplitude dose-response curve closely matches that obtained for the same agonist in the presence of 100 nM NLX (normalized to UK14304), suggesting that at least 100 nM NLX is released under these photolysis conditions (Figure S5).

DERM is known to have a high affinity for MOR and thus is expected to have a relatively slow k<sub>off</sub> in comparison to other opioid agonists. In order to compare the kinetics of agonist unbinding between agonists, deactivation kinetics of a panel of eleven agonists was examined at sub-saturating, near EC<sub>50</sub> concentrations that should avoid contributions from agonist rebinding. The opioid-evoked currents were below saturation by comparing them to the currents evoked by UK14304 (3 μM) in the same cells (Figure 7A, B). Varying deactivation rates were observed with no obvious correlation to current amplitude, suggesting that agonist rebinding was not occurring. To determine if deactivation kinetics were predictive of receptor affinity, the corresponding values for each agonist were plotted on a logarithmic scale (Figure 7C). Affinities are presented as pK<sub>i</sub> determined at rat MORs in radioligand competition experiments (Table S1 and associated discussion) (McPherson et al., 2010; Raynor et al., 1994; Raynor et al., 1995).

Although two agonists ( $\beta$ -endorphin and etorphine) exhibited dramatically slower deactivation kinetics, most agonists exhibited deactivation time constants ranging from 1-2 s, and none were faster than 1 s. Agonists with the highest affinities generally exhibited the slowest decay rates ( $\beta$ -endorphin, DERM, and etorphine), and the lowest affinity agonist codeine produced the fastest. Although DERM is reported as the highest affinity peptide ligand in the series,  $\beta$ -endorphin currents decayed >3 fold more slowly. At the other extreme, although

DAMGO, ME, and endomorphin-1 are reported to exhibit very similar affinities, the DAMGO- and ME-induced currents decayed  $\sim$ 3 fold more rapidly and were among the fastest to deactivate in the entire series. When considered according to compound class, the small molecules, which are most likely to exhibit a similar  $k_{on}$ , trend according to affinity, but with a slope of 0.3 rather than 1. For example, the decay rates measured for morphine and codeine differ by a factor of 1.6 rather than a factor of 10, as would be predicted for a constant  $k_{on}$ . The peptides showed even greater variability, consistent with their structural diversity and complexity, which could give rise to various forms of multi-step ligand-receptor interactions that determine  $k_{on}$  and  $k_{off}$ .

### **Discussion**

In this study we describe the development of a caged analog of the opioid antagonist NLX and demonstrate that it provides a means to study opioid receptor deactivation dynamics in brain tissue. This reagent lacks residual antagonist activity at concentrations up to at least 100 µM and photolysis in brain slices affords rapid and nearly complete inhibition of MOR mediated outward currents at both sub-saturating and near-saturating concentrations of agonist. Using full-field uncaging with a 405 nm LED as a light source and 5 µM CNV-NLX in the bath, light flashes ranging from one hundred milliseconds to 5 seconds in duration produced graded inhibition of the agonist-evoked currents, with the largest stimulus affording at least 100 nM NLX at the recorded neurons, corresponding to about 2% photolysis. In principle, the use of more powerful light sources would allow for even greater concentration jumps and for shorter light flashes to produce comparable levels of inhibition. At sub-saturating agonist concentrations, the inhibition produced by a single light flash was stable for several tens of seconds, presumably reflecting slow diffusion of NLX away from the illuminated area. At saturating agonist concentrations, the light-induced antagonism reversed more quickly, which is likely facilitated by competitive interactions between the ligands.

In LC neurons, MORs and  $\alpha$ -adrenergic receptors activate the same population of GIRK channels (Fiorillo and Williams, 1996; Williams and North, 1984). The convergence of  $G_{i/o}$ -coupled receptor signaling onto GIRK channels in LC neurons allowed us to examine the

selectivity of light driven antagonism with CNV-NLX for opioid receptors. Although photolysis of CNV-NLX during the application of the α-adrenoreceptor agonist UK14304 appeared to produce a small and transient inhibition in the form of an inward current, this effect persisted in the absence of CNV-NLX, indicating that it is an artifact of the intense light flash and is not due to non-specific actions of the photolysis products. The precise origin of this artifact is not clear from these data.

NLX is well established to function as a competitive antagonist at MORs and has also been characterized as an inverse-agonist under certain circumstances (Connor and Traynor, 2010). In our experiments, NLX was photoreleased in the presence of agonist, so the photoreleased NLX requires agonist dissociation before it can bind to receptors, according to the simplest models of ligand-receptor interactions that contain a single agonist binding site per functional receptor. In the absence of antagonist, the unbound agonist would simply rebind to sustain a constant level of receptor activation. However, if a sufficiently high concentration of antagonist is released to out-compete the agonist, agonist rebinding is overcome such that MORs cease to activate GIRK channels. Our race experiments (Fig. S4) involving simultaneous uncaging of agonist and antagonist indicate that NLX binding occurs much faster than GIRK activation, which occurs with a time constant of ~ 300 ms under these conditions. In acutely isolated LC neurons GIRK channels deactivated in the absence of bound agonist with a time constant of ~ 1 s (Ingram et al., 1997), so deactivation rates slower than 1 s were interpreted to reflect, at least in part, agonist dissociation from receptors. A similar time course of GIRK current deactivation following GABA-B receptor activation was also found in acutely isolated hippocampal pyramidal neurons (Sodickson and Bean, 1996). The deactivation of GIRK current is dependent on multiple mechanisms that can include agonist/receptor interactions, downstream mechanisms such as the activity of RGS proteins and the experimental conditions. The similarity in the rate of GIRK inactivation between the MOR and GABA-B receptors may be of interest given that the GABA-B receptor is an obligate dimer (Jones et al., 1998; Kaupmann et al., 1998; Pin et al., 2003; White et al., 1998). There is structural evidence that MOR can form dimers (Manglik et al., 2012), raising the possibility that NLX may initiate deactivation prior to agonist dissociation by

binding to unoccupied allosterically coupled orthosteric sites in multimeric receptors (Albizu et al., 2006; Springael et al., 2007), however the functional consequences of these associations remain to be determined. What is known is that single MORs isolated in nano discs have the ability to activate G-proteins, such that dimers of MOR are not strictly necessary at a functional level (Kuszak et al., 2009). Thus our interpretations of the measured deactivation rates are currently restricted to the simplest scenario of monomeric receptors and competitive interactions at a single orthosteric binding site.

Using the high affinity peptide agonist DERM and a fixed light-driven dose of CNV-NLX (5 µM, 5 s flash), we determined an agonist concentration window in which the measured current deactivation rates appear to reflect agonist unbinding from the receptor (10 – 100 nM), occurring with a time constant of ~ 4 s. Although the amplitude of the DERM-evoked currents varied according to agonist concentration, the current deactivation rates did not vary significantly. Apparently, the amount of NLX released by this light stimulus and dose (~100 nM, 100x the Kd) is sufficient to fully outcompete DERM and prevent agonist rebinding within this concentration range. At higher DERM concentrations, the rate of deactivation slowed significantly. The agonist concentration range over which photolysis of CNV-NLX will provide an accurate measure of dissociation kinetics without a contribution from agonist rebinding will vary with agonist affinity, concentration of CNV-NLX, and the quantity of light applied. In principle, kinetic studies can be conducted on large concentrations of high affinity agonists using large concentrations of CNV-NLX and/or high intensity light, but it may be more practical to use sub-saturating doses of agonist to conserve material and avoid phototoxicity.

Our kinetic data imply important considerations for the use of CNV-NLX to study GPCR deactivation kinetics using caged antagonists. None of the agonist-induced currents deactivated with time constants faster than 1 s, suggesting the presence of an instrumental limit. Thus, some ligands may dissociate with time constants faster than 1 s, but the rate-limiting step in signaling deactivation for such agonists is likely disassembly of the G protein-GIRK complex to enable channel closure. In contrast, several agonists, including DERM, yielded long time constants (3-10 s), suggesting that slow ligand dissociation is rate limiting in these cases, whereas those falling

between the extremes likely reflect mixed contributions from both events on approximately the same time scale. Regardless of which mechanism constitutes the rate limiting step, these deactivation time constants are intrinsic to each ligand-receptor complex and constitute a useful biophysical parameter for characterizing the receptor state and its associated signaling pathways. Thus in the context of studying mechanisms that regulate GPCR signaling, the choice of agonist determines which process is predominantly being measured: ligand dissociation or GIRK coupling (Figure 5). Monitoring current deactivation with slowly-dissociating agonists should reveal changes in receptor affinity that might occur during prolonged signaling (Birdsong et al., 2013) or during cross-talk between GPCRs. On the other hand, rapidly-dissociating agonists should reveal changes in the intrinsic rate of GIRK deactivation.

Because NLX is not selective for opioid receptor subtypes, CNV-NLX can be used to probe delta and kappa opioid receptors, in addition to MORs. Due to the necessity of the A-ring phenol for high affinity binding of opiate alkaloids in general, this caging strategy should work equally well with other opiates that contain an analogous phenol, regardless of whether they function as agonists or antagonists. When considered with our success in caging the opioid peptide agonists LE and Dynorphin A (1-8) at the N-terminal Tyrosine phenol, these findings open the door to caged opioid ligands that exhibit receptor subtype selectivity, of which many candidates are available as either peptides or small molecules.

We anticipate that CNV-NLX will also be useful to study actions of endogenously released opioid peptides in brain slices and *in vivo*. Although little is known about the quantities of endogenous opioids released *in vivo*, there is no evidence that saturating concentrations of agonist are reached. Since CNV-NLX effectively antagonizes high concentrations of exogenous agonists, it should be effective in blocking endogenous opioid actions *in vivo* and in *ex vivo* tissue with high spatiotemporal precision. Since the dissociation rate of NLX from MORs and diffusional clearance is slow, photoreleased NLX remains active for several minutes after a light flash. Together these points suggest that photolysis of CNV-NLX *in vivo* may facilitate the identification of the functions of endogenous opioid signaling in behaving animals.

# **Acknowledgements**

We thank Seksiri Arttamangkul, William Birdsong and Nicolas Tritsch for insightful discussions, S.D. Liberles, L.B. Buck and J.L. Whistler plasmids, and the ICCB-Longwood screening facility for access to plate readers.

# **Authorship Contributions**

Participated in research design: Banghart, Williams, and Sabatini

Conducted experiments: Banghart, Williams, and Shah

Contributed new reagents or analytic tools: Banghart and Lavis

Performed data analysis: Banghart and Williams

Wrote or contributed to the writing of the manuscript: Banghart, Williams and Sabatini

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# **Footnotes**

This work was supported by funds from the Howard Hughes Medical Institute to B.L.S and L.D.L; a grant from the National Institute of Mental Health to B.L.S. [MH085498]; grants from the National Institute on Drug Abuse to J.T.W [DA08163], and M.R.B [DA034648]; and postdoctoral fellowships from the Helen Hay Whitney Foundation and the Charles A. King Trust, N.A., Bank of America, Co-Trustee to M.R.B.

### **Figure Legends**

**Figure 1.** Design and synthesis of CNV-NLX, a caged opioid antagonist. A, Chemical structures of CYLE (1), morphine (2), and codeine (3). B, Two-step synthesis of CNV-NLX (6) from NLX (4).

**Figure 2.** CNV-NLX is inactive at the mu opioid receptor expressed in HEK293 cells. A, Doseresponse curves of NLX (black circles) and CNV-NLX (blue squares) in the presence of 1  $\mu$ M LE (n=12 wells). B, Dose response curves of LE alone (black squares), and in the presence of 5  $\mu$ M NLX (upward red triangles) and CNV-NLX (downward green triangles) (n=6 wells).

**Figure 3.** Light-driven antagonism of opioid receptors by CNV-NLX. A, Photolysis inhibits currents evoked by high concentrations of the MOR agonists DAMGO, DERM and methadone, but causes only an artifactual response at α-adrenergic receptors in the presence of UK14304. B, Summary plot comparing currents evoked by MOR agonists and the α-adrenergic agonist UK14304, before and after CNV-NLX photolysis (n=4-8 cells per agonist). Two concentrations were pooled for both ME and methadone. Purple triangles indicate 5 s flashes of 405 nm light.

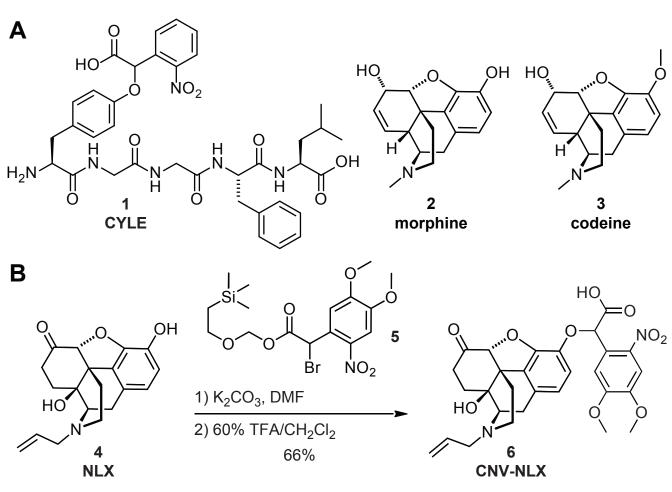
**Figure 4:** Analog photo-control of NLX release. A, Varying the duration of the light flash produces graded inhibition of the outward current evoked by bath perfusion of a subsaturating concentration of DAMGO. B, Summary plot comparing the current amplitudes produced by DAMGO (100 nM) before and after each uncaging stimulus to that produced by a saturating concentration of UK14304 (3  $\mu$ M, n=4 cells). Because DAMGO (100 nM) produced smaller peak currents than UK14304 (3  $\mu$ M), the opioid stimulus is confirmed to be sub-saturating.

**Figure 5.** Schematic illustrating the molecular events that underlie opioid signaling deactivation in response to antagonist uncaging in the presence of agonist. The photo-released competitive antagonist can bind only after agonist dissociation, which occurs with a rate constant of  $k_{\text{off}}$ . In the agonist-free state, the G-protein subunits that support GIRK channel opening dissociate from the activated channels, allowing channel closure with a rate constant of  $k_{\text{deact}}$ .

**Figure 6.** Low agonist concentrations reveal DERM ligand dissociation kinetics. A, (*left*), Superimposition of the currents evoked by 10 nM and 10 μM DERM. (*right*) Amplitude-normalized currents reveal slower deactivation at saturating agonist concentration. B, Summary plot of the DERM current, normalized to the current induced by UK14304 (3 μM, black squares), and tau decay (open circles) as a function of DERM concentration (*n*=4-8 cells per condition). The dashed line indicates the average of the time constants measured at 10, 30 and 100 nM, which did not vary significantly.

**Figure 7.** Light-induced deactivation of currents produced by various MOR agonists at subsaturating concentrations. A, Superimposition of the normalized currents evoked by subsaturating concentrations of various small molecule (*left*) and peptide (*right*) agonists. B, Summary plot of the normalized currents (white bars) and corresponding light-induced deactivation rates (black bars) at the indicated sub-saturating concentrations. Opioid agonist-induced peak currents are normalized to the peak currents produced by UK14303 (3 μM) in the same cells (*n*=4-8 cells per condition). C, Summary plot of the deactivation time constant vs. agonist affinity according to compound class (*n*=4-8 cells per condition).

# Figure 1



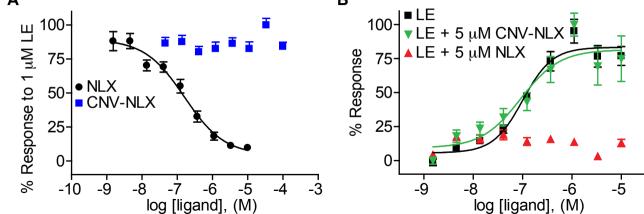


Figure 3

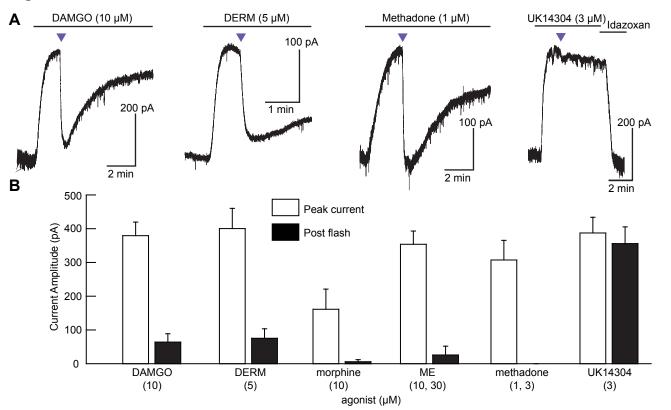


Figure 4

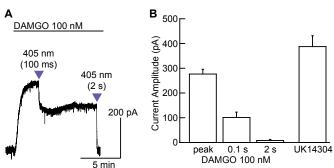


Figure 5  $O_2N$ 405 nm ′

Figure 6 405 nm DERM 10 µM normalized 200 p/ DERM 10 n 10 s В 1.5 current (UK 14304 normalized 12 current ∘ tau 10 tau decay (s) 1.0 8 6 0.5 4 2 0 0 10 100 1000 10000 [DERM] nM

Figure 7

