

Septal Glucagon-Like Peptide I Receptor Expression Determines Suppression of Cocaine-Induced Behavior

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Glucagon-like peptide I (GLP-I) and its receptor GLP-IR are a key component of the satiety signaling system, and long-acting GLP-I analogs have been approved for the treatment of type-2 diabetes mellitus. Previous reports demonstrate that GLP-I regulates glucose homeostasis alongside the rewarding effects of food. Both palatable food and illicit drugs activate brain reward circuitries, and pharmacological studies suggest that central nervous system GLP-I signaling holds potential for the treatment of addiction. However, the role of endogenous GLP-I in the attenuation of reward-oriented behavior, and the essential domains of the mesolimbic system mediating these beneficial effects, are largely unknown. We hypothesized that the central regions of highest *Glp-1r* gene activity are essential in mediating responses to drugs of abuse. Here, we show that *Glp-1r*-deficient (*Glp-1r*^{-/-}) mice have greatly augmented cocaine-induced locomotor responses and enhanced conditional place preference compared with wild-type (*Glp-1r*^{+/+}) controls. Employing mRNA *in situ* hybridization we located peak *Glp-1r* mRNA expression in GABAergic neurons of the dorsal lateral septum, an anatomical site with a crucial function in reward perception. Whole-cell patch-clamp recordings of dorsal lateral septum neurons revealed that genetic *Glp-1r* ablation leads to increased excitability of these cells. Viral vector-mediated *Glp-1r* gene delivery to the dorsal lateral septum of *Glp-1r*^{-/-} animals reduced cocaine-induced locomotion and conditional place preference to wild-type levels. This site-specific genetic complementation did not affect the anxiogenic phenotype observed in *Glp-1r*^{-/-} controls. These data reveal a novel role of GLP-IR in dorsal lateral septum function driving behavioral responses to cocaine.

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INTRODUCTION

The shift from controlled to compulsive drug use leading to addiction is poorly understood but seems to involve pathways originally evolved for natural rewards like food (Volkow and Wise, 2005). Recent preclinical studies suggested that glucagon-like peptide 1 (GLP-1) modulates the rewarding effects of both food (Alhadeff *et al*, 2012; Dickson *et al*, 2012; Dossat *et al*, 2011) and addictive drugs (Egecioglu *et al*, 2013a, b, 2013c; Erreger *et al*, 2012; Graham *et al*, 2013). This was supported by the observation that pretreatment with the synthetic GLP-1R agonist Exendin-4 (Ex4) attenuates the behavioral effects of drugs of abuse including ethanol (Egecioglu *et al*, 2013c; Shirazi *et al*, 2013), nicotine (Egecioglu *et al*, 2013a), amphetamine, and cocaine

(Egecioglu *et al*, 2013a, b; Erreger *et al*, 2012; Graham *et al*, 2013). Therefore, GLP-1R agonists have been proposed as treatment options for drug addiction. Most studies employed the systemic administration of GLP-1R ligands, obviating the dissection of peripheral *versus* central GLP-1R effects. Interestingly, GLP-1R activation in the ventral tegmental area (VTA) mediates the rewarding effects of alcohol (Shirazi *et al*, 2013) and palatable food (Mietlicki-Baase *et al*, 2013). In addition, systemic Ex4 decreases the release of dopamine in the nucleus accumbens (NAc) (Egecioglu *et al*, 2013a, b). It remains to be determined whether these effects are because of a direct activation of GLP-1R in the NAc and VTA or rather a result of modulations on a circuit level.

In the central nervous system (CNS), GLP-1 is produced by neurons in the nucleus of the solitary tract (NTS) that project to various regions, including sites that regulate stress responses, as well as reward and motivation (Llewellyn-Smith *et al*, 2011; Rinaman, 2010). The immunohistochemical detection of GLP-1R expression in the brain is unknown as specific anti-GLP-1R antibodies are not available (Pyke and Knudsen, 2013), yet ligand binding assays suggest the presence of the receptor in the amygdala and the mesolimbic

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system, including the VTA, NAc, and lateral septum (LS), providing an anatomical correlate for the reported beneficial effects of pharmacological GLP-1R stimulation (Göke *et al*, 1995).

The LS is highly interconnected with the mesolimbic system (Olds and Milner, 1954; Sheehan, 2004) primarily through γ -aminobutyric acid (GABA)ergic projections (Gallagher *et al*, 1995; Risold and Swanson, 1997a). LS neurons form a strong collateral network (Gallagher *et al*, 1995; Sheehan, 2004) that not only causes recurrent inhibition within the LS, but also tightly controls the activity of extraseptal target areas (Sheehan, 2004). The LS regulates responses to natural rewards (Noda *et al*, 1997) and is involved in stress-induced relapse to drug abuse (Highfield *et al*, 2000). Hence, the LS is a hub that integrates information arriving from central key regulators and mediates appropriate behavioral responses (Sheehan, 2004). A recent study identified a novel circuit, linking context and reward, that relies on neurons of the dorsal LS (dLS) (Luo *et al*, 2011). This circuit starts with hippocampal glutamatergic projections that stimulate GABAergic neurons in the dLS. These dLS neurons project to the VTA, leading to disinhibition of dopaminergic neurons within the VTA, and a subsequent increase of dopamine levels in the NAc (Luo *et al*, 2011; Mazei-Robison *et al*, 2011).

The identification of mechanisms that regulate LS activity is therefore important for understanding the circuits of the mesolimbic reward system as well as the neurobiology of drug-induced neuroplasticity and drug reinforcement. Here, we used a genetic mouse model with a targeted null mutation of the *Glp-1r* locus (Scrocchi *et al*, 1996), enabling the determination of behavioral responses to cocaine administration that were dependent upon GLP-1-mediated LS neuromodulation.

MATERIALS AND METHODS

Animals

Experiments were carried out in accordance with the UNSW Australia Animal Care and Ethics Committee. Heterozygous *Glp-1r*^{+/-} breeding pairs were used to produce age-matched male *Glp-1r*^{-/-} mice (Scrocchi *et al*, 1996) on C57BL/6J background and *Glp-1r*^{+/+} littermates. PCR genotyping was performed as described previously (Kim *et al*, 2013). Animals were individually housed in a temperature- and humidity-controlled room with a 12 h light/dark cycle and had access to food and water *ad libitum*.

Adeno-Associated Virus (AAV) Constructs

The rat *Glp-1r* open reading frame was cloned upstream and in-frame with the coding region of the enhanced green fluorescent protein (GFP) reporter in an AAV expression cassette containing the 1.1-kb chicken β -actin promoter (CBA), the woodchuck hepatitis virus posttranscriptional regulatory element (WPRE), and the bovine growth hormone polyadenylation sequence (bGHpA) flanked by AAV2 inverted terminal repeats (pAAV-GLP-1R-GFP). The same backbone carrying no *Glp-1r* cDNA was used as a control (pAAV-GFP). In addition, a nontagged GLP-1R construct (During *et al*, 2003) was used as a control for experiments in human embryonic kidney (HEK293) cells.

Immunoblotting

HEK293 cells were transfected with pAAV-GLP-1R-GFP, pAAV-GLP-1R, or pAAV-GFP, using the standard calcium phosphate (CaPO₄) precipitation method. At 48 h after transfection, the cells were exposed to Ex4 (10 nM) for 0–50 min before harvesting in solubilization buffer (30 mM Tris-Cl, pH 7.4, and 5 mM MgCl₂ containing 4 mg of CHAPS (Sigma) and 20% glycerol). Then, 10 μ g of total protein was resolved by 12% SDS-PAGE and electroblotted onto nitrocellulose membrane. The membrane was incubated with primary antibodies against phosphorylated extracellular signal-regulated kinases (mouse anti-pERK1/2; 1 : 2500; Cell Signaling, Danvers, MA) and the GFP-tag (rabbit anti-GFP; 1 : 2000 (von Jonquieres *et al*, 2013)). The membrane was reprobed using a rabbit anti-ERK1/2 antibody (1 : 2500; Cell Signaling). Secondary antibodies were horseradish peroxidase conjugated as required (1 : 5000; Dianova, Hamburg, Germany). Immunoreactivity was detected using the enhanced chemiluminescence system (Bio-Rad, Gladesville, NSW, Australia) and digitalized (GelDoc, Bio-Rad).

AAV Vector Generation and Administration

Packaging of neurotropic AAVrh10 vectors was performed as described previously (Klugmann *et al*, 2005). Briefly, following triple transfection of HEK293 cells, AAV vectors were purified using iodixanol (OptiPrep, Sigma-Aldrich, Taufkirchen, Germany) gradient ultracentrifugation (Zolotukhin *et al*, 2002). Titers were determined by quantitative real-time PCR of vector genomes using primers for WPRE. Then, 1 μ l of either AAV-GLP-1R-GFP or AAV-GFP (8×10^{11} vg/ml) was injected bilaterally into the dLS (+0.5 mm AP, \pm 0.3 mm ML, –2.6 mm DV from bregma). Vectors were delivered to 4 week-old mice at a rate of 150 nl/min using a microprocessor-controlled mini-pump (World Precision Instruments, Sarasota, FA) with 34G beveled needles (World Precision Instruments) in a stereotaxic frame (Kopf Instruments, Tujunga, CA).

Immunohistochemistry

Brains of all animals were assessed for transgene expression at the end of behavioral testing. Immunostaining of brain sections was performed as previously described (Klugmann *et al*, 2005). Briefly, mice were transcardially perfused with 10% buffered neutral formalin (Sigma-Aldrich), removed, and post-fixed for 2 h in the same fixative before cryoprotection in 30% sucrose/PBS. Free-floating cryosections (40 μ m) were rinsed with PBS–Triton X-100 (0.2%), blocked in immunobuffer (4% horse serum in PBS, pH 7.4, 0.4% Triton X-100), followed by incubation with rabbit anti-GFP (von Jonquieres *et al*, 2013) (1 : 1000). Sections were then incubated with donkey anti-rabbit Alexa 488 antibody (1 : 1000; Invitrogen, Karlsruhe, Germany) and counterstained with 4',6-diamidino-2-phenylindole (DAPI). Sections were mounted and coverslipped with Mowiol (Sigma-Aldrich). Fluorescence was visualized using a confocal microscope (Zeiss Z1 Axio Examiner NLO710; Carl Zeiss, North Ryde, NSW, Australia).

In Situ Hybridization (ISH)

ISH was performed as previously described (Marsicano and Lutz, 1999). Both digoxigenin (DIG)-labeled and fluorescein isothiocyanate (FITC)-labeled riboprobes were used as described elsewhere (Yang *et al*, 1999). The cDNAs encoding glutamic acid decarboxylase (GAD65) and mouse GLP-1R (GenBank accession number: NM_021332) in the pBlue-script plasmid backbone, including a *Glp-1r* sense probe, were used for *in vitro* transcription. Hybridization was carried out overnight at 55 °C in 100 µl of buffer containing DIG-labeled (300 ng/ml) *Glp-1r* riboprobe for single ISH or, in case of double ISH, additional FITC-labeled *Gad65* riboprobe (300 ng/ml). Sections were counterstained with DAPI or toluidin blue and coverslipped in Mowiol or DPX (Sigma-Aldrich). Fluorescent sections were visualized using a confocal microscope; DIG-labeled sections were visualized using an Olympus BX51 microscope. Semiquantitative analysis, in the coronal plane cut through bregma, was performed by determining the number of *Glp-1r* mRNA-expressing cells among the total population of dLS neurons identified by soma size or *Gad65* mRNA-expressing neurons. Samples were collected from three *Glp-1r*^{-/-} and *Glp-1r*^{+/-} animals (three sections per animal, spaced 80 µm). Digitized images were processed with Adobe Photoshop CS5 or ZEN lite (Zeiss) where appropriate.

Electrophysiology

Coronal brain slices (300 µm) were prepared from mice (4–8 weeks) using a VT1200 vibratome (Leica) according to standard techniques (Power and Sah, 2007). Briefly, mice were anesthetized with isoflurane (5%) and decapitated, and the brains removed and sectioned in oxygenated (95% O₂, 5% CO₂) ice-cold, modified artificial cerebrospinal fluid (ACSF) containing the following (in mM): 125 NaCl, 3 KCl, 1.25 NaH₂PO₄, 25 NaHCO₃, 6 MgCl₂, 1 CaCl₂, and 10 glucose. Slices were incubated at 34 °C for 30 min and thereafter maintained at room temperature in ACSF solution (in mM): 125 NaCl, 3 KCl, 1.25 NaH₂PO₄, 25 NaHCO₃, 1 MgCl₂, 2 CaCl₂, and 25 glucose (equilibrated with 95% O₂, 5% CO₂). Whole-cell patch-clamp recordings were made from visually identified neurons (Zeiss Axio Examiner D1) in slices continuously superfused with ACSF (1–2 ml/min) heated to 30 ± 1 °C using an in-line heater (TC-324B Warner Instrument Corporation). Patch pipettes (2–4 MΩ) were filled with internal solution (in mM): 135 KMeSO₄, 7 NaCl, 10 HEPES, 2 Mg₂ATP, and 0.3 Na₃GTP, pH 7.3 with KOH (osmolarity, 280–300 mOsm/kg). Electrophysiological recordings utilized an Axopatch 200B amplifier (Molecular Devices), filtered at 2–5 kHz, digitized at 20 kHz with a Digidata1440 (Molecular Devices) interface, controlled using PClamp 10.2 software (Molecular Devices). Current was injected to adjust the membrane potential to –65 mV, and the voltage response to somatic current injections (600 ms, –50 to +150 pA, 10 pA increments) was used to measure the neuron's AP firing pattern. Action potential latency was calculated from the first action potential evoked at the threshold current injection. Neurons with resting potentials > –55 mV were excluded from the data set. Voltage-clamp was employed to dissect dLS membrane conductances, in particular A-type K⁺-currents (Supplementary Data).

Behavioral Testing

All animals were adult (2–4 months) and individually housed 2 weeks before behavioral testing to prevent abnormal behavior caused by fighting or suppression by the alpha male. Experience in our lab shows that reward-related behavior does not change in C57bl6/J mice between 2 and 6 months. All animals were subjected to the following battery of tests. The elevated plus maze (EPM) was employed to assess inherent anxiety-related behavior. The maze consisted of two opposite open arms and two opposite enclosed arms, interconnected by a central platform (center 6 × 6 cm²; open arms, 35 × 6 cm²; closed arms, 35 × 6 × 15 cm³; elevation 1 m; illumination 40 lux). Mice were placed on the maze, facing a closed arm and were allowed to freely explore the EPM for 5 min. Mouse behavior was analyzed using ANY-maze tracking software (Stoelting, Wood Dale, IL). The ratio of time spent in open arms was used to estimate open arm aversion (reflecting anxiety). The same cohorts of animals were subjected to two noncontingent models of substance abuse (Kalivas, 2009). Briefly, clear Plexiglas boxes (41 × 41 × 40 cm; Photobeam Activity System: MedAssociates, St Albans, VT) were used as open field boxes for recording the locomotor sensitization to cocaine (30 min sessions per test day). After an initial day with saline injections (baseline activity), animals received cocaine (10 mg/kg, IP) for 5 consecutive days (days 1–5; with day 5 representing the development of sensitization). After 7 consecutive drug- and test-free days, all mice received a priming injection of cocaine (day 12; representing the expression of sensitization). The last day of sensitization was followed by 14 test-free days before the same animals were subjected to the conditioned place preference (CPP) test. CPP was evaluated using a modified unbiased protocol as described previously (Bilbao *et al*, 2008). Briefly, CPP was performed in a custom-made box with two equally sized compartments (25 × 25 × 40 cm) interconnected by a rectangular corridor. The compartments were differentiated by the pattern of the walls and texture of the floor. Mice received one session of preconditioning (15 min), followed by 4 days of conditioning. During the conditioning phase, mice were treated with eight alternating injections of saline (morning) and cocaine (10 mg/kg; IP; afternoon) and then confined to the corresponding compartment immediately after the injection for 30 min. For the postconditioning test (performed at noon), mice were allowed to explore the apparatus without any treatment for 15 min. The CPP score represents the difference between the time spent in the cocaine compartment before and following conditioning.

Data Analysis and Statistics

Current clamp data were analyzed off-line using custom routines written in LabVIEW (National Instruments, Austin, TX). Data are presented as means ± SEM and were analyzed as appropriate using unpaired, two-tailed *t*-test, or two-way repeated measures ANOVA followed by the Bonferroni *post hoc* test to evaluate statistical significance. In all cases, *P* < 0.05 was considered statistically significant.

RESULTS

Abundant Expression of Endogenous *Glp-1r* in the LS

Glp-1r-expressing cells were found in the amygdala as well as the hypothalamus (Figure 1a). In order to gain an overview

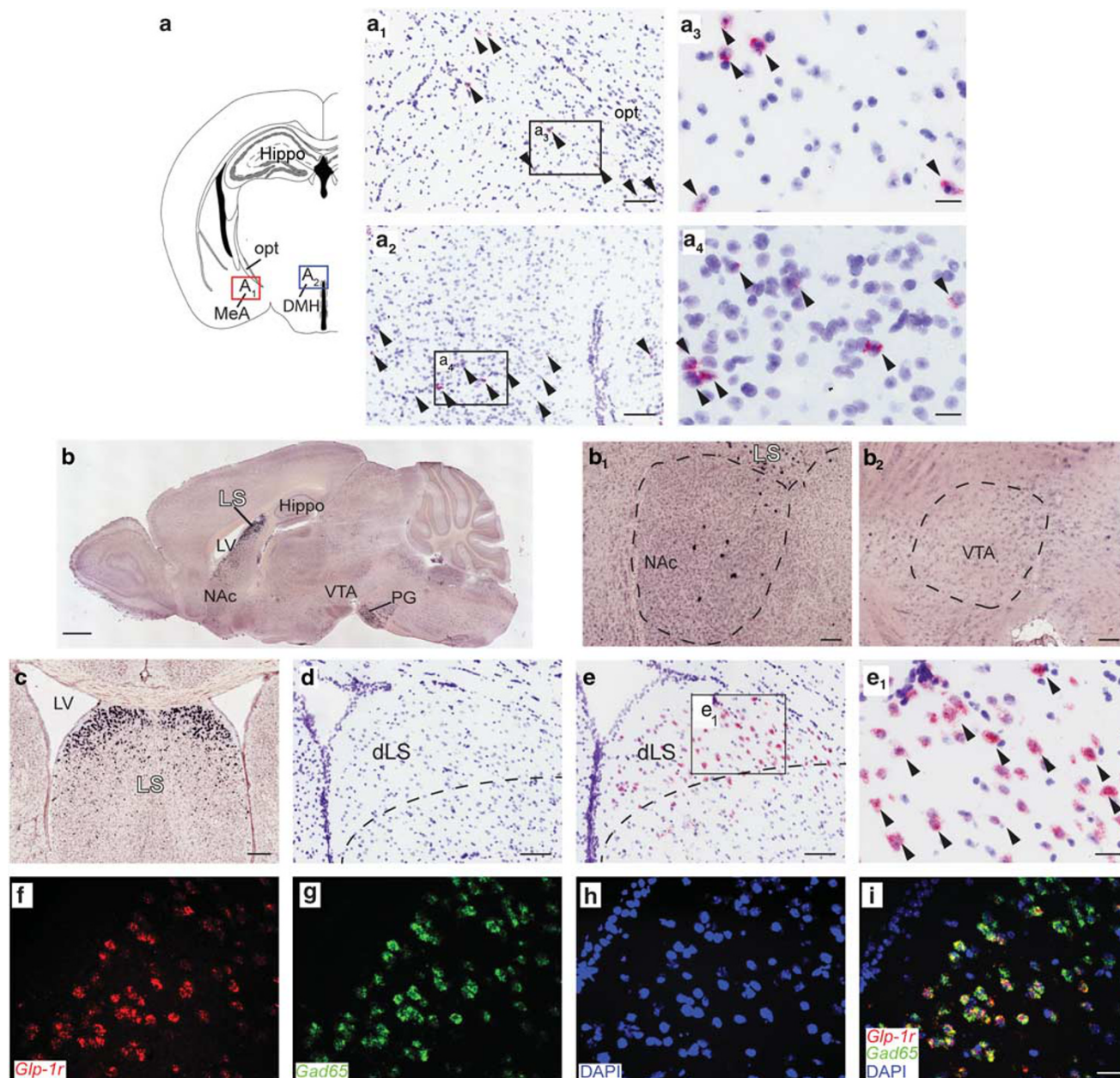


Figure 1 *Glp-1r* mRNA is enriched in the lateral septum. (a) Illustration of a coronal mouse brain section. (a₁, a₂) *Glp-1r* mRNA-expressing cells (purple) in the medial amygdaloid nucleus (red box in (a)) or the hypothalamus (blue box in (a)) labeled by arrowheads. (a₃, a₄) High magnification of individual *Glp-1r* mRNA-positive cells. (b) Overview of *Glp-1r* mRNA expression (black staining) domains in the sagittal plane (Allen Brain Atlas ID 73606497_125). (b₁, b₂) Close-ups of the NAc and the VTA showing little *Glp-1r* expression. (c) In the coronal plane, highest *Glp-1r* expression (silver impregnation) was identified in the LS (Allen Brain Atlas ID 74511737_345). (d) The *Glp-1r* sense probe produced no signal. (e) Abundant *Glp-1r* mRNA expression (pink) was confirmed in the dLS using a specific DIG-labeled riboprobe. (e₁) Higher magnification of the dLS indicated by the box in (e) depicting strong cellular signal detection of *Glp-1r* mRNA. (f) Fluorescent detection of *Glp-1r* (red) and (g) *Gad65* (green) mRNA in the dLS. (h) Nuclear counterstaining with DAPI (blue). (i) Merged picture reveals that almost all *Glp-1r*-positive septal neurons are GABAergic. Bars: (a₁, a₂, d, e) 75 μ m; (a₃, a₄, e, f) 10 μ m; (b) 1 mm; (b₁, b₂) 100 μ m; (c) 200 μ m. dLS, dorsal lateral septum; DMH, dorsomedial hypothalamic nucleus; Hippo, hippocampus; LV, lateral ventricle; MeA, medial amygdaloid nucleus; NAc, nucleus accumbens; opt, optic tract; PG, pontine gray; VTA, ventral tegmental area.

of *Glp-1r* gene activity, we utilized expression data publicly available from the Allen Brain Atlas database (Lein *et al*, 2007). High-density *Glp-1r* mRNA labeling was detected not only in the pontine gray but also in the LS (Figure 1b). In contrast, only few or no *Glp-1r*-positive cells were found in the NAc (Figure 1b₁) or VTA (Figure 1b₂). Given that LS

neurons project to both areas, we further elucidated *Glp-1r* mRNA expression in the LS. Abundant *Glp-1r* expression levels in the LS were corroborated assessing an independent Allen Brain Atlas data set (Figure 1c). We then performed ISH to identify *Glp-1r* expression in the LS in more detail. Hybridization with the sense riboprobe did not result in

signal detection, confirming absence of nonspecific signals (Figure 1d). The antisense riboprobe confirmed strong *Glp-1r* mRNA expression in the dLS as shown in a representative image (Figure 1e). Semiquantitation of these results revealed that >95% of dLS neurons express *Glp-1r* mRNA. To identify the phenotype of *Glp-1r*-positive dLS neurons, we performed double fluorescent ISH using probes for *Glp-1r* and *Gad65*, the enzyme responsible for GABA synthesis. Virtually all *Glp-1r*-positive cells expressed *Gad65*, indicating that these cells are inhibitory GABAergic neurons (Figures 1f–i). No signal was detected with the antisense probe using brain sections from GLP-1R-deficient mice (not shown).

Ablation of GLP-1R Signaling Alters Intrinsic Excitability of Septal Neurons

Our anatomical data suggested that septal GLP-1Rs may play an important role in the mesolimbic reward system. We hypothesized that this is caused by GLP-1R-dependent changes in dLS neuronal excitability and therefore analyzed the firing properties of *Glp-1r*^{+/+} and *Glp-1r*^{-/-} neurons employing brain slice whole-cell patch-clamp. The dLS neurons from *Glp-1r*^{+/+} and *Glp-1r*^{-/-} did not differ in their passive membrane properties (resting potential, input resistance, membrane time constant) or their AP waveform (threshold, amplitude, width), their fast AHP amplitude, or the minimum current required to elicit an AP (Supplementary Table S1). However, there were differences in the neuronal firing properties. In response to a threshold current injection, the onset of the AP was delayed in *Glp-1r*^{-/-} neurons compared with *Glp-1r*^{+/+} controls (Figure 2a). Increasing the injected current revealed that septal *Glp-1r*^{-/-} neurons were more excitable, firing more APs than *Glp-1r*^{+/+} neurons to twice threshold current injections (Figure 2d) and to current injections >120 pA (Figure 2e and f). The AP onset difference suggested that fast activating voltage-dependent K⁺ currents might be altered in *Glp-1r*^{-/-} neurons. Dissecting these currents with voltage-clamp protocols in combination with the pharmacological blockage isolated prominent A-type K⁺ currents that were retained with *Glp-1r* deletion (Supplementary Figure S1).

GLP-1R in the dLS Controls Cocaine-Induced Behaviors

Our anatomical and electrophysiological data indicated that GLP-1R contributes to dLS function and therefore to behavioral outputs modulated by the dLS (Sheehan, 2004). We hypothesized that assessments of anxiety-related and drug-induced behaviors would reveal deficits in *Glp-1r*^{-/-} mice that were tied to the dLS and hence could be rescued by genetic complementation of GLP-1R in the dLS.

To achieve precise anatomical accuracy for expression of transgenic GLP-1R *in vivo*, we employed the AAV platform to induce robust long-term transgene expression following intracranial delivery (Klugmann et al, 2005). The AAV expression construct included a GFP-epitope tag at the GLP-1R C-terminus for immunodetection of the recombinant receptor (Figure 3a). Biological functionality of the tagged receptor was confirmed in heterologous HEK293 cells by monitoring the activation of ERK1/2, a known downstream event of GLP-1R signaling (During et al, 2003). Levels

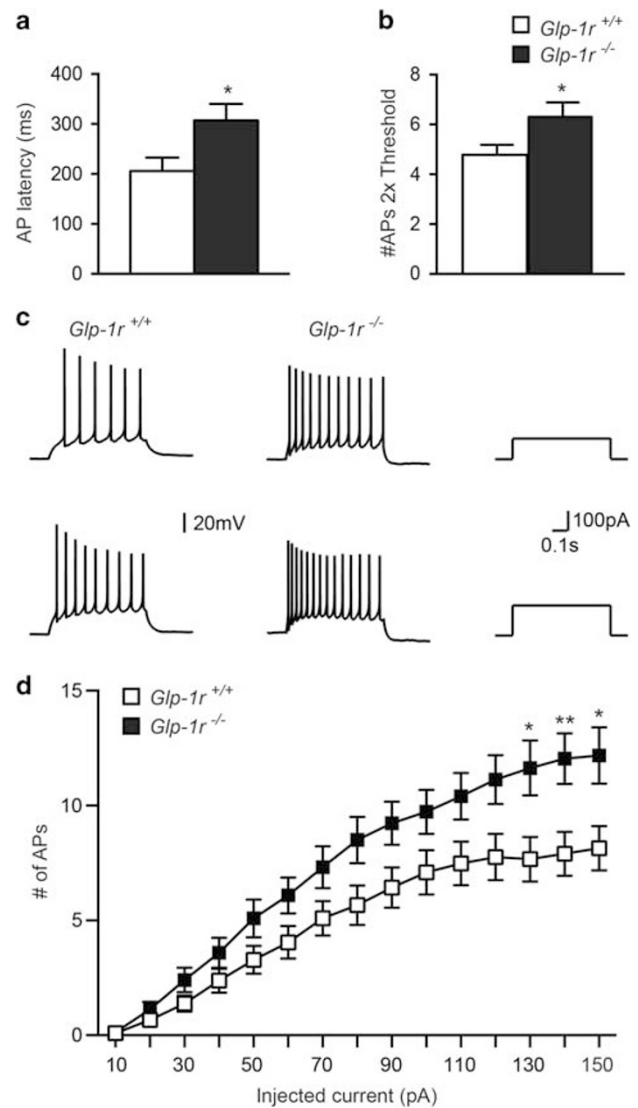


Figure 2 Increased firing of *Glp-1r*^{-/-} septal neurons. (a) The time to fire an AP is delayed in *Glp-1r*^{-/-} (total of 23 cells from 10 animals) compared with *Glp-1r*^{+/+} (total of 21 cells from 11 animals) neurons ($P=0.02$). (b) The number of APs at twice threshold was elevated in *Glp-1r*^{-/-} compared with *Glp-1r*^{+/+} controls ($P=0.04$). (c) Representative responses of *Glp-1r*^{+/+} (left) and *Glp-1r*^{-/-} (right) neuron, respectively, to 600 ms depolarizing current steps at 100 and 150 pA. (d) Responses (number of APs evoked by 600 ms stimulus) of *Glp-1r*^{+/+} and *Glp-1r*^{-/-} cells across a range of step current injections from 10 to 150 pA revealed an increased activity of *Glp-1r*^{-/-} neurons (two-way repeated measures ANOVA \times genotype effect $F(1, 43) = 6.48$, $P=0.01$). * $P<0.05$, ** $P<0.01$.

of phosphorylated ERK1/2 peaked at 10 min and declined after 20 min following administration of Ex4 (Figure 3b). A doublet at 80 and 100 kD detected by immunoblot (Figure 3b) indicated posttranslational modifications. Immunocytochemical detection of recombinant GLP-1R revealed cell surface localization (Figure 3c). These data suggest that the presence of the GFP tag does not interfere with receptor signaling, posttranslational processing, or trafficking.

We then packaged AAV-GLP-1R-GFP or AAV-GFP for bilateral delivery into the mouse dLS. Immunohistochemical

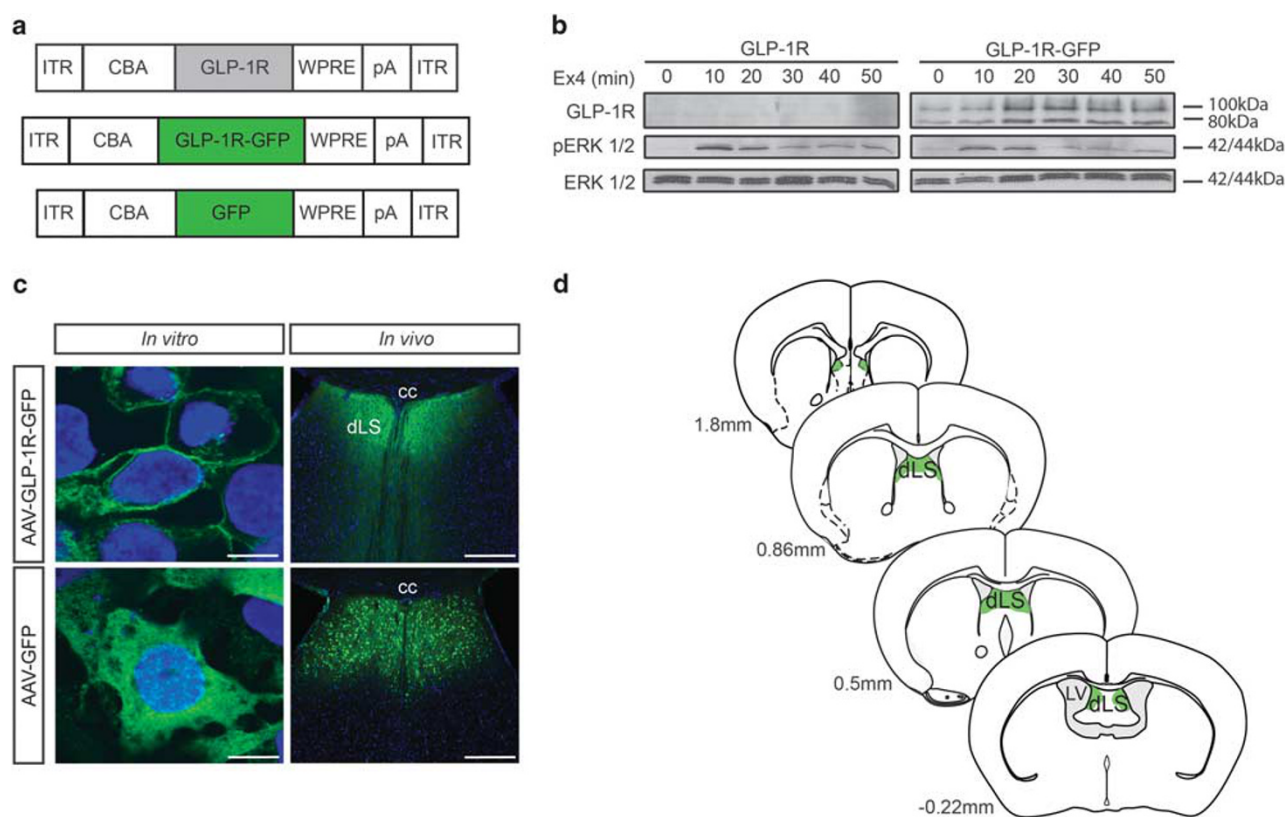


Figure 3 AAV-mediated *Glp-1r* gene delivery to the dLS. (a) AAV expression cassettes. ITR, inverted terminal repeat; CBA, chicken β -actin promoter; pA, polyadenylation signal; WPRE, woodchuck hepatitis virus posttranscriptional regulatory element. (b) Biological activity of GLP-1R-GFP was confirmed by expression in heterologous HEK293 cells and exposure to Ex4 (10 nM) for the indicated durations (0–50 min). The dynamics of ERK phosphorylation were similar following activation of the tagged GLP-1R or the native receptor. Total ERK served as loading control. Note that the GFP antibody exclusively recognized the tagged recombinant receptor. (c) Recombinant GLP-1R-GFP traffics to the cell surface in HEK293 cells and in the mouse LS (top row), whereas GFP controls show cytosolic expression (bottom row). Anti-GFP immunofluorescence (green); DAPI (blue) nuclear counterstain. Bars: 10 μ m. (d) Schematic illustration of vector spread assessed by AAV-transduced neuronal somata; cc, corpus callosum; dLS, dorsal lateral septum; LV, lateral ventricle.

assessment of transduction was performed 3 weeks following vector delivery, when AAV-mediated transgene expression is known to reach maximum and stable levels (Klugmann *et al*, 2005), to confirm accurate anatomical targeting; the subcellular expression patterns matched our observations *in vitro* (Figure 3c). As anatomical confinement of the rescue experiment is essential, we investigated the possibility of off-targeting in the lateral ventricles. Transduced ependymal cells, glia lining the walls of the ventricles, are an inevitable consequence of intraventricular AAV delivery. We critically examined all 15 AAV animals ($n = 9$ for AAV-GLP-1R; $n = 6$ for AAV-GFP) for this feature and observed GFP-positive ependymal cells in only one case, an AAV-GFP brain, as depicted in Figure 3c.

The vector spread extended 2 mm in the rostrocaudal dimension (Supplementary Figure S2) and is illustrated in Figure 3d. AAVrh10 can be transported retrogradely along axonal projections (Cearley and Wolfe, 2007); except a small number of transduced CA3 pyramidal neurons in the dorsal hippocampus (not shown), we did not detect GFP-positive cell bodies in extraseptal regions.

For behavioral assessment we tested AAV-GFP-treated and naive *Glp-1r*^{-/-} or *Glp-1r*^{+/+} mice. Data of the corresponding AAV-GFP and naive mice were pooled as they were not statistically different (not shown). A separate cohort

entailed *Glp-1r*^{-/-} mice bilaterally injected with AAV-GLP-1R-GFP into the dLS. We employed the EPM test to determine levels of inherent anxiety. *Glp-1r*^{-/-} mice and *Glp-1r*^{-/-} mice injected with AAV-GLP-1R-GFP spent less time in the center of the maze, showed fewer open arm entries, and spent less time in the open arms compared with *Glp-1r*^{+/+} controls (Figure 4a). All groups displayed comparable locomotor activity (Supplementary Figure S3A). These results suggest that the loss of GLP-1R in brain regions outside the dLS mediate anxiety-related behavior.

The same cohorts were then subjected to the locomotor sensitization test as described (see Materials and Methods). *Glp-1r*^{-/-} mice, but not *Glp-1r*^{-/-} mice injected with AAV-GLP-1R-GFP, showed an augmented acute response to cocaine (Figure 4b; day 1). Increased locomotor activity was detected in *Glp-1r*^{+/+} controls and *Glp-1r*^{-/-} mice between days 1 and 5, indicating intact development of sensitization. Although development was absent in *Glp-1r*^{-/-} mice injected with AAV-GLP-1R-GFP, this group was not statistically different to *Glp-1r*^{+/+} controls. Following a drug-free period of 7 days, the animals received a cocaine challenge, and *Glp-1r*^{+/+} controls and *Glp-1r*^{-/-} mice, but not *Glp-1r*^{-/-} mice injected with AAV-GLP-1R-GFP, showed expression of sensitization (day 12). Baseline locomotion in the open field box was not different among groups

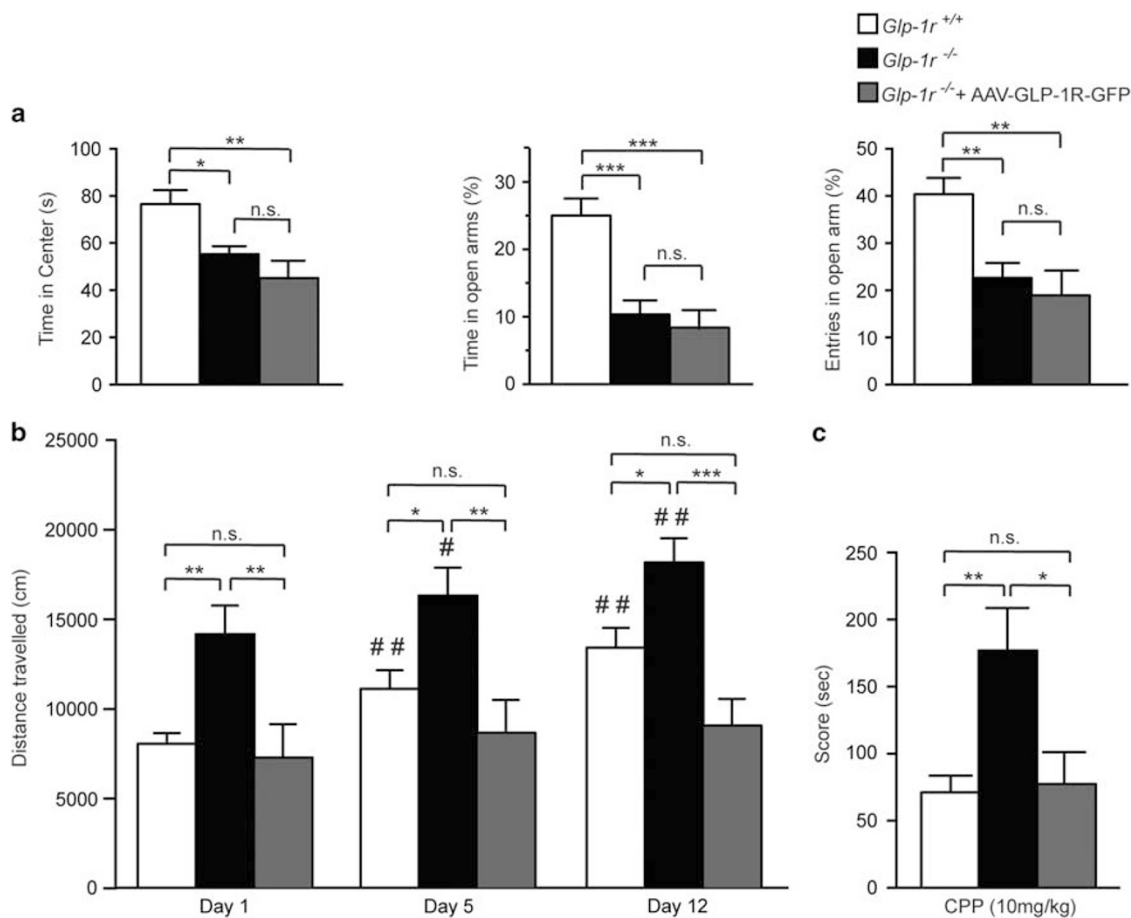


Figure 4 Genetic complementation of GLP-IR expression in the dLS rescues cocaine-induced, but not anxiety-related, behaviors in *Glp-1r*-deficient mice. (a) *Glp-1r*^{-/-} mice and *Glp-1r*^{-/-} mice injected with AAV-GLP-1R-GFP spent less time in the center of the EPM (left panel, one-way ANOVA $F=8.049$; $P=0.0012$); *Glp-1r*^{+/+} controls ($n=17$) compared with *Glp-1r*^{-/-} mice ($n=15$; $P<0.05$), or *Glp-1r*^{+/+} controls compared with *Glp-1r*^{-/-} + AAV-GLP-1R-GFP mice ($n=9$; $P<0.01$). Middle panel: *Glp-1r*^{-/-} mice and *Glp-1r*^{-/-} mice injected with AAV-GLP-1R-GFP spent less time in the open arms (one-way ANOVA $F=14.34$; $P<0.0001$) compared with *Glp-1r*^{+/+} controls ($P<0.0001$) and entered the open arms less frequently (right panel; one-way ANOVA $F=9.65$; $P<0.001$; *post hoc* test $P<0.01$). (b) Altered locomotor response to cocaine for all groups over the course of the paradigm (two-way repeated measures ANOVA genotype effect $F(2, 41)=9.771$, $P=0.0003$). There was no difference between genetically complemented *Glp-1r*^{-/-} mice ($n=9$) and *Glp-1r*^{+/+} controls ($n=17$) following acute cocaine exposure (day 1; $P>0.9999$). Intact development and expression of cocaine sensitization (two-way repeated measures ANOVA cocaine effect $F(2, 82)=21.00$, $P<0.0001$) was present in *Glp-1r*^{+/+} and *Glp-1r*^{-/-} mice ($n=15$) but absent in *Glp-1r*^{-/-} + AAV-GLP-1R-GFP mice (day 1–day 5, $P=0.5141$ and day 1–day 12, $P=0.2919$). Note that there was no difference between *Glp-1r*^{+/+} controls and *Glp-1r*^{-/-} + AAV-GLP-1R-GFP on any given day. # *P* compared with day 1; * *P* compared with all groups on a given day. (c) Genetically complemented *Glp-1r*^{-/-} mice displayed a normal CPP phenotype (one-way ANOVA $F=7.037$, $P=0.0023$). The CPP score represents the difference between the time spent in the cocaine compartment before and following conditioning. * or # $P<0.05$, ** or ## $P<0.01$, *** $P<0.001$.

(Supplementary Figure S3B). All groups showed a similar magnitude of change upon comparison of the difference between days 5 and 1 or between days 12 and 1 (Supplementary Figure S4A). Therefore, the sensitization process *per se* was not enhanced in *Glp-1r*^{-/-} mice compared with *Glp-1r*^{+/+} controls, probably because of the augmented initial sensitivity to cocaine and a ceiling effect in *Glp-1r*^{-/-} mice. Sensitization as well as initial responsiveness to psychomotor stimulants can be exhibited by stereotypy that is usually associated with a drop in locomotion. However, no alterations between groups were observed when comparing stereotypic time over the course of the experiment (Supplementary Figure S4B).

The rewarding effects of cocaine were investigated in the CPP paradigm. *Glp-1r*^{-/-} mice exhibited enhanced CPP compared with *Glp-1r*^{+/+} controls following conditioning

using cocaine doses of 10 mg/kg (Figure 4c) or 15 mg/kg (Supplementary Figure S5), suggesting augmented cocaine-seeking behavior. This response was restored in *Glp-1r*^{-/-} mice treated with AAV-GLP-1R-GFP (Figure 4c). To rule out that the observed CPP of *Glp-1r*^{-/-} mice is caused by an enhanced conditioned locomotor effect (Huston *et al*, 2013) we analyzed the locomotor activity during CPP expression and detected no difference between groups (Supplementary Figure S6).

Glp-1r mRNA expression was increased by fourfold in the dLS of AAV-GLP-1R-GFP-treated *Glp-1r*^{-/-} mutants compared with AAV-GFP-treated *Glp-1r*^{+/+} mice using qRT-PCR analysis of brain slice punches (Supplementary Figure S7). Although the corresponding GLP-1R protein quantities could not be determined, it was shown previously that even substantial AAV-mediated transgene expression at

mRNA levels translates to physiological protein amounts (Blume *et al*, 2013).

DISCUSSION

This study provides clear evidence that GLP-1R-positive dLS GABAergic neurons are critical for the mediation of drug-induced behavior. Several reports have shown that feeding-related peptides, such as leptin or ghrelin, influence the mesolimbic system (Hommel *et al*, 2006; Skibicka *et al*, 2011). The role of GLP-1 and its analogs in this context has only recently been investigated by pharmacological studies, supporting the concept that GLP-1R signaling alters behaviors induced by drugs of abuse (Egecioglu *et al*, 2013a, b, 2013c; Erreger *et al*, 2012; Graham *et al*, 2013; Shirazi *et al*, 2013). The preclinical potential of GLP-1-based compounds for the treatment of dependence has been broadly appreciated (Skibicka, 2013), and the translational relevance of this research is emerging (van Bloemendaal *et al*, 2014).

Acute injections of GLP-1 or Ex4 into the VTA or NAc reduce reward-driven behavior, an effect that was attributed to GLP-1R activation in these brain areas (Dickson *et al*, 2012; Dossat *et al*, 2011). The direct action of GLP-1 appears conceivable in light of classical ligand binding data reporting the presence of GLP-1R in these areas (Göke *et al*, 1995). Moreover, GLP-1-producing NTS neurons project into key areas of the mesolimbic system including the VTA and NAc (Merchenthaler *et al*, 1999; Rinaman, 2010). However, only mRNA ISH data inform specifically on the neuroanatomical sites of *Glp-1r*-producing neurons; whereas sparse *Glp-1r* mRNA levels were reported in the aforementioned parts of the mesolimbic system (Merchenthaler *et al*, 1999), this study identified most abundant central *Glp-1r* mRNA expression in the dLS (Figure 1b and c), the functional relevance of which has not been appreciated yet. Broadly, the LS is connected with brain regions implicated in regulating mood and reward (Risold and Swanson, 1997b) and strongly influences motivated behaviors (Olds and Milner, 1954; Prado-Alcala *et al*, 1984).

Here, we utilized genetic GLP-1R models on the C57BL/6J background and showed that the complete ablation of GLP-1R caused an anxiogenic phenotype that was not restored by dLS directed AAV-*Glp-1r* gene delivery, suggesting that extra septal GLP-1Rs (ie, amygdala) mediate anxiety-related behavior. Inconclusive results have previously been reported for EPM test performance of GLP-1R-deficient mice on a CD1 background (MacLusky, 2000). Considering strain differences, that are known to play a critical role in mediating behavioral responses, CD1 mice showed an decrease in anxiety-related behavior compared with the C57BL/6J background (Michalikova *et al*, 2010).

Glp-1r^{-/-} mice showed augmented cocaine-induced acute locomotor response and increased CPP, indicating enhanced reinforcement and stimulatory effects of cocaine in these animals. This phenotype was corrected by the genetic rescue confined to the dLS, indicating that GLP-1R in neurons from that brain region is essential for motivated behavior.

The peripheral delivery of Ex4 in mice and rats reduces cocaine and amphetamine-induced hyperlocomotion (Egecioglu *et al*, 2013b; Erreger *et al*, 2012; Graham *et al*,

2013); yet these pharmacological studies utilized protocols limited to monitor acute psychostimulant responses. In contrast, we employed an established sensitization paradigm testing both acute and chronic responses (Bilbao *et al*, 2008), revealing that complete GLP-1R ablation enhanced cocaine-induced locomotor activity throughout the test. As such, our findings are in line with the previously reported ability of Ex4 to block the expression of nicotine-induced locomotor sensitization (Egecioglu *et al*, 2013a). The persistent nature of the sensitized motor response has been proposed to reflect neurochemical alteration similar to the enduring changes associated with addiction, indicating potentially shared neuronal substrates (Kalivas, 2009; Robinson and Berridge, 1993).

The LS is crucial for performance in the CPP paradigm (Sartor and Aston-Jones, 2012) and our anatomical and behavioral data strongly suggest that dLS GLP-1Rs contribute to this phenomenon. Although GLP-1R-deficiency is implicated in diminished hippocampus-dependent learning (During *et al*, 2003), and the CPP paradigm relies on contextual memory (Cunningham *et al*, 2006), we detected increased cocaine-induced conditioning of *Glp-1r*^{-/-} animals providing evidence that the contextual impairment does not confound the interpretation of the CPP performance. Given that human addiction involves drug self-administration, future experiments could employ contingent models to examine the role of septal GLP-1Rs.

The neurochemical phenotype of GLP-1R-expressing neurons is broadly unknown for most brain regions. We provide evidence that GLP-1R-positive cells in the dLS are predominantly GABAergic and it is known that LS projections terminate in the VTA and NAc (Onteniente *et al*, 1987; Risold and Swanson, 1997b; Swanson, 1982). Attenuated dopamine release in the NAc was reported following systemic Ex4 administration (Egecioglu *et al*, 2013a, c). Moreover, intra-VTA GLP-1R activation is mediated in part by nonglutamatergic mechanisms (Mietlicki-Baase *et al*, 2013). These data raise the possibility that the effects mediated by direct activation of GLP-1R in the mesolimbic system can be attributed to modulations on a circuit level. We propose that GLP-1R neurons in the dLS are essential for mesolimbic function. The underlying mechanisms are unclear but may involve downstream effects of GLP-1R signal transduction such as the modulation of cAMP levels (Drucker *et al*, 1987), activation of Ca²⁺-channels (Acuna-Goycolea and van den Pol, 2004), or regulation of transcription factors (Klugmann *et al*, 2006).

In summary, we show that the dLS has the most prominent GLP-1 receptor expression in the brain. GLP-1R-null mice exhibited increased excitability in GABAergic dorsal lateral septum neurons. These mutants had strongly augmented behavioral responses to cocaine when tested in two widely used noncontingent models of drug abuse. In a rescue experiment, we genetically complemented GLP-1R expression exclusively in the dLS of GLP-1R-deficient mice and observed restoration of cocaine-induced behavior to wild-type levels. This dLS-directed rescue of the increased cocaine-induced behavioral measures in GLP-1R-null mice indicates that GLP-1 transmission in this brain region is a key suppressor of addictive drug-induced behavior. These findings will shift attention to this brain region and GLP-1 transmission as a novel target for therapeutic intervention in drug addiction.

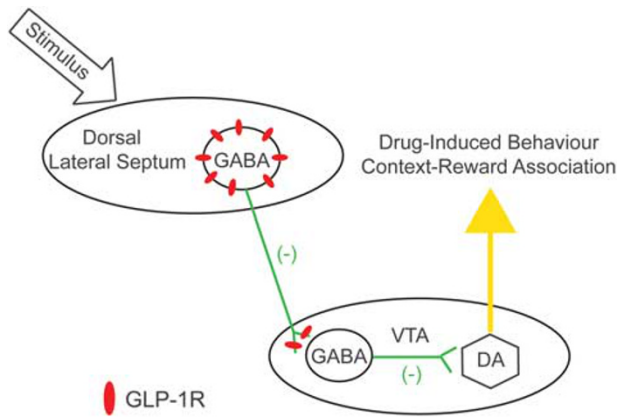


Figure 5 Septal GLP-IR regulates motivated behavior. GLP-IR-expressing GABAergic septal neurons modulate cellular activity following stimulation. GLP-IRs are present in both somata and terminals of these neurons (Göke et al, 1995; Shirazi et al, 2013) that form inhibitory synapses with GABAergic interneurons in the VTA (Luo et al, 2011). Their inhibition, in turn, leads to activation of dopaminergic VTA neurons via disinhibition. Hence, this dLS–VTA pathway controls drug-induced behaviors such as locomotor sensitization and drug seeking. DA, dopamine. Adapted from Luo et al (2011).

Our working model (Figure 5) integrates our findings into a recently proposed CA3–dLS–VTA pathway that connects dLS activity to dopamine release from VTA neurons and, in turn, modulates motivated behavior (Luo et al, 2011). Most GABAergic dLS neurons coexpress *Glp-1r* mRNA (Figure 1). LS cells get activated by a range of stimuli (Sheehan, 2004), leading to context-dependent (Luo et al, 2011) and drug-induced adaptations of motivated behavior. GLP-1R appears to play an essential role for these adaptations, suggested by our electrophysiological (Figure 2) and genetic rescue data (Figure 4b and c).

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