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Altered neurotransmission in the mesolimbic reward system of *Girk1*^{-/-} mice

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Abstract

Mice lacking the *Girk2* subunit of G protein-gated inwardly-rectifying K⁺ (*Girk*) channels exhibit dopamine-dependent hyperactivity and elevated responses to drugs that stimulate dopamine neurotransmission. The dopamine-dependent phenotypes seen in *Girk2*^{-/-} mice could reflect increased intrinsic excitability of or diminished inhibitory feedback to midbrain dopamine neurons, or secondary adaptations triggered by *Girk2* ablation. We addressed these possibilities by evaluating *Girk*^{-/-} mice in behavioral, electrophysiological, and cell biological assays centered on the mesolimbic dopamine system. Despite differences in the contribution of *Girk1* and *Girk2* subunits to *Girk* signaling in midbrain dopamine neurons, *Girk1*^{-/-} and *Girk2*^{-/-} mice exhibited comparable baseline hyperactivities and enhanced responses to cocaine. *Girk* ablation also correlated with altered afferent input to dopamine neurons in the ventral tegmental area. Dopamine neurons from *Girk1*^{-/-} and *Girk2*^{-/-} mice exhibited elevated glutamatergic neurotransmission, paralleled by increased synaptic levels of AMPA glutamate receptors. In addition, synapse density, AMPA receptor levels, and glutamatergic neurotransmission were elevated in medium spiny neurons of the nucleus accumbens from *Girk1*^{-/-} and *Girk2*^{-/-} mice. We conclude that dopamine-dependent phenotypes in *Girk2*^{-/-} mice are not solely attributable to a loss of *Girk* signaling in dopamine neurons, and likely involve secondary adaptations facilitating glutamatergic signaling in the mesolimbic reward system.

Keywords

Cocaine; dopamine; glutamate; plasticity; knockout

INTRODUCTION

The mesolimbic dopamine (DA) system consists of projections from the ventral tegmental area (VTA) to the nucleus accumbens (NAcc) and mediates in part the motor-stimulatory and reinforcing effects of drugs of abuse (Nestler 2004; Wise 2004; Bjorklund and Dunnett 2007). Though drugs of abuse have distinct molecular targets, they share the ability to increase DA levels in the NAcc (Luscher and Ungless 2006). Cocaine, for example, elevates

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NAcc DA levels by blocking the re-uptake of released DA. The physiological relevance of this effect and location is underscored by the impact of destruction of the dopaminergic projection from the VTA to NAcc, which includes hypoactivity and insensitivity to the motor-stimulatory effects of cocaine (Koob et al. 1981).

In addition to potentiating DA signaling within the NAcc, cocaine also triggers “long-loop” and auto-inhibitory feedback to the VTA mediated by GABA_B (GABA_BR) and D₂ DA (D₂R) receptors, respectively. G protein-gated inwardly-rectifying K⁺ (Girk/K_{IR3}) channels are expressed in VTA DA neurons and mediate the postsynaptic inhibitory effect of GABA_BR and D₂R activation (Beckstead et al. 2004; Cruz et al. 2004; Labouebe et al. 2007). The prototypical neuronal Girk channel is a heterotetramer formed by Girk1 and Girk2 subunits (Karschin et al. 1996; Liao et al. 1996; Luscher et al. 1997; Koyrakh et al. 2005). Midbrain DA neurons of the VTA and substantia nigra (SN), however, do not express *Girk1* (Karschin et al. 1996; Inanobe et al. 1999; Cruz et al. 2004; Labouebe et al. 2007). Thus, genetic ablation of *Girk2* correlates with blunted GABA_BR- and D₂R-dependent postsynaptic inhibition in VTA and SN DA neurons, whereas loss of *Girk1* has no effect (Beckstead et al. 2004; Cruz et al. 2004; Koyrakh et al. 2005; Labouebe et al. 2007).

Girk2^{-/-} mice exhibit a basal hyperactivity that was normalized by the D₁ DA receptor (D₁R) antagonist SCH 23390 (Blednov et al. 2001; Blednov et al. 2002), indicating that *Girk2*^{-/-} mice exhibit elevated DA signaling. *Girk2*^{-/-} mice also exhibit enhanced responses to the acute motor-stimulatory effects of cocaine and the D₁R partial agonist SKF 38393 (Blednov et al. 2002; Morgan et al. 2003). One interpretation of these data is that *Girk2* ablation diminishes inhibitory feedback to VTA DA neurons and/or triggers elevated intrinsic excitability of VTA DA neurons, leading to elevated DA release in the NAcc and striatum. Secondary adaptations, perhaps linked to alterations in excitatory signaling, might also explain the DA-related phenotypes reported in *Girk2*^{-/-} mice. Indeed, such adaptations are triggered by drugs of abuse and stress, and can promote persistent adaptations in the NAcc that are implicated in behavioral sensitization (Kauer and Malenka 2007; Engblom et al. 2008; Thomas et al. 2008; Zweifel et al. 2008; Heshmati 2009).

Here, we sought insight into the DA-related phenotypes reported in *Girk2*^{-/-} mice. We reasoned that if these phenotypes reflect the loss of Girk signaling in VTA DA neurons, they would be observed in *Girk2*^{-/-} but not *Girk1*^{-/-} mice. We applied a combination of behavioral, electrophysiological, and cell biological approaches to examine the impact of *Girk1* and *Girk2* ablation on neurotransmission within the mesolimbic reward pathway. Our findings indicate that loss of Girk signaling in midbrain DA neurons is not the primary cause of the DA-related phenotypes reported in *Girk2*^{-/-} mice, and suggest instead that secondary adaptations that facilitate glutamatergic neurotransmission may play a role.

MATERIALS AND METHODS

Drugs

Cocaine hydrochloride, baclofen, tetrodotoxin, picrotoxin, lidocaine hydrochloride and kynurenic acid were purchased from Sigma (St. Louis, MO). CGP54626 was purchased from Tocris (Ellisville, MO). rTertiapinQ was purchased from Alomone Labs, Inc. (Jerusalem, Israel).

Animals

All animal use was reviewed and approved by the Institutional Animal Care and Use Committee of the University of Minnesota. Efforts were made to minimize the pain and discomfort of the animals throughout the study. Adult mice were housed on a 12 h light/dark cycle, with food and water available *ad libitum*. The generation of *Girk*^{-/-} was described

previously (Signorini et al. 1997; Bettahi et al. 2002; Torrecilla et al. 2002). The *Girk* null mutations were backcrossed through >12 rounds against the C57BL/6 strain prior to experimentation.

Locomotor activity

Baseline and cocaine-induced motor activity were assessed in wild-type and *Girk*^{-/-} mice (4-10 wk) using automated open field environments, as described (Pravetoni and Wickman 2008). Baseline activity data were obtained during 60-min daily sessions over a consecutive 3-d period. On day 3, subjects were given injections of saline (i.p.) and placed in the open field for 60-min. Subsequently, subjects were randomly assigned to groups that received saline or one of four cocaine doses (3, 15, 30, or 60 mg/kg, i.p.). Total distance traveled after the injection was monitored for 60 min.

Electrophysiology

Coronal slices containing the VTA (200-300 μ m) and sagittal slices containing the NAcc (240 μ m) were prepared from drug-naïve wild-type and *Girk*^{-/-} mice (4-6 weeks), and allowed to recover in oxygenated ACSF at room temperature for >1 h. Slices were transferred to recording chambers and perfused with oxygenated ACSF (+/- drugs) at a flow rate of approximately 2-2.5 mL/min during experiments. Currents, resistances, and potentials were measured using Multiclamp 700A amplifiers and pCLAMP v.9 software (Molecular Devices; Foster City CA) and stored on hard disk. All measured and command potentials factored in junction potentials predicted using JPCalc software (Molecular Devices).

Studies involving VTA DA neurons were performed at 32-34°C, with borosilicate (3-5 M Ω) electrodes containing (in mM): 140 K-gluconate, 2 MgCl₂, 1.1 EGTA, 5 HEPES, 2 Na-ATP, 0.3 Na₃GTP, and 5 phosphocreatine, pH 7.4. For sIPSC measurements, KCl replaced K-gluconate in the pipette solution. Upon achieving whole-cell access, the current response to a 1-s voltage-ramp (-60 to -120 mV) was measured to assess I_h presence and size. Following a 5-min stabilization period, spontaneous action potential frequency and duration were measured in whole-cell current clamp mode (I=0). Cells exhibiting relatively large capacitance (>50 pF), I_h current (>100 pA at -120 mV), moderate and stable spontaneous activity (1-5 Hz), and long action potential durations (>2.5 s) were considered to be putative DA neurons. Indeed, these properties correlate well with expression of tyrosine hydroxylase and *Pitx3*, selective markers of DA neurons (Grace and Onn 1989; Johnson and North 1992; Momiyama et al. 1996; Cameron et al. 1997; Klink et al. 2001; Neuhoff et al. 2002; Mathon et al. 2005; Labouebe et al. 2007).

Changes in whole-cell holding current evoked by baclofen were measured at a holding potential (V_{hold}) of -60 mV. The holding current, input resistance (R_{in}), and series resistance (R_A) values were monitored throughout these experiments by tracking responses to periodic (0.2 Hz) voltage steps (-5 mV, 800 ms). Only experiments with stable (<20% variation) and low series resistances (<20 M Ω) were included in the final data set. The GABA_BR antagonist CGP54626 (2 μ M) was used to verify the receptor-dependence of baclofen-evoked responses. sIPSCs were measured over a 2-3 min period at V_{hold} = -70 mV, in the presence of kynurenic acid (2 mM). sEPSCs were measured over 2-3 min period at V_{hold} = -80 mV, in the presence of picrotoxin (100 μ M). All data were low-pass filtered at 2 kHz, digitized at 10 kHz, and analyzed using pCLAMP v.9 software (Molecular Devices).

Studies involving NAcc medium spiny neurons were conducted at room temperature, and recording electrodes contained the K-gluconate pipette solution described above or an

alternative solution lacking K^+ (in mM): 117 cesium gluconate, 2.8 NaCl, 20 HEPES, 0.4 EGTA, 5 tetraethylammonium-Cl, 2 MgATP, and 0.3 MgGTP, pH 7.2–7.4. mEPSC amplitudes and frequencies measured in wild-type NAcc neurons did not differ for these two pipette solutions and as such, data were pooled. Medium spiny neurons were identified by their morphology and hyperpolarized resting membrane potential (-75 to -85 mV). mEPSCs were measured at $V_{\text{hold}} = -80$ mV in the presence of picrotoxin (100 μM) and either lidocaine hydrochloride (0.6–0.8 mM) or TTX (1 μM). mEPSC data were filtered at 2 kHz, digitized at 5–10 kHz, and collected and analyzed using custom software (Igor Pro; Wavemetrics, Lake Oswego, OR).

AMPA receptor subunit quantification

Synaptic levels of GluR1 and GluR2/3 subunits were measured in the VTA and NAcc from wild-type and *Girk*^{-/-} mice (8–12 wk) via post-embedding immunoelectron microscopy. Tissue was prepared as described (Lujan et al. 1996); sections containing of the VTA and NAcc were then cut at 500 μm incubated in a 1 M sucrose/PBS solution overnight, slammed onto copper blocks cooled in liquid nitrogen, and embedded in Lowicryl HM20 (TAAB Laboratories; Aldermaston, UK) after freeze substitution with methanol. Ultrathin sections (80 nm) from three Lowicryl-embedded blocks were incubated for 45 min on coated nickel grids with drops of blocking solution containing: 0.05 M TBS, 0.9% NaCl, 0.03% Triton X-100, and 2% albumin. Grids were then incubated at room temperature overnight in blocking solution containing rabbit polyclonal antibodies (10 $\mu\text{g}/\text{ml}$ each) against GluR1 (AB1504, Millipore; Billerica, MA) or GluR2/3 (AB1506, Millipore). For VTA experiments, a mouse monoclonal antibody directed against TH (mAb 6D7, EMD Biosciences; San Diego, CA) was used to identify DA neurons. After washing in TBS, grids were incubated for 2 hr in drops of goat anti-rabbit IgG conjugated to 10 nm colloidal gold particles (BioCell International; Cardiff, UK) for experiments involving the NAcc, or a mixture of goat anti-rabbit IgG conjugated to 10 nm colloidal gold particles and goat anti-mouse IgG conjugated to 20 nm colloidal gold particles for experiments involving the VTA; in all cases, secondary antibodies were diluted 1:80 in a 0.05 M TBS solution containing 2% normal human serum and 0.5% polyethylene glycol. Grids were then washed in TBS for 30 min and counterstained for electron microscopy with saturated aqueous uranyl acetate and lead citrate.

Ultrastructural analyses were performed with a Jeol 1010 electron microscope (Peabody, MA). Electron photomicrographs were captured with a CCD camera (MegaView III Soft Imaging System; Munster, Germany). Digitized electron images were modified for brightness and contrast with Adobe Photoshop version 7.0. Quantification of GluR1 and GluR2/3 immunolabeling was performed in ultrathin sections obtained from 3 separate panels of wild-type, *Girk1*^{-/-}, and *Girk2*^{-/-} mice. Areas in the VTA and Nacc were chosen at random and images were captured at a magnification of 50,000X. In the VTA, only those synapses consisting of an axon terminal apposed to a PSD located on a dendritic shaft labelled for TH were included in the analysis. In the Nacc, only those synapses consisting of an axon terminal apposed to a PSD present on a dendritic spine were included in the analysis. In both locations, the length of the PSD was measured and the number of immunoparticles per PSD was tabulated. There was no impact of *Girk* ablation on PSD length (not shown).

Synapse quantification

Mice (8–10 weeks) were anesthetized and perfused with a solution containing 2% paraformaldehyde and 1.5% glutaraldehyde, as described (Marker et al. 2005). Coronal sections (60 μm) were cut through the level of the NAcc using a vibrating microtome. After several washes in PB, sections were post-fixed with 1% osmium tetroxide in PB and block-

stained with 1% uranyl acetate in distilled water. Sections were then dehydrated in ascending series of ethanol (to 100%) followed by propylene oxide and flat-embedded on glass slides in Durcupan (Sigma; St. Louis, MO). For quantitative analysis of the density of spines/synapses in the NAcc, sampling fields were chosen by using the systematic random sampling method (Geinisman et al. 1996). After collection of semi-thin sections and determination of sample sites, serial ultrathin sections (80 nm thick) of the neuropil containing the sampling fields were collected from three different parts of the NAcc of three different animals. Sections were placed on slot grids and stained with Reynolds' lead citrate.

Ultrastructural analyses were performed on a Jeol-1010 electron microscope from sections at a magnification of 40,000X. Spine/synapse density was estimated by using the unbiased physical dissector method. In brief, data were collected from pairs of serial sections (dissectors). Synapses were identified on the reference section by the presence of: 1) a postsynaptic density (PSD), 2) synaptic vesicles at the presynaptic terminal, and 3) opposing membranes between the pre- and the postsynaptic terminals. Only PSDs found on dendritic spines and only one PSD per spine were analyzed. An unbiased counting frame was superimposed over each of the two micrographs, and the PSD was used as a counting unit. Synapses were labeled on the reference section micrograph if their PSD profiles were located either entirely or partly within the frame and did not intersect the forbidden edges of the frame and its extensions. Finally, only synapses that had a PSD profile in the reference, but not in the look-up section, were counted. At least seven neuropil fields (each 163.65 μm^2) were photographed from each animal, corresponding to a total section area of 3.437 μm^2 . Synapse density was calculated as the number of synapses per counting divided by the product of area of the counting frame and the height of the dissector.

Data analysis

Data are presented throughout as the mean \pm SEM. Statistical analyses were performed using GraphPad Prism (GraphPad Software; La Jolla, CA). Quantal events were analyzed using Minianalysis software (Synaptosoft, Decatur, GA). The impact of genotype on VTA DA neuron characteristics, baclofen-induced current amplitude, EPSC and IPSC frequency and amplitude, synaptic level of AMPA subunits, synapse density, and cocaine-induced motor activity were assessed using one-way ANOVA, followed by Tukey's Multiple Comparison post hoc test when appropriate. For all statistical comparisons, *P* values of less than 0.05 were considered to reflect significant differences between groups.

RESULTS

We first measured basal activity levels and the motor-stimulatory effect of systemic cocaine (0-60 mg/kg) in wild-type and *Girk*^{-/-} mice in a standard open-field environment (Fig. 1). *Girk1*^{-/-} and *Girk2*^{-/-} mice were similarly hyperactive during their initial and subsequent exposures to the novel environment (Fig. 1A). Cocaine triggered dose-dependent increases in motor activity in all groups (Fig. 1B). While none of the genotypes responded significantly to 3 mg/kg cocaine, *Girk1*^{-/-} and *Girk2*^{-/-} mice showed comparable and enhanced responses to 15 mg/kg cocaine. Maximal responses did not differ across genotypes, though they were observed at 15 mg/kg for *Girk2*^{-/-} mice and at 30 mg/kg for the other three groups. Responses of *Girk2*^{-/-} mice to 30 mg/kg were notably, though not significantly, smaller than those seen at the 15 mg/kg. Similarly, 60 mg/kg cocaine evoked sub-maximal responses in wild-type, *Girk1*^{-/-}, and *Girk3*^{-/-} mice.

Given the differential expression patterns of *Girk1* and *Girk2* in midbrain DA neurons, the comparable profiles for *Girk1*^{-/-} and *Girk2*^{-/-} mice with respect to basal and cocaine-induced motor activity argue that the loss of *Girk* signaling in these neurons is not the primary cause. Thus, we asked next whether *Girk* ablation altered other aspects of midbrain

DA neuron physiology. We first examined the impact of *Girk* subunit ablation on the spontaneous activity and afferent input to putative DA neurons in acutely-isolated slices of the VTA. VTA DA neurons were identified using morphological and electrophysiological criteria (**Materials and Methods**) that we demonstrated previously correlated well with expression of *Pitx3* (Labouebe et al. 2007), a DA neuron-specific transcription factor. These criteria also correlated well with robust GABA_BR-dependent somatodendritic outward currents in slices from wild-type mice (Fig. 2A). Peak outward currents evoked by the GABA_BR agonist baclofen measured in slices from *Girk1*^{-/-} and *Girk3*^{-/-} mice were indistinguishable from wild-type, whereas currents in slices from *Girk2*^{-/-} mice were dramatically attenuated (Fig. 2A,B).

Girk ablation had no impact on apparent capacitance, input resistance, I_h current amplitude, or action potential duration in putative VTA DA neurons (not shown). Genotype-dependent differences in spontaneous rates, however, were evident. Surprisingly, VTA DA neurons from *Girk2*^{-/-} mice exhibited lower rates than wild-type, *Girk1*^{-/-}, and *Girk3*^{-/-} counterparts (Fig. 2C). Firing rates of VTA DA neurons from wild-type slices were consistently and reversibly lowered in the presence of the *Girk* channel blocker tertiapin, suggesting that the lower firing rates observed in DA neurons from *Girk2*^{-/-} mice were largely attributable to the loss of *Girk* channel function (Fig. 2C). Indeed, tertiapin application had no effect on firing rates of VTA DA neurons in slices from *Girk2*^{-/-} mice (not shown).

We hypothesized that VTA DA neurons in *Girk2*^{-/-} mice receive elevated inhibitory input from local GABA interneurons. To test this hypothesis, we first recorded spontaneous inhibitory postsynaptic currents (sIPSCs) in VTA DA neurons from wild-type and *Girk*^{-/-} mice (Fig. 3). The frequency and amplitude of sIPSCs measured in wild-type mice were consistent with previous reports (Mathon et al. 2005). Consistent with our prediction, sIPSC frequency was elevated in DA neurons from *Girk2*^{-/-} mice, relative to frequencies measured in wild-type, *Girk1*^{-/-}, and *Girk3*^{-/-} mice (Fig. 3A,B). sIPSC amplitudes did not differ across genotypes (Fig. 3C). Second, we examined the impact of picrotoxin on the spontaneous activity of VTA DA neurons in slices from wild-type and *Girk*^{-/-} mice, reasoning that GABA_A receptor blockade should normalize firing rates if the lower basal firing rates seen in *Girk2*^{-/-} mice were attributable to elevated GABAergic input. Indeed, VTA DA neuron firing rates did not differ across genotypes in the presence of picrotoxin application (Fig. 3D).

Elevated inhibitory input to, and lower *ex vivo* firing rates of, VTA DA neurons in *Girk2*^{-/-} mice is inconsistent with the DA-related phenotypes shown in Fig. 1 and reported previously (Blednov et al. 2001; Blednov et al. 2002; Morgan et al. 2003). We reasoned that *Girk2* ablation may also trigger adaptations facilitating excitatory neurotransmission in VTA DA neurons, and that the influence of these adaptations would be muted due to the transection of excitatory afferents that occurs during slice preparation. To address this possibility, we first measured spontaneous excitatory postsynaptic currents (sEPSCs) in VTA DA neurons from wild-type and *Girk*^{-/-} mice (Fig. 4). sEPSCs were readily observed in VTA DA neurons from wild-type mice (Fig. 4A), with amplitudes and frequencies consistent with previous reports using similar conditions (Engblom et al. 2008; Zweifel et al. 2008). sEPSC frequency was elevated significantly and similarly in VTA DA neurons from *Girk1*^{-/-} and *Girk2*^{-/-} mice, but not *Girk3*^{-/-} mice (Fig. 4B). Interestingly, sEPSC amplitudes were elevated significantly in *Girk2*^{-/-} mice (Fig. 4C).

We next examined the synaptic levels of AMPA receptor subunits in VTA DA neurons using double-labeling post-embedding immunoelectron microscopy (Fig. 5). Specifically, the relative amounts of the AMPA receptor subunits GluR1 (Fig. 5A-D) and GluR2/3 (Fig.

5E-H) in the postsynaptic density (PSD) in VTA DA neurons (identified by tyrosine hydroxylase labeling) were compared in wild-type, *Girk1*^{-/-}, and *Girk2*^{-/-} mice. Synaptic densities of GluR1 and GluR2/3 levels were significantly elevated (~2-fold) in VTA DA neurons from *Girk1*^{-/-} and *Girk2*^{-/-} mice, suggesting that the elevated frequency of sEPSCs measured in VTA DA neurons from *Girk1*^{-/-} and *Girk2*^{-/-} mice is due to the increased synaptic distribution of AMPA receptors.

Persistent elevation in glutamatergic signaling in VTA DA neurons can trigger adaptations in the NAcc, including increased glutamatergic input, dendritic spine density, and expression of the AMPA receptor subunit GluR1 (Robinson and Kolb 2004; Boudreau and Wolf 2005; Pulipparacharuvil et al. 2008; Mameli et al. 2009). Given the evidence for increased glutamatergic neurotransmission in VTA DA neurons from *Girk1*^{-/-} and *Girk2*^{-/-} mice, we next probed for adaptations related to excitatory neurotransmission in the NAcc of these mice. Specifically, we measured miniature EPSCs (mEPSCs) in NAcc shell neurons from wild-type and *Girk*^{-/-} mice (Fig. 6). We found that mEPSC frequency and amplitude were elevated in NAcc neurons from *Girk1*^{-/-} and *Girk2*^{-/-} mice, relative to wild-type controls. Interestingly, mEPSC frequency and amplitude were significantly higher in NAcc neurons from *Girk2*^{-/-} mice as compared to neurons from *Girk1*^{-/-} mice.

We next examined and quantified key cellular and molecular markers of excitatory neurotransmission in the NAcc of wild-type and *Girk*^{-/-} mice. First, we quantified excitatory synapse density in medium spiny neurons of the NAcc using an optical dissector approach (Sterio 1984). Synapse density in the NAcc from *Girk1*^{-/-} (1.31 synapses/ μm^3) and *Girk2*^{-/-} (1.35 synapses/ μm^3) mice was ~15% higher as compared to wild-type control (1.12 synapses/ μm^3) (Fig. 7A-D). Moreover, synaptic densities of GluR1 (Fig. 7E-H) & GluR2/3 (Fig. 7I-L) were significantly elevated (~2-fold) in medium spiny neurons in the NAcc from *Girk1*^{-/-} and *Girk2*^{-/-} mice. Similar increases in synapse density and AMPA subunit labeling were observed in both the NAcc shell and core.

DISCUSSION

Long-loop (GABA_BR-dependent) and autocrine (D₂R-dependent) inhibitory feedback to VTA DA neurons is triggered by cocaine-induced increases in DA levels in the NAcc and VTA, respectively, and lead to tempered VTA DA neuron output and motor stimulation (Wolf et al. 1978; Einhorn et al. 1988; Napier and Potter 1989; Chen and Reith 1994). Simultaneous pharmacological inhibition of both feedback pathways evokes hyperactivity and exaggerated responses to psychostimulants, including cocaine (Steketee et al. 1991, 1992; Narayanan et al. 1996). Since *Girk* channels mediate the postsynaptic inhibitory effects linked to GABA_BR and D₂R activation, *Girk* ablation in midbrain DA neurons should promote hyperactivity and exaggerated motor-stimulatory responses to cocaine due to the weakening of these inhibitory feedback mechanisms. Moreover, given that the *Girk* channel in midbrain DA neurons contains *Girk2* but not *Girk1*, one would predict that hyperactivity and enhanced responses to cocaine would be seen in *Girk2*^{-/-} but not *Girk1*^{-/-} mice. To the contrary, *Girk1*^{-/-} and *Girk2*^{-/-} mice showed comparable baseline hyperactivities and enhanced responses to acute cocaine. Thus, the loss of inhibitory feedback to VTA DA neurons cannot explain the DA-dependent phenotypes manifest in *Girk2*^{-/-} mice.

By the same logic, one can conclude that alterations in intrinsic excitability of VTA DA neurons are not the primary cause of the DA-dependent phenotypes in *Girk2*^{-/-} mice. Indeed, ablation of *Girk2* but not *Girk1* should enhance the intrinsic excitability of VTA DA neurons, particularly if these neurons exhibit a high degree of basal *Girk* channel activity. While intrinsic excitability of VTA DA neurons may be elevated in *Girk2*^{-/-} mice, these

neurons exhibited lower spontaneous activity *ex vivo*. Firing rates were normalized by picrotoxin, indicating that VTA DA neurons from *Girk2*^{-/-} mice receive elevated inhibitory input from local GABA neurons (Sugita et al. 1992). While this elevated inhibitory tone may be offset *in vivo* by elevated excitatory input (as our data suggest), these data do not support the contention that the DA-dependent hyperactivity in *Girk2*^{-/-} mice reflects the enhanced intrinsic excitability of VTA DA neurons.

Chronic elevation of glutamatergic signaling in VTA DA neurons promotes persistent adaptations in glutamatergic signaling in the NAcc (Mameli et al. 2009). These adaptations are thought to underlie drug-seeking behavior and behavioral sensitization (Kalivas 2004; Self 2004). Interestingly, *Girk1*^{-/-} and *Girk2*^{-/-} mice show enhanced responses to acute cocaine administration, and the NAcc adaptations seen in *Girk1*^{-/-} and *Girk2*^{-/-} mice, including increased synapse density and elevated excitatory signaling, overlap with those in animals treated chronically with psychostimulants. Enhanced excitatory signaling in the NAcc, however, is thought to be a response to cocaine withdrawal rather than a consequence of repeated cocaine exposure (Kourrich et al. 2007). As such, adaptations in drug-naïve, constitutive *Girk1*^{-/-} and *Girk2*^{-/-} mice may be distinct from those linked to drug exposure.

The adaptations observed in the mesolimbic DA system of *Girk1*^{-/-} and *Girk2*^{-/-} mice may reflect synaptic scaling, a form of homeostatic plasticity triggered by persistent alterations in factors that influence neuronal excitability (Turrigiano and Nelson 2004; Kato et al. 2007; Thiagarajan et al. 2007). Synaptic scaling, manifest as an increase in the synaptic expression of GluR1 and GluR2-containing AMPA receptors, has been reported in the NAcc during cocaine withdrawal (Boudreau and Wolf 2005; Boudreau et al. 2007; Kourrich et al. 2007; Boudreau et al. 2009). Similarly, we observed elevated synaptic levels of both GluR1 and GluR2/3 in VTA DA and NAcc medium spiny neurons in *Girk1*^{-/-} and *Girk2*^{-/-} mice. It is possible, therefore, that the enhanced synaptic AMPA receptor distribution and elevated glutamatergic neurotransmission in VTA DA neurons and NAcc medium spiny neurons from *Girk1*^{-/-} and *Girk2*^{-/-} mice reflects compensatory responses of these neurons to persistent inhibition.

While the similarities between *Girk1*^{-/-} and *Girk2*^{-/-} mice with respect to adaptations relevant to excitatory neurotransmission are striking, some differences were evident. Notably, sEPSC amplitudes measured in VTA DA neurons were greater in *Girk2*^{-/-} mice as compared to *Girk1*^{-/-} mice. Similarly, mEPSC frequency and amplitude were both significantly elevated in NAcc medium spiny neurons in *Girk2*^{-/-} mice relative to *Girk1*^{-/-} mice. These differences may result directly from the loss of Girk signaling in VTA DA neurons, or may reflect more global differences linked to the relative impact of *Girk1* and *Girk2* ablation on neuronal physiology. While the genetic ablation of *Girk1* and *Girk2* yields similar outcomes in terms of somatodendritic currents in hippocampal pyramidal, locus ceruleus, and spinal cord dorsal horn neurons (Koyrakh et al. 2005; Marker et al. 2006; Cruz et al. 2008), *Girk2* ablation may have a more dramatic impact than *Girk1* ablation on synaptic Girk currents, which likely account for a small fraction of the somatodendritic currents measured in most studies. In support of this contention, a significant fraction of Girk2 protein, but not Girk1, is found within the postsynaptic specialization at excitatory synapses (Koyrakh et al. 2005; Marker et al. 2006). Clearly, understanding the more subtle differences between *Girk1*^{-/-} and *Girk2*^{-/-} mice reported in this study and other studies will require more selective genetic and/or pharmacologic approaches.

In summary, we demonstrate that genetic ablation of *Girk1* and *Girk2* promotes adaptations in the mesolimbic DA system that facilitate excitatory glutamatergic neurotransmission. The adaptations observed are reminiscent of those reported for synaptic scaling and following drug administration, suggesting that Girk channels may normally serve as a barrier to such

adaptations. Accordingly, the regulation of Girk signaling strength, perhaps in VTA DA neurons and/or in neurons that provide input to the VTA, may constitute in part the mechanism by which drugs of abuse evoke adaptations that promote chronic drug intake, or provide a target for therapeutic interventions aimed at preventing such adaptations.

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Abbreviations used

VTA	ventral tegmental area
NAcc	nucleus accumbens
DA	dopamine
Girk	G protein-gated inwardly-rectifying K ⁺ channel
SN	substantia nigra
GABA_BR	GABA _B receptor
D₁R	D1 dopamine receptor
D₂R	D2 dopamine receptor
AMPA	α-amino-3-hydroxyl-5-methyl-4-isoxazole-propionate
EPSC	excitatory postsynaptic current
IPSC	inhibitory postsynaptic current
PSD	postsynaptic density

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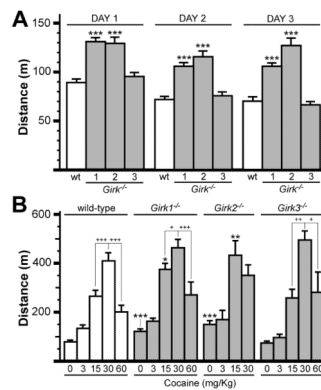


Figure 1. Baseline and cocaine-induced motor activity in wild-type and *Girk*^{-/-} mice
A) Total distance traveled (m) for wild-type (wt), *Girk1*^{-/-}, *Girk2*^{-/-}, and *Girk3*^{-/-} mice in open-field environments during 1-h sessions on 3 consecutive days (n=36-81 per genotype). Statistical symbols: *** – $P < 0.001$, vs. wild-type (within day). **B)** Cocaine-induced motor activity in wild-type and *Girk*^{-/-} mice. Subjects were challenged with saline or one of 4 cocaine doses (3, 15, 30, 60 mg/kg i.p.) and total distance traveled (m) was measured during the 60-min post-injection window. Cocaine triggered dose-dependent increases in motor activity in wild-type ($F_{4,125}=54.6$; $P < 0.001$), *Girk1*^{-/-} ($F_{4,130}=65.3$; $P < 0.001$), *Girk2*^{-/-} ($F_{3,52}=14.4$; $P < 0.001$), and *Girk3*^{-/-} ($F_{4,81}=65.3$; $P < 0.001$) mice. Genotype-dependent differences in cocaine-induced responses were observed following injection of 0 ($F_{3,192}=23.0$; $P < 0.001$) and 15 mg/kg cocaine ($F_{3,87}=6.1$; $P < 0.001$), but not after the 3 ($F_{3,41}=2.0$; $P=0.1$), 30 ($F_{3,43}=1.8$; $P=0.2$), or 60 ($F_{3,41}=2.0$; $P=0.1$) mg/kg injections (n=6-30 per genotype and dose). Statistical symbols: *, **, *** – $P < 0.05$, 0.01, and 0.001, respectively, vs. wild-type (within dose); within-genotype comparisons are displayed to convey differences observed at the higher cocaine doses: +, ++, +++ – $P < 0.05$, 0.01, and 0.001, respectively, relative to the response at 30 mg/kg cocaine.

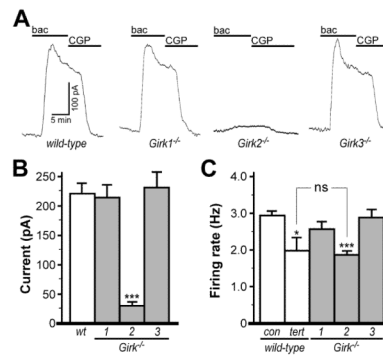


Figure 2. Functional characterization of VTA DA neurons from wild-type and *Girk*^{-/-} mice
A) Outward currents evoked by baclofen (bac, 200 μ M) in VTA DA neurons from wild-type, *Girk1*^{-/-}, *Girk2*^{-/-}, and *Girk3*^{-/-} mice. Outward currents corresponded with a decrease in input resistance and were sensitive to 0.3 external Ba²⁺ (not shown), and were reversed by the GABA_BR antagonist CGP54626 (CGP, 2 μ M). **B)** Summary of baclofen-induced outward currents in VTA DA neurons from wild-type (219±17 pA), *Girk1*^{-/-} (212±21 pA), *Girk2*^{-/-} mice (31±4 pA), and *Girk3*^{-/-} (229±37 pA) mice (n=6-22 per genotype). Genotype-dependent differences in peak baclofen-induced current amplitude were evident ($F_{3,39}=11.9$, $P<0.001$). Symbols: ***, $P<0.001$ vs. wild-type. **C)** A significant effect of group on firing rate was observed ($F_{4,147}=10.5$, $P<0.001$) with respect to *ex vivo* firing frequencies of VTA DA neurons from wild-type mice at baseline (con, 2.93±0.12 Hz) and following application of the Girk channel blocker tertiapin (tert, 1.98±0.34 Hz), and at baseline in slices from *Girk1*^{-/-} (2.55±0.20 Hz), *Girk2*^{-/-} mice (1.84±0.11 Hz) and *Girk3*^{-/-} (2.87±0.22 Hz) mice (n=9-49 per group). Note that there was no significant (ns) difference in firing rates of VTA DA neurons from *Girk2*^{-/-} mice and wild-type neurons treated with tertiapin. Symbols: *, ***, $P<0.05$ and 0.001, respectively, vs. wild-type.

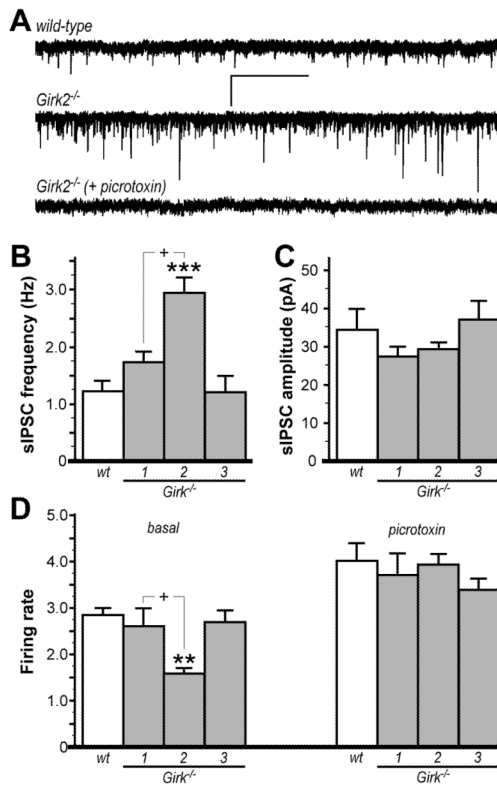


Figure 3. Inhibitory input to VTA DA neurons in wild-type and *Girk*^{-/-} mice

A) Representative traces of sIPSCs measured in VTA DA neurons from wild-type (upper) and *Girk2*^{-/-} mice (middle); sIPSCs were blocked by the GABA receptor antagonist picrotoxin (100 μ M, lower). Scale bars: 40 pA/5s. Summary histograms of sIPSC frequency and amplitude in VTA DA neurons from wild-type (wt, 1.2 \pm 0.2 Hz, 34.4 \pm 5.7 pA), *Girk1*^{-/-} (1.7 \pm 0.2 Hz, 26.9 \pm 3.1 pA), *Girk2*^{-/-} (2.9 \pm 0.3 Hz, 29.2 \pm 2.2 pA), and *Girk3*^{-/-} (1.2 \pm 0.3 Hz, 36.6 \pm 5.5 pA), mice are shown in panels **B** and **C**, respectively (n=7-10 per genotype). A significant effect of genotype was observed for sIPSC frequency ($F_{3,29}=10.1$; $P<0.001$) but not amplitude ($F_{3,29}=1.1$; $P=0.4$). Symbols: *** $P<0.001$ vs. wild-type and *Girk3*^{-/-}; + $P<0.05$, *Girk1*^{-/-} vs. *Girk2*^{-/-}. **D)** Firing frequency of VTA DA neurons from wild-type and *Girk*^{-/-} mice (n=8-21 per genotype) were measured at baseline (left) and following administration of 100 μ M picrotoxin (right). An effect of genotype on firing rates was observed at baseline ($F_{3,56}=10.0$; $P<0.001$) but not after picrotoxin application ($F_{3,56}=1.0$; $P=0.4$). Symbols: ** $P<0.01$ vs. wild-type and *Girk3*^{-/-}; + $P<0.05$, *Girk1*^{-/-} vs. *Girk2*^{-/-}.

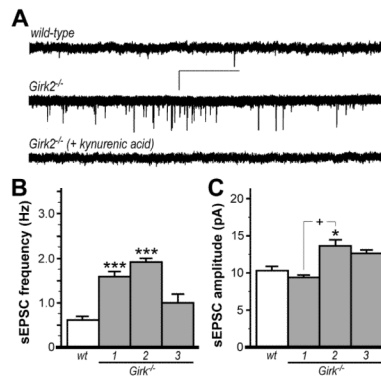


Figure 4. Excitatory input to VTA DA neurons in wild-type and *Girk*^{-/-} mice

A) Representative traces of sEPSCs measured in VTA DA neurons from wild-type (upper) and *Girk2*^{-/-} mice (middle); sEPSCs were blocked by the non-selective ionotropic glutamate receptor antagonist kynurenic acid (2 mM; lower). Scale bars: 20 pA/5 s. Summary histograms of sEPSC frequency and amplitude in VTA DA neurons from wild-type (wt, 0.6±0.1 Hz, 10.4±0.8 pA), *Girk1*^{-/-} (1.6±0.1 Hz, 8.9±0.4 pA), *Girk2*^{-/-} (1.9±0.1 Hz, 13.5±0.8 pA), and *Girk3*^{-/-} (1.0±0.2 Hz, 12.7±0.4 pA), mice are shown in panels B and C, respectively (n=7-12 per genotype). A significant effect of genotype was observed for both sEPSC frequency ($F_{3,31}=18.4$; $P<0.001$) and amplitude ($F_{3,31}=8.8$; $P<0.001$). Symbols: *,*** $P<0.05$ and 0.001, respectively, vs. wild-type; + $P<0.001$, *Girk1*^{-/-} vs. *Girk2*^{-/-}.

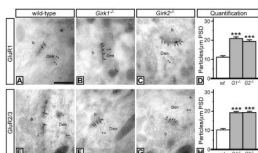


Figure 5. AMPA subunit labeling in VTA DA neurons from wild-type and *Girk*^{-/-} mice
 Electron micrographs showing GluR1 (**A-C**) and GluR2/3 (**E-G**) immunoreactivities in VTA DA neurons from wild-type, *Girk1*^{-/-}, and *Girk2*^{-/-} mice, as detected using double-labeling, post-embedding immunogold electron microscopy. In sections from wild-type mice, immunoparticles for GluR1 (**A**) and GluR2/3 (**E**) subunits (arrows, 10 nm gold particles) were found along the PSDs of dendritic shafts (Den) of DA neurons (TH-positive cells, arrowheads, 20 nm gold particles) establishing contact with axon terminals (b). In sections from *Girk1*^{-/-} mice (**B,F**) and *Girk2*^{-/-} mice (**C,G**), an increase in the number of immunoparticles for both GluR1 and GluR2/3 subunits along the PSD was observed. These images are representative of data from three separate panels of wild-type and *Girk*^{-/-} mice. Scale bar: 0.2 μ m. **D,H**) Quantification of GluR1 and GluR2/3 synaptic density in VTA DA neurons from wild-type, *Girk1*^{-/-} (*G1*^{-/-}), and *Girk2*^{-/-} (*G2*^{-/-}) mice. Synaptic density is expressed as number of immunoparticles per 1 μ m length of PSD. Symbols: *** $P < 0.001$ vs. wild-type.

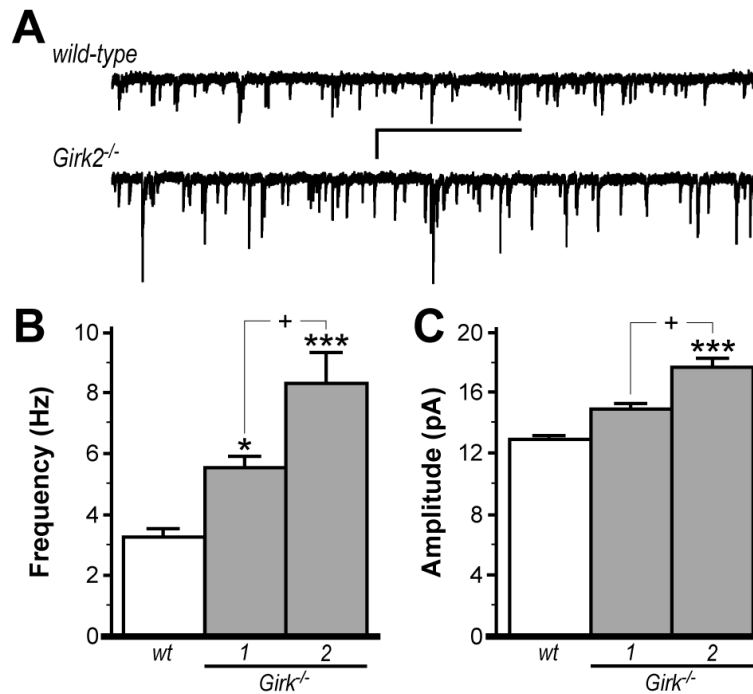


Figure 6. Excitatory input to NAcc medium spiny neurons in wild-type and *Girk*^{-/-} mice
A) Representative traces of mEPSCs in NAcc DA neurons from wild-type (upper) and *Girk2*^{-/-} (lower) mice. Scale bars: 2 s/5 pA. mEPSCs were blocked completely by the non-selective ionotropic glutamate receptor antagonist kynurenic acid (2 mM, not shown). Summary of mEPSC frequency and amplitude for medium spiny neurons in the NAcc shell from wild-type (wt, 3.3±0.3 Hz, 12.9±0.4 pA), *Girk1*^{-/-} (5.5±0.4 Hz, 14.6±0.7 pA), and *Girk2*^{-/-} (8.3±1.0 Hz, 17.1±0.8 pA) mice are shown in panels **B** and **C**, respectively (n=11-23 per genotype). Genotype-dependent differences were observed for both mEPSC frequency ($F_{2,44}=21.8$; $P<0.001$) and amplitude ($F_{2,44}=14.6$; $P<0.001$). Symbols: *,*** $P<0.05$ and 0.001 , respectively, vs. wild-type; + $P<0.05$, *Girk1*^{-/-} vs. *Girk2*^{-/-}.

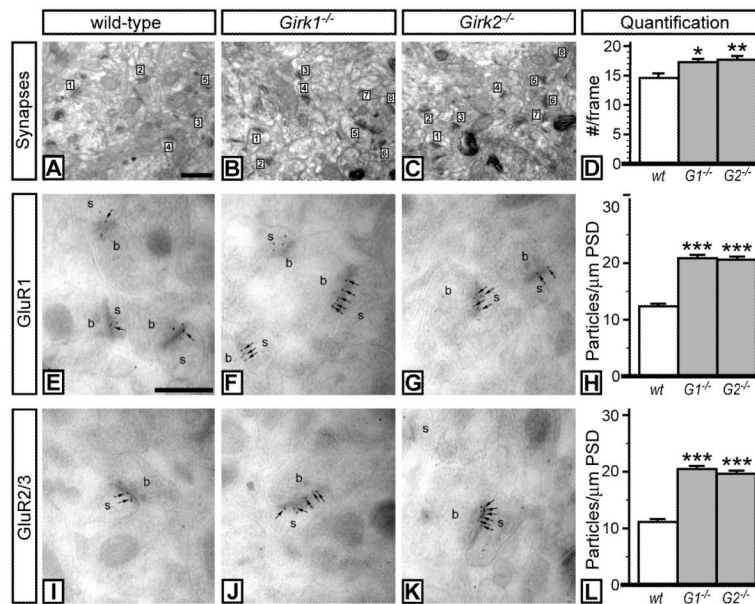


Figure 7. Markers of excitatory neurotransmission in the NAcc of wild-type and *Girk*^{-/-} mice
 Electron micrographs taken from the NAcc core of wild-type (A), *Girk1*^{-/-} (B), and *Girk2*^{-/-} (C) mice, showing excitatory synapses (numbered) in the neuropil. Synapses between dendritic spines and axon terminals that exhibited a clear PSD were counted. Scale bar: 0.2 μm. **D**) Quantification of synapses from wild-type, *Girk1*^{-/-} (*G1*^{-/-}), and *Girk2*^{-/-} (*G2*^{-/-}) mice. The number of synapses per frame (#/frame) is plotted as a function of genotype. Synapses were tabulated in 21 frames per genotype, from 3 different animals per genotype. A significant effect of genotype on synapse density was observed ($F_{2,60}=5.4$; $P<0.01$). Symbols: *,** $P<0.05$ and 0.01 , respectively, vs. wild-type. **E-L**) AMPA subunit labeling in NAcc medium spiny neurons from wild-type and *Girk*^{-/-} mice. Electron micrographs showing GluR1 (**E-G**) and GluR2/3 (**I-K**) immunoreactivity in medium spiny neurons from wild-type, *Girk1*^{-/-}, and *Girk2*^{-/-} mice, as detected using post-embedding immunogold electron microscopy. In sections from wild-type mice, immunoparticles for GluR1 (**E**) and GluR2/3 (**I**) subunits (arrows, 10 nm gold particles) were found along PSDs on individual spines (s) in contact with axon terminals (b). In sections from *Girk1*^{-/-} (**F,J**) and *Girk2*^{-/-} (**G,K**) mice, an increase in the number of immunoparticles for both GluR1 and GluR2/3 subunits along the PSD was observed. Scale bar: 0.2 μm. **H,L**) Quantification of GluR1 and GluR2/3 synaptic density in Nacc medium spiny neurons from wild-type, *Girk1*^{-/-} (*G1*^{-/-}), and *Girk2*^{-/-} (*G2*^{-/-}) mice. Synaptic density is expressed as number of immunoparticles per 1 μm length of PSD. Symbols: *** $P<0.001$ vs. wild-type.