

# Delta glutamate receptor conductance drives excitation of mouse dorsal raphe neurons

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**Abstract** The dorsal raphe nucleus is the predominant source of central serotonin, where neuronal activity regulates complex emotional behaviors. Action potential firing of serotonin dorsal raphe neurons is driven via  $\alpha$ 1-adrenergic receptors ( $\alpha$ 1-A<sub>R</sub>) activation. Despite this crucial role, the ion channels responsible for  $\alpha$ 1-A<sub>R</sub>-mediated depolarization are unknown. Here, we show in mouse brain slices that  $\alpha$ 1-A<sub>R</sub>-mediated excitatory synaptic transmission is mediated by the ionotropic glutamate receptor homolog cation channel, delta glutamate receptor 1 (GluD1). GluD1<sub>R</sub>-channels are constitutively active under basal conditions carrying tonic inward current and synaptic activation of  $\alpha$ 1-A<sub>R</sub>s augments tonic GluD1<sub>R</sub>-channel current. Further, loss of dorsal raphe GluD1<sub>R</sub>-channels produces an anxiogenic phenotype. Thus, GluD1<sub>R</sub>-channels are responsible for  $\alpha$ 1-A<sub>R</sub>-dependent induction of persistent pacemaker-type firing of dorsal raphe neurons and regulate dorsal raphe-related behavior. Given the widespread distribution of these channels, ion channel function of GluD1<sub>R</sub> as a regulator of neuronal excitability is proposed to be widespread in the nervous system.

## Introduction

Recent reports estimate that 1 in 5 adults worldwide are affected by a mental health disorder, with anxiety and depression being the most common affecting more than 260 million people (*GBD 2017 Disease and Injury Incidence and Prevalence Collaborators, 2018*). Most current pharmacotherapies to treat these disorders target serotonin receptors or serotonin clearance. The dorsal raphe nucleus is the largest serotonergic nucleus in the brain and the predominant source of central serotonin (5-HT). In vivo, tonic noradrenergic input to the dorsal raphe that activates G $\alpha$ <sub>q/11</sub> protein-coupled  $\alpha$ 1-adrenergic receptors ( $\alpha$ 1-A<sub>R</sub>s) is required for 5-HT neurons to fire action potentials (*Baraban and Aghajanian, 1980; Baraban et al., 1978*) and release 5-HT (*Clement et al., 1992*). In dorsal raphe brain slices, synaptic activation of  $\alpha$ 1-A<sub>R</sub>s produces a slow membrane depolarization lasting tens of seconds (*Yoshimura et al., 1985*). Despite having a crucial role in regulating 5-HT neuron excitability, the ion channels responsible for the depolarization remain unknown.

Throughout the central and peripheral nervous system, activation of G $\alpha$ <sub>q/11</sub> protein-coupled receptors (G<sub>q</sub>PCRs), namely metabotropic glutamate mGlu<sub>R</sub>s, muscarinic acetylcholine M1 (mACh<sub>R</sub>s), or  $\alpha$ 1-A<sub>R</sub>s produces slow, noisy inward currents. Multiple mechanisms have been reported to underlie the inward current including: inhibition of K<sup>+</sup> current (including leak, Ca<sup>2+</sup>-activated, and Kv7/M-current) (*Benson et al., 1988; Halliwell and Adams, 1982; Madison et al., 1987; Shen and North, 1992*), modulation of TTX-sensitive persistent Na<sup>+</sup> current (*Yamada-Hanff and Bean, 2013*), and activation of transient potential receptor canonical (TRPC) (*Hartmann et al., 2008; Kim et al., 2003*), Na<sup>+</sup>-leak (NALCN) (*Lu et al., 2009*), or delta glutamate receptor-channels (*Ady et al., 2014; Benamer et al., 2018*).

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**eLife digest** Serotonin is a chemical that allows cells to communicate in the nervous system of many animals. It is also particularly important in the treatment of mental health disorders: a large number of antidepressants work by preventing nerve cells from clearing away serotonin, therefore increasing the overall level of the molecule in the brain. Yet, exactly how serotonin is released remains unclear.

When a serotonin-producing cell is activated, a series of biochemical processes lead to the creation of an electric current that, ultimately, is required for the cell to release serotonin. This mechanism starts when the  $\alpha$ 1-adrenergic receptor, a protein at the surface of the cell, detects noradrenaline molecules. However, on its own, the  $\alpha$ 1-adrenergic receptor is unable to create an electric current: this requires ion channels, another type of protein which can let charged particles in and out of the cell. Here, Gantz et al. set out to determine the identity of the ion channel that allows noradrenaline signals to generate electrical activity in cells which can release serotonin.

Electrical and chemical manipulation of mouse brain slices revealed that an ion channel called delta-glutamate 1 was active in serotonin-producing cells exposed to noradrenaline. In fact, applying toxins that specifically blocked the activity of this channel also prevented the cells from responding electrically to noradrenaline.

Further experiments used mice whose serotonin-producing cells were genetically modified to turn off delta-glutamate 1. In turn, these animals showed anxiety-like behaviors, which could be consistent with a drop in serotonin levels. This is in line with previous human studies showing that patients with depression and other mental health conditions have mutations in the gene for delta-glutamate 1.

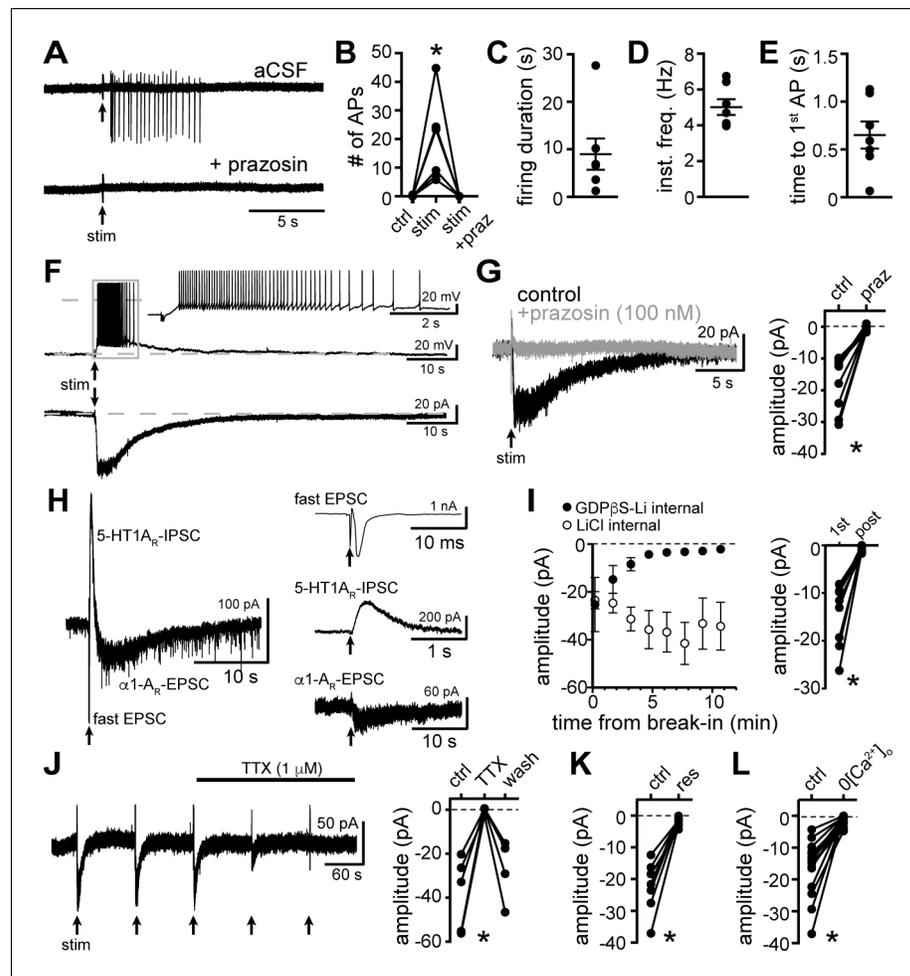
Taken together, these results give an insight into the electrical activity of serotonin-producing cells. Further work is now required to examine how changes in the gene that codes for delta-glutamate 1 ultimately affect the release of serotonin. This could potentially help to understand if certain individuals may not be able to properly produce this chemical. As many antidepressants work by retaining serotonin that is already present in the brain, this knowledge could ultimately help patients who do not currently respond to treatment.

The delta glutamate receptors, GluD1<sub>R</sub> and GluD2<sub>R</sub>, are mysterious members of the ionotropic glutamate receptor family in that they are not gated by glutamate (Araki et al., 1993; Lomeli et al., 1993). One theory is that they are strictly scaffolding proteins or synaptic organizers, rather than ion conducting channels. But wild-type channels have been reported to conduct in response to activation of mGlu<sub>R</sub>s (Ady et al., 2014; Benamer et al., 2018). GluD1<sub>R</sub> (*Grid1*) mRNA is expressed widely throughout the brain, with notably high levels in the dorsal raphe (Hepp et al., 2015; Konno et al., 2014). Here, we used a combination of in vitro patch-clamp electrophysiology and pharmacology with a CRISPR/Cas9 viral genetic strategy to determine that activation of  $\alpha$ 1-A<sub>R</sub>s in the dorsal raphe depolarizes neurons via GluD1<sub>R</sub>-channel conductance. We utilize the  $\alpha$ 1-A<sub>R</sub>-GluD1<sub>R</sub>-EPSC to explore conduction and biophysical properties of GluD1<sub>R</sub>-channels, to ultimately glean a greater understanding of GluD1<sub>R</sub>-channel gating. Lastly, we demonstrate that functional deletion of GluD1<sub>R</sub>-channels in the dorsal raphe produces an anxiogenic behavioral phenotype.

## Results

### Synaptic activation of $\alpha$ 1-adrenergic receptors produces an EPSC

Electrophysiological recordings were made from dorsal raphe neurons in acute brain slices from wild-type mice at 35° C in the presence of NMDA<sub>R</sub>, AMPA<sub>R</sub>, Kainate<sub>R</sub>, GABA-A<sub>R</sub>, and 5-HT1A<sub>R</sub> antagonists. With cell-attached recordings, a train of 5 electrical stimuli (60 Hz), delivered to the brain slice via a monopolar stimulating electrode, produced firing in previously quiescent neurons, which was blocked by application of the  $\alpha$ 1-A<sub>R</sub> antagonist, prazosin (100 nM, **Figure 1A**). The excitation produced 20±5 action potentials that lasted 9.0±3.0 s, with a latency of 650.6±0.1 ms from onset of stimulation to the first action potential (**Figure 1B-E**). In whole-cell recording using a potassium-based internal solution, the same train of electrical stimuli produced prolonged action potential



**Figure 1.** Electrical stimulation evokes long-lasting action potential firing produced by an  $\alpha$ 1-adrenergic receptor-dependent EPSC. (A) Representative traces of cell-attached recording where stimulation of the brain slice (5 stim at 60 Hz) produced action potential firing that was abolished by application of the  $\alpha$ 1-adrenergic receptor antagonist, prazosin (100 nM). (B) Plot of number of action potentials showing the stimulation-induced increase in frequency ( $p=0.004$ ,  $n=6$ ). (C) Plot of duration of action potential firing. (D) Plot of mean instantaneous frequency of action potential firing over the first 10 s of firing. (E) Plot of the latency from stimulation onset to the first action potential. (F) Example whole-cell recordings in the same cell, where electrical stimulation of the slice produced prolonged action potential firing in current-clamp (upper trace) and a slow EPSC in voltage-clamp mode (lower trace). (G) Bath application of prazosin eliminated the slow EPSC shown in a representative trace (left, baseline adjusted) and in grouped data (right,  $p=0.002$ ,  $n=10$ ). (H) Representative traces of a whole-cell recording when the brain slice was stimulated in the absence of antagonists showing the kinetics of the  $\alpha$ 1- $A_R$ -EPSC relative to the fast EPSC (peak has been truncated) and 5HT1 $A_R$ -IPSC (left). Subsequent addition of AMPA $_R$ /Kainate $_R$  and GABA $_R$  and 5-HT1 $A_R$  antagonists revealed the remaining synaptic current produced by  $\alpha$ 1- $A_R$  activation (right). Time of stimulations are marked by arrows. (I) With GDP $\beta$ S-Li-containing internal solution, the amplitude of the  $\alpha$ 1- $A_R$ -EPSC ran down within ~5–20 min of break-in to whole-cell mode; shown in a plot compared with control internal solution containing LiCl only (left) and in grouped data (right,  $p=0.004$ ,  $n=9$ , 1 $^{st}$ : first EPSC; post: post-dialysis). (J) Bath application of tetrodotoxin (TTX, 1  $\mu$ M) reversibly eliminated the  $\alpha$ 1- $A_R$ -EPSC shown in a representative trace (left,  $\alpha$ 1- $A_R$ -EPSC evoked every 90 s) and in grouped data (right,  $p=0.009$ ,  $n=4$ ). (K) Plot of the inhibition of  $\alpha$ 1- $A_R$ -EPSC amplitude by application of reserpine (res, 1  $\mu$ M,  $p=0.016$ ,  $n=7$ ). (L) Plot of the inhibition of  $\alpha$ 1- $A_R$ -EPSC amplitude by removal of external  $Ca^{2+}$  ( $0[Ca^{2+}]_o$ ,  $p=0.0001$ ,  $n=14$ ). Line and error bars represent mean  $\pm$  SEM, \* denotes statistical significance.

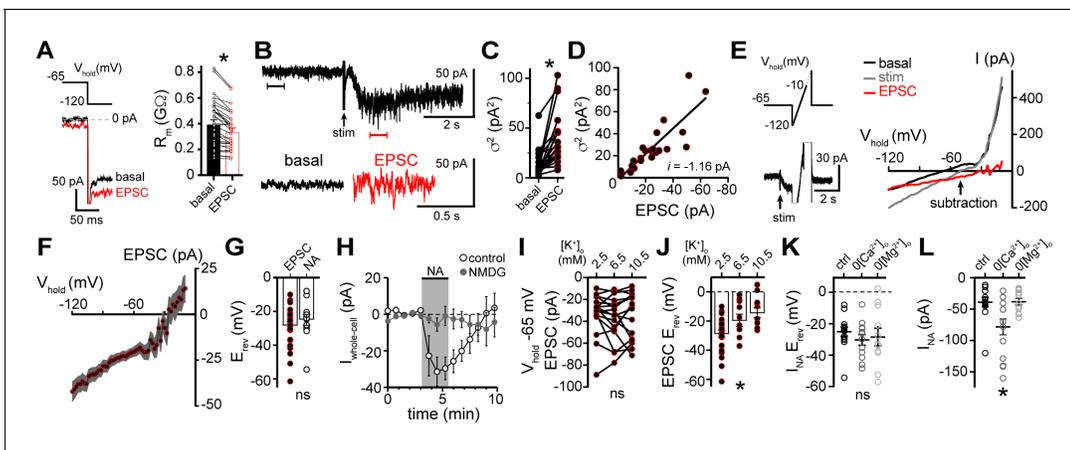
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**Source data 1.** Numerical data that were used to generate graphs in **Figure 1**.

firing (**Figure 1F**). In voltage-clamp mode ( $V_{\text{hold}} -65$  mV), the same stimulation produced a slow and long-lasting ( $27.4 \pm 2.3$  s,  $n=10$ ) excitatory postsynaptic current (EPSC, **Figure 1F**) that was eliminated by the application of prazosin (**Figure 1G**). Prazosin had no effect on basal whole-cell current ( $-3.8 \pm 3.4$  pA,  $p=0.232$ ,  $n=10$ , data not shown) indicating a lack of persistent inward current due to noradrenaline tone. On average, the duration of the  $\alpha_1$ -A<sub>R</sub>-EPSC was orders of magnitude longer than fast AMPA<sub>R</sub> channel-mediated EPSCs ( $\sim 10^{3.5} \times$ ) and  $\sim 18 \times$  longer than 'slow' 5-HT<sub>1A</sub> receptor-G protein-coupled inwardly rectifying potassium channel (GIRK)-dependent IPSCs (**Gantz et al., 2015a; Figure 1H**). To test whether  $\alpha_1$ -A<sub>R</sub>-EPSCs were dependent on G protein-signaling, recordings were made with an internal solution containing GDP $\beta$ S-Li<sub>3</sub> (1.8 mM) in place of GTP. Disruption of G protein signaling with intracellular dialysis of GDP $\beta$ S-Li<sub>3</sub> eliminated the  $\alpha_1$ -A<sub>R</sub>-EPSC within 5-20 mins post-break-in ( $p=0.004$ ,  $n=9$ ), whereas dialysis with LiCl alone had no effect on the amplitude of the  $\alpha_1$ -A<sub>R</sub>-EPSC ( $p=0.625$ ,  $n=4$ , **Figure 1I**). These findings demonstrate a cell-autonomous requirement of G protein signaling in the generation of the  $\alpha_1$ -A<sub>R</sub>-EPSC. Application of tetrodotoxin (1  $\mu$ M) reversibly abolished the  $\alpha_1$ -A<sub>R</sub>-EPSC, demonstrating a dependence on presynaptic action potentials (**Figure 1J**). Disruption of the vesicular monoamine transporter with reserpine (1  $\mu$ M) or removal of external  $\text{Ca}^{2+}$  also eliminated the  $\alpha_1$ -A<sub>R</sub>-EPSC, indicating noradrenaline release is vesicular (**Figure 1K and L**).

### Biophysical properties of the channel

Under our recording conditions, resistance of the membrane ( $R_m$ ) significantly decreased during the  $\alpha_1$ -A<sub>R</sub>-EPSC, indicative of opening of ion channels (**Figure 2A**). Membrane noise variance ( $\sigma^2$ ) increased significantly during the EPSC compared to membrane noise under basal conditions



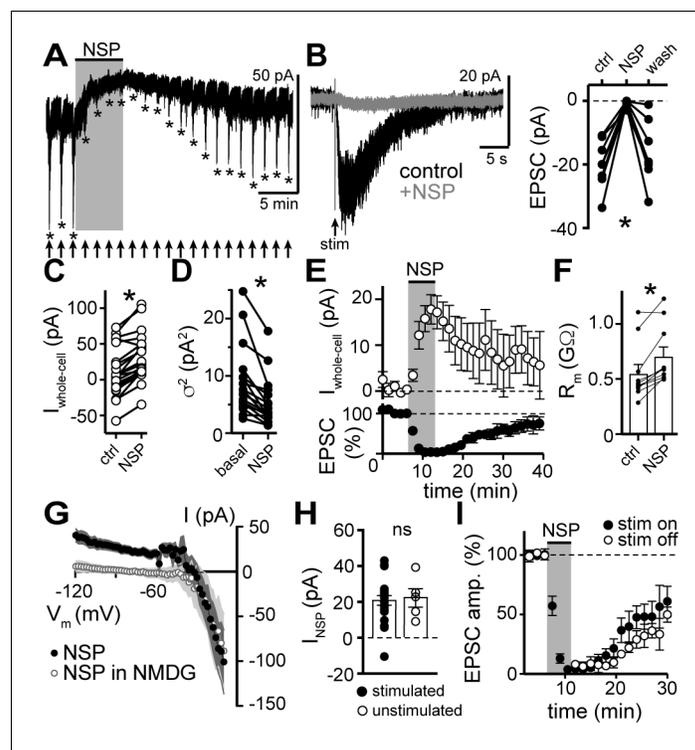
**Figure 2.**  $\alpha_1$ -adrenergic receptor-dependent inward current is carried by sodium entry. (A) Membrane resistance ( $R_m$ ,  $\Delta V$   $-65$  to  $-120$  mV) decreased during the  $\alpha_1$ -A<sub>R</sub>-EPSC indicating an opening of ion channels, as shown in an example trace (left) and in grouped data (right,  $p < 0.0001$ ,  $n = 31$ ). (B) Representative trace of membrane noise during the  $\alpha_1$ -A<sub>R</sub>-EPSC, brackets denote segments shown below on an expanded scale. (C) Membrane noise (variance,  $\sigma^2$ ) increased during the  $\alpha_1$ -A<sub>R</sub>-EPSC ( $p < 0.0001$ ,  $n = 22$ ). (D) Plot of  $\alpha_1$ -A<sub>R</sub>-EPSC variance versus mean amplitude, linear fit represents mean unitary current ( $i$ ,  $r^2 = 0.713$ ,  $p < 0.0001$ ). (E) Slow voltage ramps (1 mV/10 ms, analyzed from  $-120$  to  $-10$  mV) were used to determine the current-voltage relationship of the  $\alpha_1$ -A<sub>R</sub>-EPSC (subtraction), determined by subtracting current at the peak of the  $\alpha_1$ -A<sub>R</sub>-EPSC (stim) from current measured in control conditions just prior to stimulation (basal). Current generated during ramps were truncated for clarity. (F) Current-voltage relationship of the  $\alpha_1$ -A<sub>R</sub>-EPSC from grouped data. Shaded area represents mean  $\pm$  SEM. (G) Plot of reversal potentials ( $E_{\text{rev}}$ ) of the  $\alpha_1$ -A<sub>R</sub>-EPSC and  $I_{\text{NA}}$  ( $p > 0.999$ ,  $n = 26$  and 14). (H) Replacing 126 mM NaCl with NMDG eliminated inward  $I_{\text{NA}}$ , shown in a time-course plot ( $V_{\text{hold}} -65$  mV,  $p < 0.0001$ ,  $n = 14$  and 13). (I) Plot of  $\alpha_1$ -A<sub>R</sub>-EPSC amplitudes measured at  $V_{\text{hold}} -65$  mV, in 2.5, 6.5, and 10.5 mM  $[\text{K}^+]_o$  ( $p = 0.162$ ,  $n = 17$ ). (J) Plot of  $\alpha_1$ -A<sub>R</sub>-EPSC reversal potential ( $E_{\text{rev}}$ ) with varying concentration of external  $\text{K}^+$  ( $[\text{K}^+]_o$ ), demonstrating a depolarizing shift in  $E_{\text{rev}}$  as external  $\text{K}^+$  was increased ( $p = 0.010$ ,  $n = 26$ , 10, and 11). (K) Plot of reversal potentials ( $E_{\text{rev}}$ ) of  $I_{\text{NA}}$ , demonstrating no significant difference between control conditions (ctrl), and after removal of external  $\text{Ca}^{2+}$  ( $0[\text{Ca}^{2+}]_o$ ,  $p = 0.49$ ,  $n = 14$  and 12) or  $\text{Mg}^{2+}$  ( $0[\text{Mg}^{2+}]_o$ ,  $p = 0.73$ ,  $n = 14$  and 11). (L) Plot of the amplitude of  $I_{\text{NA}}$  ( $V_{\text{hold}} -65$  mV) demonstrating an augmented  $I_{\text{NA}}$  amplitude in  $0[\text{Ca}^{2+}]_o$  ( $p = 0.017$ ,  $n = 14$ ), but not in  $0[\text{Mg}^{2+}]_o$  ( $p > 0.9999$ ,  $n = 11$ ) as compared with control conditions ( $n = 14$ ). Line and error bars represent mean  $\pm$  SEM, \* denotes statistical significance, ns denotes not significant.

The online version of this article includes the following source data and figure supplement(s) for figure 2:

**Source data 1.** Numerical data that were used to generate graphs in **Figure 2**.

**Figure supplement 1.** Tail current analysis reveals voltage-dependence of the  $\alpha_1$ -adrenergic receptor-dependent inward current.

(Figure 2B and C). The  $\alpha_1$ -A<sub>R</sub>-EPSC  $\sigma^2$  – amplitude relationship was well fit by linear regression, suggestive of a consistent conductance state, yielding an estimate of a -1.16 pA unitary current (Figure 2D). Voltage ramps from -120 to -10 mV (1 mV/10 ms) before and during the  $\alpha_1$ -A<sub>R</sub>-EPSC (Figure 2E) showed that the current reversed polarity at  $-28.6 \pm 2.4$  mV (Figure 2E-G). Exogenous application of noradrenaline (30  $\mu$ M, in the presence of  $\alpha_2$ -A<sub>R</sub> antagonist, idazoxan, 1  $\mu$ M) produced an inward current ( $I_{NA}$ ) with a similar reversal potential ( $-25.1 \pm 2.9$  mV, Figure 2G). Replacing extracellular Na<sup>+</sup> (126 mM) with N-methyl D-glucamine (NMDG) completely abolished inward  $I_{NA}$ , suggesting Na<sup>+</sup> as the prominent charge carrier (Figure 2H). Increasing extracellular K<sup>+</sup> from 2.5 to 6.5 or 10.5 mM, expected to shift  $E_k$  from -107 to -81 and -69 mV, respectively, had no effect on the amplitude of the  $\alpha_1$ -A<sub>R</sub>-EPSC at  $V_{hold}$  -65 mV (Figure 2I) nor -120 mV ( $p=0.692$ ,  $n=11$ , data not shown), but produced a significant depolarizing shift in  $E_{rev}$  of the  $\alpha_1$ -A<sub>R</sub>-EPSC (Figure 2J), suggesting the channel is also permeable to K<sup>+</sup>, and may be 2-3 $\times$  as permeable to K<sup>+</sup> as Na<sup>+</sup>. Removal of



**Figure 3.** NASPM blocks the  $\alpha_1$ -A<sub>R</sub>-EPSC and a tonic sodium inward current. (A) Example whole-cell voltage-clamp recording of the basal whole-cell current and the  $\alpha_1$ -A<sub>R</sub>-EPSC evoked every 90 s prior to, during, and after bath application of NASPM (NSP, 100  $\mu$ M). Time of stimulations are marked by arrows and peak of the  $\alpha_1$ -A<sub>R</sub>-EPSC are marked by asterisks. (B) NASPM completely eliminated the  $\alpha_1$ -A<sub>R</sub>-EPSC shown in representative traces (left, baseline adjusted) and in grouped data (right,  $p=0.001$ ,  $n=8$ ). (C) Bath application of NASPM produced an apparent outward current ( $p<0.0001$ ,  $n=21$ ). (D) Membrane noise (variance,  $\sigma^2$ ) decreased following NASPM (NSP,  $p<0.0001$ ,  $n=19$ ). (E) Time course of the inhibition of the  $\alpha_1$ -A<sub>R</sub>-EPSC amplitude (bottom) and apparent outward current (top) by application of NASPM. (F) Membrane resistance ( $R_m$ ,  $\Delta V$  -65 to -75 mV) increased during bath application of NASPM, indicating the apparent outward current was due to ion channels closing ( $p=0.004$ ,  $n=10$ ). (G) Current-voltage relationship of apparent outward current produced by NASPM ( $n=8$ ). Replacing 126 mM NaCl eliminated the apparent outward current ( $n=11$ ), suggesting a block of a tonic inward Na<sup>+</sup> current. Shaded area represents mean  $\pm$  SEM. (H) Plot of amplitude of NASPM-induced apparent outward current in stimulated and unstimulated brain slices demonstrating no effect of prior electrical stimulation ( $p=0.850$ ,  $n=21$  and 5). (I) Time course of the inhibition of the  $\alpha_1$ -A<sub>R</sub>-EPSC amplitude by application of NASPM, demonstrating identical block of the  $\alpha_1$ -A<sub>R</sub>-EPSC whether or not  $\alpha_1$ -A<sub>R</sub>-EPSCs were evoked during NASPM application. Line and error bars represent mean  $\pm$  SEM, ns indicates not significant, \* denotes statistical significance.

The online version of this article includes the following source data for figure 3:

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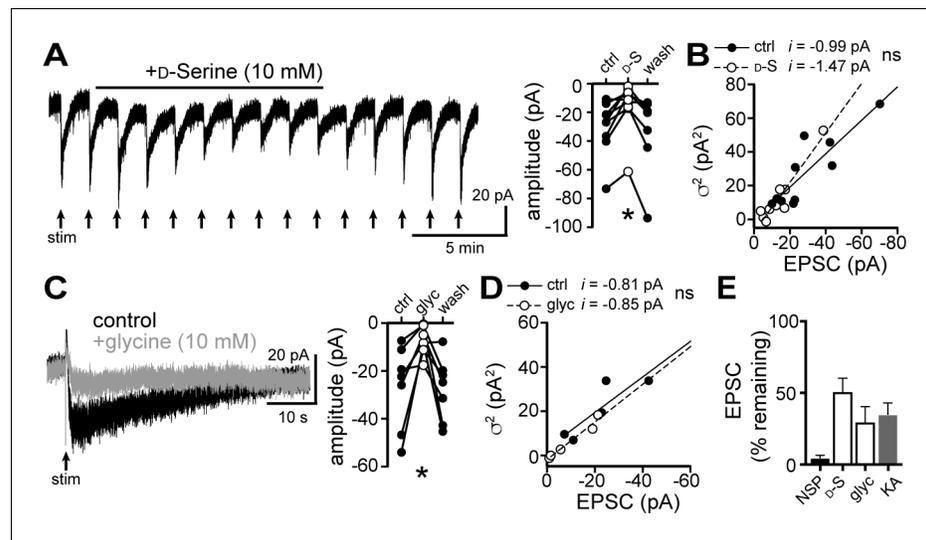
external  $\text{MgCl}_2$  had no significant effect on  $E_{\text{rev}}$  ( $-28.5 \pm 5.7$  mV), nor on the amplitude of  $I_{\text{NA}}$  (**Figure 2K and L**). Removal of external  $\text{CaCl}_2$  also had no effect on  $E_{\text{rev}}$  ( $-30.3 \pm 3.5$  mV) but significantly augmented inward  $I_{\text{NA}}$ , (**Figure 2K and L**). Taken together, the data suggest that  $\alpha 1\text{-A}_R$ -dependent current, whether produced by vesicular release of noradrenaline or exogenous noradrenaline application is carried through a mixed cation channel, with inward current carried predominantly by  $\text{Na}^+$  entry. Here, measurements of  $E_{\text{rev}}$  assume voltage-independence of the channel and the signaling mechanism by which  $\alpha 1\text{-A}_R$  signal to the channel. To test for voltage-dependence, we employed a two-pulse voltage-step protocol. Current was measured at  $V_{\text{hold}} -120$  mV following a conditioning pre-pulse ( $-120$  to  $30$  mV,  $150$  ms) before and after application of noradrenaline (**Figure 2—figure supplement 1A and B**).  $I_{\text{NA}}$  was isolated by subtracting the current under basal conditions from the current during noradrenaline. Conductance ( $G_{\text{NA}}$ ) was calculated, using an  $E_{\text{rev}}$  of  $-25.1$  mV. Conditioning depolarizing pre-pulses incrementally reduced  $G_{\text{NA}}$  and the increase in membrane noise induced by noradrenaline measured at  $V_{\text{hold}} -120$  mV (**Figure 2—figure supplement 1C and D**), demonstrating voltage-dependence of inward  $I_{\text{NA}}$ , such that depolarization reduced conductance.

### $\alpha 1\text{-adrenergic}$ receptors modulate tonic $\text{GluD1}_R$ -channel current

To assess involvement of  $\text{GluD1}_R$ -channels in carrying the  $\alpha 1\text{-A}_R$ -EPSC, we applied 1-Naphthyl acetyl spermine (NASPM), a synthetic analogue of Joro spider toxin that is an open-channel blocker of some other  $\text{Ca}^{2+}$ -permeable ionotropic glutamate receptors (*Blaschke et al., 1993; Guzmán et al., 2017; Koike et al., 1997*) and of  $\text{GluD}_R$ -channels (*Benamer et al., 2018; Kohda et al., 2000*). Application of NASPM ( $100 \mu\text{M}$ ,  $6$  min) blocked the  $\alpha 1\text{-A}_R$ -EPSC ( $96.0 \pm 12.5\%$  reduction), which recovered to baseline after a wash of  $>30$  mins (**Figures 3A, B, E and I**). NASPM also produced an apparent outward current ( $I_{\text{NSP}}$ ) of  $20.5 \pm 3.7$  pA with an  $E_{\text{rev}}$  of  $-31.4 \pm 4.8$  mV (**Figures 3A, C, E and G**) and a reduction in membrane noise (**Figure 3A and D**). After washout,  $I_{\text{NSP}}$  reversed with a similar time course of recovery of the  $\alpha 1\text{-A}_R$ -EPSC (**Figure 3E**).  $I_{\text{NSP}}$  was associated with an increase in  $R_m$  (**Figure 3F**) indicating a closure of channels. Replacing extracellular  $\text{Na}^+$  ( $126$  mM) with NMDG eliminated  $I_{\text{NSP}}$  (**Figure 3G**). Thus,  $I_{\text{NSP}}$  was due to block of tonic  $\text{Na}^+$ -dependent inward current.  $I_{\text{NSP}}$  was not dependent on prior electrical stimulation of the brain slice, as the magnitude of  $I_{\text{NSP}}$  was similar between stimulated and unstimulated brain slices (**Figure 3H**). Given that NASPM is an open-channel blocker (*Koike et al., 1997*), we tested whether electrically evoking an  $\alpha 1\text{-A}_R$ -EPSC during the application of NASPM was required for block. After obtaining a steady  $\alpha 1\text{-A}_R$ -EPSC baseline, NASPM was applied for  $6$  min without stimulating the brain slice. The  $\alpha 1\text{-A}_R$ -EPSC was blocked when stimulation was reapplied (**Figure 3I**), indicating that the channels underlying the  $\alpha 1\text{-A}_R$ -EPSC were already blocked. Thus, the  $\alpha 1\text{-A}_R$ -EPSC is mediated by channels that are at least transiently open at rest and may be the same channels underlie the apparent outward current induced by NASPM.

$\text{GluD}_R$ s bind the amino acids D-serine and glycine, both of which partially reduce constitutively open mutant and wild-type  $\text{GluD}_R$  channel current (*Ady et al., 2014; Benamer et al., 2018; Naur et al., 2007; Yadav et al., 2011*), likely by inducing a conformational change in the channel that resembles a desensitized state (*Hansen et al., 2009*). Application of D-serine ( $10$  mM,  $13.5$  min) reduced the amplitude of the  $\alpha 1\text{-A}_R$ -EPSC by  $49.7 \pm 9.6\%$  (**Figure 4A and E**), without affecting unitary channel current (**Figure 4B**). Application of glycine ( $10$  mM,  $4.5$  mins, in the presence of the glycine receptor antagonist, strychnine ( $10 \mu\text{M}$ ), also reduced the amplitude of the  $\alpha 1\text{-A}_R$ -EPSC by  $70.9 \pm 11.0\%$  (**Figure 4C and E**), without affecting unitary channel current (**Figure 4D**). Lastly, we found that application of the glutamate receptor antagonist kynurenic acid ( $1$  mM,  $10.5$  min) reduced the  $\alpha 1\text{-A}_R$ -EPSC amplitude by  $65.6 \pm 8.3\%$  (**Figure 4E**).

Next, a viral genetic strategy was used to functionally delete  $\text{GluD1}_R$ -channels by targeting the encoding gene, *Grid1*, via CRISPR/Cas9 (**Figure 5—figure supplement 1A–C**). In brief, one of two cocktails of AAV1 viruses were microinjected into the dorsal raphe of wild-type mice. The *Grid1* cocktail that targeted  $\text{GluD1}_R$ -channels included AAV1 viruses encoding Cas9, and mouse *Grid1* guide RNA with a nuclear envelope-embedded enhanced green fluorescent protein (eGFP) reporter. A separate cohort received a control cocktail of AAV1 viruses encoding Cas9 and eGFP reporter (control). Brain slices were prepared after  $>4$  weeks and the dorsal raphe was microdissected and frozen on dry ice to assess the mutation of *Grid1*. Restriction enzyme site-digested PCR confirmed *in vivo* mutation of *Grid1* at the expected site (**Figure 5—figure supplement 1D**). In separate *Grid1*



**Figure 4.** D-serine and glycine reduce the  $\alpha 1$ -AR-EPSC. (A) Bath application of D-serine (10 mM) reversibly reduced the  $\alpha 1$ -AR-EPSC, shown in a representative trace (left) and in grouped data (right,  $p=0.001$ ,  $n = 7$ ). (B) Plot of  $\alpha 1$ -AR-EPSC variance versus mean amplitude prior to (ctrl) and after reduction by D-serine (D-S), linear fit represents mean unitary current ( $i$ ), demonstrating no change in  $i$  with D-serine ( $p=0.165$ ,  $n = 10$  and 10). (C) Bath application of glycine (10 mM) reversibly reduced the  $\alpha 1$ -AR-EPSC, shown in representative traces (left, baseline adjusted) and in grouped data (right,  $p=0.015$ ,  $n = 7$ ). (D) Plot of  $\alpha 1$ -AR-EPSC variance versus mean amplitude prior to (ctrl) and after reduction by glycine (glyc), linear fit represents mean unitary current ( $i$ ), demonstrating no change in  $i$  with glycine ( $p=0.895$ ,  $n = 5$  and 5). (E) Summarized data of percent remaining in  $\alpha 1$ -AR-EPSC after NASPM (NSP, 100  $\mu$ M), D-serine (D-S, 10 mM), glycine (glyc, 10 mM), or kynurenic acid (KA, 1 mM). Line and error bars represent mean  $\pm$  SEM, ns indicates not significant, \* denotes statistical significance.

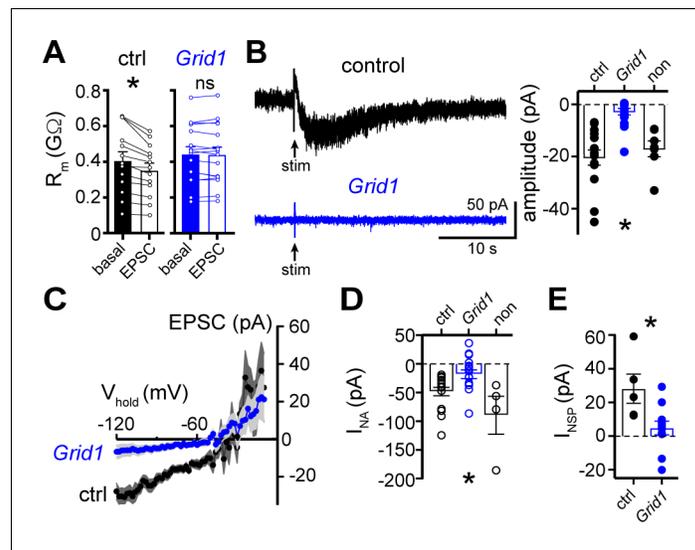
The online version of this article includes the following source data for figure 4:

**Source data 1.** Numerical data that were used to generate graphs in **Figure 4**.

and control cohorts, brain slices were prepared and whole-cell voltage-clamp recordings were made from transduced and non-transduced neurons visualized in brain slices by expression of eGFP. In eGFP<sup>+</sup> neurons from control mice, electrical stimulation produced a decrease in  $R_m$  and an  $\alpha 1$ -AR-EPSC, and bath application of noradrenaline caused inward  $I_{NA}$  (**Figure 5**). However, in eGFP<sup>+</sup> neurons from *Grid1* mice, electrical stimulation did not change  $R_m$  (**Figure 5A**) and no  $\alpha 1$ -AR-EPSC was detected above baseline noise (**Figure 5B and C**). In addition, inward  $I_{NA}$  was substantially smaller in eGFP<sup>+</sup> neurons from *Grid1* mice, as compared to eGFP<sup>+</sup> neurons from control mice (**Figure 5D**). In the same slices from *Grid1* mice, eGFP<sup>-</sup> neurons still had an  $\alpha 1$ -AR-EPSC and inward  $I_{NA}$  (**Figure 5B and D**). Lastly, bath application of NASPM produced an apparent outward current in eGFP<sup>+</sup> neurons from control mice, but not from *Grid1* mice (**Figure 5E**). Taken together, these results demonstrate that conduction through GluD1<sub>R</sub>-channels is necessary for the  $\alpha 1$ -AR-EPSC and the NASPM-sensitive tonic inward current.

## Functional deletion of GluD1<sub>R</sub>-channels in the dorsal raphe produces a behavioral phenotype

To assay a functional role of GluD1<sub>R</sub>-channels in dorsal raphe-related behavior, wild-type mice received a microinjection into the dorsal raphe of either *Grid1* or control virus cocktails. Behavioral assays were conducted >4 weeks post-injection, then the accuracy of the dorsal raphe injection and limited-spread of transduction was verified post-hoc by immunohistochemistry (**Figure 6A**). Basal locomotion was assayed in a dark arena. There was no difference between the two groups in the total distance traveled (**Figure 6B**) nor in the velocity of movements between control and *Grid1* mice ( $p=0.772$ ,  $n = 18$  and 16, data not shown). Next, mice were tested on an elevated plus maze in a well-lit room, an experimental assay of rodent anxiety behavior (**Walf and Frye, 2007**) known to involve both serotonergic and non-serotonergic neurons in the dorsal raphe (**Lawther et al., 2015**). *Grid1* mice spent less time in the open arms when compared to control mice (**Figure 6C–E**). Control



**Figure 5.** The  $\alpha_1$ -AR-EPSC is eliminated by targeting of GluD1<sub>R</sub>-channels via CRISPR/Cas9. (A) Membrane resistance ( $R_m$ ,  $\Delta V$   $-65$  to  $-120$  mV) decreased after stimulation in transduced neurons from mice injected with AAV-Cas9 and AAV-empty (ctrl,  $p=0.0002$ ,  $n = 13$ ), but not in transduced neurons from mice injected with AAV-Cas9 and AAV-*Grid1* (*Grid1*,  $p=0.562$ ,  $n = 16$ ). (B) Representative traces (left) and grouped data (right,  $p<0.0001$ ,  $n = 15$  and  $16$  and  $7$ ) demonstrating the presence of an  $\alpha_1$ -AR-EPSC in transduced neurons from control mice, but not from *Grid1* mice. Neighboring non-transduced neurons from mice injected with AAV-Cas9 and AAV-*Grid1* (non) had an  $\alpha_1$ -AR-EPSC that was indistinguishable from transduced neurons from control mice ( $p>0.999$ ). (C) Current-voltage relationship of the  $\alpha_1$ -AR-EPSC from control ( $n = 13$ ) and *Grid1* ( $n = 16$ ) grouped data. Shaded area represents mean  $\pm$  SEM. (D) Targeting GluD1<sub>R</sub> reduced the inward current to noradrenaline (NA,  $I_{NA}$ ,  $30 \mu\text{M}$ ) as compared to transduced neurons from control mice and neighboring non-transduced neurons ( $p=0.004$ ,  $n = 16$  and  $16$  and  $4$ ). Inward  $I_{NA}$  in non-transduced neurons from mice injected with AAV-Cas9 and AAV-*Grid1* was similar to transduced neurons from control mice ( $p=0.631$ ). (E) Targeting GluD1<sub>R</sub> reduced the tonic inward current revealed by bath application of NASPM ( $100 \mu\text{M}$ ,  $I_{NSP}$ ) as compared to transduced neurons from control mice ( $p=0.009$ ,  $n = 5$  and  $11$ ). Line and error bars represent mean  $\pm$  SEM, \* denotes statistical significance, ns denotes not significant.

The online version of this article includes the following source data and figure supplement(s) for figure 5:

**Source data 1.** Numerical data that were used to generate graphs in **Figure 5**.

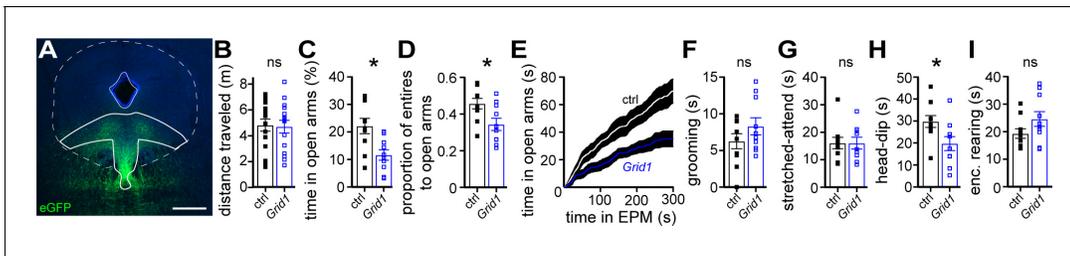
**Figure supplement 1.** Design and testing of guide RNA targeting mouse *Grid1*.

and *Grid1* mice made a similar total number of entries to either open or enclosed arms (control:  $39.4 \pm 2.0$ ; *Grid1*:  $36.0 \pm 2.3$ ,  $p=0.697$ ), but *Grid1* mice made proportionally fewer entries to the open arms (**Figure 6D**). Time spent grooming or in stretched-attend postures were similar between control and *Grid1* mice (**Figure 6F and G**). Since movement in the elevated plus maze reflects conflict between innate drive to explore of a novel environment and natural avoidance of open spaces (**Walf and Frye, 2007**), we also examined exploratory behaviors. *Grid1* mice spent less time lowering their head over the edge of the open arms than control mice (head-dipping, **Figure 6H**), suggestive of decreased exploratory behavior. However, *Grid1* mice spent a similar amount of time rearing in the enclosed arms compared to control mice (**Figure 6I**) suggesting innate exploratory drive in the enclosed arms was intact. Taken together, these results are indicative of heightened anxiety after functional deletion of GluD1<sub>R</sub>-channels in the dorsal raphe.

## Discussion

### Physiological relevance of GluD1<sub>R</sub>-channels to dorsal raphe function

In vivo, 5-HT neurons in the dorsal raphe require noradrenaline release and subsequent activation of  $\alpha_1$ -ARs to maintain persistent action potential firing (**Baraban and Aghajanian, 1980**). The activation



**Figure 6.** Loss of functional GluD1 receptors in the dorsal raphe produces an anxiogenic behavioral phenotype in mice. (A) Example maximum intensity projection confocal image of spread of viral transduction following dorsal raphe microinjection of AAV-Cas9 and AAV-*Grid1* using an eGFP reporter; scale bars, 0.5 mm. Image was registered and aligned with plate 69 (Franklin and Paxinos mouse brain atlas) with dorsal raphe outlined in solid white. (B) Plot of distance traveled in 30 mins in a dark arena, demonstrating no difference in horizontal locomotion between mice with dorsal raphe microinjections with AAV-Cas9 and AAV-empty (ctrl) versus with AAV-Cas9 and AAV-*Grid1* ( $p=0.762$ ,  $n = 18$  and  $15$ ). (C) In an elevated plus maze, *Grid1* mice spent less time in the open arms as compared with control-transduced mice ( $p=0.007$ ,  $n = 10$  and  $10$ ). (D) *Grid1* mice made proportionally fewer entries to the open arms compared to control-transduced mice ( $p=0.033$ ,  $n = 10$  and  $10$ ). (E) Plot of cumulative time spent in the open arms of an elevated plus maze (EPM). (F) Plot of time spent grooming, demonstrating no difference between control-transduced and *Grid1* mice ( $p=0.481$ ,  $n = 10$  and  $10$ ). (G) Plot of time spent in stretched-attend posture, demonstrating no difference between control-transduced and *Grid1* mice ( $p=0.968$ ,  $n = 10$  and  $9$ ). (H) *Grid1* mice spent less time looking over the edge of the open arms (head-dip) than control-transduced mice ( $p=0.018$ ,  $n = 10$  and  $10$ ). (I) Plot of time spent rearing in the enclosed arms, demonstrating no difference between control-transduced and *Grid1* mice ( $p=0.143$ ,  $n = 10$  and  $10$ ). Line and error bars represent mean  $\pm$  SEM,  $n =$  number of mice, \* denotes statistical significance, ns denotes not significant.

The online version of this article includes the following source data for figure 6:

**Source data 1.** Numerical data that were used to generate graphs in **Figure 6**.

of  $\alpha 1$ -ARs in the dorsal raphe by exogenous agonist was thought to depolarize neurons through net reduction of  $K^+$  conductance, transiently activating calcium-activated  $K^+$  current while persistently decreasing another  $K^+$  current, and by activation of an unidentified non- $K^+$  conductance (Pan et al., 1994). In a more recent study in the dorsal raphe, Brown et al. (2002) reported that activation of  $\alpha 1$ -ARs, induces  $Na^+$ -dependent inward current with an  $E_{rev}$  of  $-23$  mV, similar to our findings. Our study identifies GluD1 receptor-channels as the ion channel that carries this mixed cation current, indicating that modulation of GluD1<sub>R</sub>-channels is a key constituent in driving persistent action potential firing of the 5-HT neurons. In principle, inward GluD1<sub>R</sub>-channel current may bring the membrane potential to threshold, but recruitment of other voltage-gated ion channels is expected to underlie the persistent pacemaker-like activity. Intriguingly, Brown et al. (2002) demonstrated that activation of  $G_q$  protein-coupled histamine  $H_1$  and orexin  $OX_2$  receptors also produced an inward current that was occluded by the  $\alpha 1$ -AR-dependent current. Whether these receptors, and other  $G_q$ PCRs, augment GluD1<sub>R</sub>-channel current remains to be determined.

Dysregulation of the 5-HT signaling neuropsychiatric disorders is well-established. Pharmacotherapies to boost serotonin signaling are common and often efficacious in some of these conditions. Genetic association studies have identified *GRID1* as a susceptibility gene for psychiatric conditions, including schizophrenia, major depressive disorder, bipolar disorder, autism spectrum disorder, and alcohol dependence (Edwards et al., 2012; Fallin et al., 2005; Griswold et al., 2012). Global *Grid1* knock-out mice display abnormal social behaviors, including heightened aggression and decreased social interaction, as well as altered emotional behaviors (Yadav et al., 2012) that are analogous to features of neuropsychiatric conditions in humans. Our study found that functional deletion of GluD1<sub>R</sub>-channels, specifically in the dorsal raphe, produces a heightened anxiety-like response in the elevated plus maze without changing basal locomotion and exploratory behaviors in non-threatening environments. Previous studies have demonstrated that both 5-HT and non-5-HT/GABAergic dorsal raphe neurons are activated by aversive, anxiety or fear-producing stimuli (Seo et al., 2019; Silveira et al., 1993), with regional subpopulation specificity (Grahm et al., 1999; Grahm et al., 2019; Lawther et al., 2015). Our viral strategy functionally deleted GluD1<sub>R</sub>-channels in a non-specific manner, targeting all dorsal raphe neurons, including 5-HT and GABAergic neurons. Given the rich diversity of dorsal raphe neuron subtypes and subdivisions within the 5-HT neurons (Huang et al., 2019; Luo et al., 2015; Ren et al., 2018), future work will be needed to parse the behavioral role of GluD1<sub>R</sub>-channels with subnuclei/subpopulation specificity.

## Metabotropic-ionotropic receptor crosstalk modulates ion channel function of GluD1<sub>R</sub>

GluD<sub>R</sub> have been characterized as scaffold proteins or synaptic organizers, regulating LTD, endocytosis and trafficking of AMPA<sub>R</sub>, formation of excitatory and inhibitory synapses, and spine density, independent of ion conduction through the pore (Fossati et al., 2019; Hirai et al., 2003; Schmid and Hollmann, 2008; Tao et al., 2018). Similarly, NMDA<sub>R</sub> are known to signal through non-ionotropic or 'metabotropic' mechanisms where ion conduction is not required, to regulate LTD, AMPA<sub>R</sub> endocytosis, and spine morphology (Dore et al., 2016). The ability of GluD<sub>R</sub>-channels to carry ionic current does not conflict with its known role as a synaptic organizer, but rather expands the similarities between NMDA<sub>R</sub> and GluD<sub>R</sub>.

The largest obstacle in advancing the understanding of the ionotropic nature of GluD<sub>R</sub> is the lack of known agonist and inability to gate the intact channel. The majority of studies have been performed on constitutively open mutant or chimeric channels. In domain-swapped chimeric channels, agonist binding to the ligand-binding domain (LBD) of AMPA<sub>R</sub> or Kainate<sub>R</sub> opens the GluD<sub>R</sub>-channel pore and generates a substantial current, but the LBD of GluD<sub>R</sub> on the pore region of AMPA<sub>R</sub> or Kainate<sub>R</sub>-channels fails to generate current (Orth et al., 2013; Schmid et al., 2009). Two prior studies have demonstrated that in heterologous systems and brain slices, activation of metabotropic glutamate receptors (mGlu<sub>R</sub>) produces an inward current carried by GluD1<sub>R</sub>- (Benamer et al., 2018) or GluD2<sub>R</sub>-channels (Ady et al., 2014), concluding that mGlu<sub>R</sub> activation triggers gating of GluD<sub>R</sub> channels. The congruous explanation of our results is that, in dorsal raphe neurons, GluD1<sub>R</sub>-channels are functional and open under basal conditions, carrying subthreshold, tonic Na<sup>+</sup> current. Activation of α1-A<sub>R</sub>s, by exogenous agonist or synaptic release of noradrenaline modulates gating of GluD1<sub>R</sub>-channels and excites dorsal raphe neurons by increasing tonic GluD1<sub>R</sub>-channel inward current.

In general, the kinetics of iGlu<sub>R</sub> synaptic currents are controlled by the lifetime of the receptor-agonist complex and the rate of desensitization and deactivation. The presence of ambient levels of glutamate and glycine along with slow desensitization activate NMDA<sub>R</sub> to produce a tonic inward current (Sah et al., 1989). Our results demonstrate that GluD1<sub>R</sub> are functional ion channels, but whether they function as ligand-gated receptor-channels that open in response to a chemical signal, is not yet determined. What remains to be understood are the conditions that permit GluD1<sub>R</sub>-channel opening and why their activation has been largely elusive in heterologous expression systems. Reminiscent of times before the discovery of glycine as a necessary co-agonist at NMDA<sub>R</sub> (Johnson and Ascher, 1987; Kleckner and Dingledine, 1988), it may be that an endogenous agonist needed for gating is present in brain slices. Alternatively, it is possible GluD1<sub>R</sub>-channels are gated by an intracellular factor or require expression of accessory or interacting protein (Tomita, 2010). Tonic activation of α1-A<sub>R</sub>s cannot explain the tonic inward current as α1-A<sub>R</sub> antagonism did not change basal whole-cell current.

The mechanism by which α1-A<sub>R</sub>s increase GluD1<sub>R</sub>-channel current also remains to be described and may be distinct from the tonic activation. It is well-established that G<sub>q</sub>PCRs, especially mGlu<sub>R</sub> and mACh<sub>R</sub>, bidirectionally change NMDA<sub>R</sub> and AMPA<sub>R</sub> ionic currents, producing the two major forms of synaptic plasticity, long-term potentiation (LTP) and long-term depression (LTD), in part through a variety of distinct postsynaptic mechanisms (Hunt and Castillo, 2012). To our knowledge, the duration of the α1-A<sub>R</sub>-EPSC (~27 s) is exceptional for any known synaptic current and more closely resembles the duration of short-term synaptic plasticity; for instance, endocannabinoid-mediated short-term depression (Lu and Mackie, 2016). Canonically, G<sub>q</sub>PCRs activate phospholipase C which hydrolyzes the integral membrane lipid phosphatidylinositol 4,5-bisphosphate (PIP<sub>2</sub>) into inositol triphosphate (IP<sub>3</sub>) and diacylglycerol. PIP<sub>2</sub> stabilizes Kv7 channels such that PIP<sub>2</sub> hydrolysis following mACh<sub>R</sub> activation accounts for inhibition of M-current (Suh and Hille, 2002). In contrast, PIP<sub>2</sub> inhibits TRPV4 channels, such that G<sub>q</sub>PCR-dependent PIP<sub>2</sub> depletion allows for TRPV4 channels to open (Harraz et al., 2018). By the same signaling cascade, G<sub>q</sub>PCRs stimulate the production of the endocannabinoid, 2-AG, that can act directly on ion channels in the membrane (Gantz and Bean, 2017). Thus, one possibility is that α1-A<sub>R</sub>s modulate GluD1<sub>R</sub>-channels through membrane lipid signaling, involving PIP<sub>2</sub>, diacylglycerol, or 2-AG, as it can take tens-of-seconds to minutes for ion channels to recover from modulation by membrane lipids (Gantz and Bean, 2017; Suh and Hille, 2002). Alternatively, there may be direct modulation of GluD1<sub>R</sub>-channels by G protein subunits or activation of protein kinase signaling cascades. The inclusion of the calcium-chelator BAPTA in the internal

recording solution makes it unlikely that  $\alpha 1$ -ARs modulate GluD1<sub>R</sub>-channels via IP<sub>3</sub> and calcium release from intracellular stores (Hoesch *et al.*, 2004). Largely, it remains to be seen whether these intracellular signaling cascades, many of which are known to affect NMDA<sub>R</sub>- and AMPA<sub>R</sub>-channels, modify GluD<sub>R</sub>-channels.

In heterologous systems and constitutively open mutant GluD<sub>R</sub>-channels, the current reverses polarity around 0 mV (Zuo *et al.*, 1997), akin to AMPA<sub>R</sub>- and NMDAR-channels, while our results show E<sub>rev</sub> of ~ -30 mV. While slow voltage-ramps were employed to minimize space-clamp error, we cannot rule out that some of the difference may be attributed to space-clamp error in brain slices, especially since the magnitude of subtracted current is small relative to total membrane current at depolarized potentials. However, there are many reports of inward currents produced by activation of many different G<sub>q</sub>PCRs with reversal potentials between -40 and -23 mV (Awad *et al.*, 2000; Brown *et al.*, 2002; Yamada-Hanff and Bean, 2013) under different recording conditions; a commonality that is unlikely to be accounted for by space-clamp error alone. Tail current analysis revealed voltage-dependence of I<sub>NA</sub>, such that depolarization reduced conductance. These data may reflect block of GluD1<sub>R</sub>-channels by endogenous intracellular polyamines, as established for calcium-permeable AMPA<sub>R</sub>- and Kainate<sub>R</sub>-channels (Bowie and Mayer, 1995). Another important consideration is that our measurements may be subject to voltage-dependence of the signaling pathway between  $\alpha 1$ -ARs and GluD1<sub>R</sub>-channels. Taken together, measurements here should be considered an estimate of GluD1<sub>R</sub>-channels, and more precisely as the current-voltage relationship of the  $\alpha 1$ -ARs-GluD1<sub>R</sub>-channel signaling complex.

## Summary

In summary, the  $\alpha 1$ -AR-mediated depolarization of dorsal raphe neurons that drives action potential firing *in vivo* is carried by the mixed cation channel, GluD1<sub>R</sub>. Thus in addition to their role as a scaffold protein, GluD1<sub>R</sub> are functional ion channels that critically regulate neuronal excitability. Many of the biophysical properties of the GluD1<sub>R</sub>-channel are like other members of the ionotropic glutamate receptor family. Given the widespread distribution of these receptors throughout the brain (Hepp *et al.*, 2015), ion channel function of GluD1<sub>R</sub> may be prevalent and relevant to neuronal excitability and circuit function in different parts of the throughout the nervous system. This study lays the foundation to investigate the ion channel function of GluD1<sub>R</sub> in excitatory G<sub>q</sub>PCR-dependent synaptic transmission and regulation of neuronal excitability, expanding upon the wealth of knowledge of pharmacology and regulatory elements established for NMDA<sub>R</sub> and AMPA<sub>R</sub> signaling.

## Materials and methods

### Key resources table

Reagent type (species) or resource	Designation	Source or reference	Identifiers	Additional information
Strain, strain background ( <i>Mus musculus</i> )	C57BL/6J	The Jackson Laboratory	Stock# 000664 RRID:IMSR_JAX:000664	males and females
Strain, strain background ( <i>E. coli</i> )	NEB Stable	New England Biolabs	Cat# C3040H	-
Genetic reagent (adeno-associated virus)	AAV-Cas9	PMIDs:25326897 30792150	NIDA IRP Core Facility, AAV1, pX551, RRID:Addgene_60957	Lot# AAV692
Genetic reagent (adeno-associated virus)	AAV-Grid1	This paper	NIDA IRP Core Facility, AAV1, pOTTTC1706, Addgene 131683	Lot# AAV732
Genetic reagent (adeno-associated virus)	AAV-empty	This paper	NIDA IRP Core Facility, AAV1, pOTTTC1553, Addgene 131682	Lot# AAV746

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Reagent type (species) or resource	Designation	Source or reference	Identifiers	Additional information
Chemical compound, drug	Bgl I restriction enzyme	New England Biolabs	Cat# R0143S	-
Chemical compound, drug	D-serine	Millipore Sigma	Cat# S4250	10 mM
Chemical compound, drug	GDP $\beta$ S-Li <sub>3</sub>	Millipore Sigma	Cat# G7637	1.8 mM
Chemical compound, drug	Glycine	Millipore Sigma	Cat# G7126	10 mM
Chemical compound, drug	Idazoxan	Millipore Sigma	Cat# I6138	1 $\mu$ M
Chemical compound, drug	Kynurenic acid	Millipore Sigma	Cat# K3375	1 mM
Chemical compound, drug	MK-801	Tocris Bioscience	Cat #0924	5 $\mu$ M
Chemical compound, drug	NASPM	Tocris Bioscience	Cat #2766	100 $\mu$ M
Chemical compound, drug	NBQX	Tocris Bioscience	Cat #1044	3 $\mu$ M
Chemical compound, drug	NMDG	Millipore Sigma	Cat# 66930	126 mM
Chemical compound, drug	Noradrenaline	Tocris Bioscience	Cat #5169	30 $\mu$ M
Chemical compound, drug	Picrotoxin	Tocris Bioscience	Cat #1128	100 $\mu$ M
Chemical compound, drug	Prazosin	Millipore Sigma	Cat# P7791	100 nM
Chemical compound, drug	Reserpine	Millipore Sigma	Cat# R0875	1 $\mu$ M
Chemical compound, drug	Strychnine	Millipore Sigma	Cat# S8753	10 $\mu$ M
Chemical compound, drug	Tetrodotoxin	Tocris Bioscience	Cat# 1069	1 $\mu$ M
Chemical compound, drug	WAY-100635	Tocris Bioscience	Cat# 4380	300 nM
Sequence-based reagent	Forward amplification primer	IDTDNA	tgattacgccaagctt GGTGGAGCTGTG TGGATGAAGC	-

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Reagent type (species) or resource	Designation	Source or reference	Identifiers	Additional information
Sequence-based reagent	Forward sequence primer	IDTDNA	CCAGCCTGTGACCTCATGACC	-
Sequence-based reagent	Reverse amplification primer	IDTDNA	gacggccagtgaattc CTTCAGCTGTCATG ATAAGGTGATGTTG	-
Commercial assay, kit	In-Fusion HD Cloning	Takara Bio Clontech	Cat# 639647	-
Software, algorithm	Clampfit 10.7	Axon Instruments	RRID:SCR_011323	<a href="https://www.moleculardevices.com/products/axon-patch-clamp-system">https://www.moleculardevices.com/products/axon-patch-clamp-system</a>
Software, algorithm	EthoVision XT	Noldus Information Technology	RRID:SCR_000441	<a href="https://www.noldus.com/ethovision-xt">https://www.noldus.com/ethovision-xt</a>
Software, algorithm	Fiji	PMID:22743772	RRID:SCR_002285	<a href="http://fiji.sc">http://fiji.sc</a>
Software, algorithm	Prism 8	GraphPad	RRID:SCR_002798	<a href="http://www.graphpad.com">http://www.graphpad.com</a>
Software, algorithm	VersaMax Analyzer	Omnitech-electronics, Inc	<a href="http://www.omnitech-electronics.com/product/VersaMax-Legacy-Open-Field-Locomotor-Activity/1930">http://www.omnitech-electronics.com/product/VersaMax-Legacy-Open-Field-Locomotor-Activity/1930</a>	-

## Animals

All studies were conducted in accordance with the National Institutes of Health Guide for the Care and Use of Laboratory animals with the approval of the National Institute on Drug Abuse Animal Care and Use Committee. Wild-type C57BL/6J (>3 months old) mice of either sex were used. Mice were group-housed on a 12:12 hr reverse light cycle.

## Brain slice preparation and electrophysiological recordings

The methods for brain slice preparation and electrophysiological recordings were almost identical to previous reports in the dorsal raphe (*Gantz et al., 2015a*) and ventral midbrain (*Gantz et al., 2015b*). In brief, mice were deeply anesthetized with isoflurane and killed by decapitation. Brains were removed quickly and placed in warmed artificial cerebral spinal fluid (modified Krebs' buffer) containing (in mM): 126 NaCl, 2.5 KCl, 1.2 MgCl<sub>2</sub>, 1.2 CaCl<sub>2</sub>, 1.2 NaH<sub>2</sub>PO<sub>4</sub>, 21.5 NaHCO<sub>3</sub>, and 11 D-glucose with 5 μM MK-801 to reduce excitotoxicity and increase viability, bubbled with 95/5% O<sub>2</sub>/CO<sub>2</sub>. In the same solution, coronal dorsal raphe slices (220 μm) were obtained using a vibrating microtome (Leica 1220S) and incubated at 32°C > 30 min prior to recording.

Slices were then mounted in a recording chamber and perfused ~3 mL/min with ~35°C modified Krebs' buffer. Electrophysiological recordings were made with a Multiclamp 700B amplifier (Molecular Devices), Digidata 1440A A/D converter (Molecular Devices), and Clampex 10.4 software (Molecular Devices) with borosilicate glass electrodes (King Precision Glass) wrapped with Parafilm to reduce pipette capacitance (*Gantz and Bean, 2017*). Pipette resistances were 1.8–2.8 MΩ when filled with an internal solution containing, (in mM) 104.56 K-methylsulfate, 5.30 NaCl, 4.06 MgCl<sub>2</sub>, 4.06 CaCl<sub>2</sub>, 7.07 HEPES (K), 3.25 BAPTA(K4), 0.26 GTP (sodium salt), 4.87 ATP (sodium salt), 4.59 creatine phosphate (sodium salt), pH 7.32 with KOH, mOsm ~285, for whole-cell patch-clamp recordings. Series resistance was monitored throughout the experiment. Transmitter release was evoked by trains of electrical stimuli delivered via a Krebs' buffer-filled monopolar stimulating electrode positioned in the dorsal raphe, within 200 μm of the recorded neuron (*Gantz et al., 2015a*). Cell-attached recordings were made from quiescent neurons in slice, using pipettes filled with modified Krebs' buffer. For experiments involving viral microinjections, transduced neurons were identified in the slice by visualization of eGFP. Reported voltages are corrected for a liquid junction

potential of  $-8$  mV between the internal solution and external solution. All drugs were applied by bath application. All experiments were conducted following incubation in an NMDA<sub>R</sub> channel blocker (MK-801, 5  $\mu$ M, >1 hr), and then with AMPA<sub>R</sub> and Kainate<sub>R</sub> (NBQX, 3  $\mu$ M), GABA-A<sub>R</sub> (picrotoxin, 100  $\mu$ M), and 5-HT1A<sub>R</sub> (WAY-100635, 300 nM) antagonists in the external solution. In addition, a  $\alpha$ 2-adrenergic receptor antagonist (idazoxan, 1  $\mu$ M) was added for experiments where noradrenaline was applied and a glycine receptor antagonist (strychnine, 10  $\mu$ M) was added when glycine was applied. Unitary current was calculated from fluctuation analysis, as previously described (Bean *et al.*, 1990), assuming the macroscopic current arises from independent, identical channels with a low probability of opening, according probability theory;  $i = \sigma^2/[l(1-p)]$  where  $i$  is unitary current,  $\sigma^2$  is the variance,  $l$  is mean current amplitude, and  $p$  is probability of opening.

### Vector construction gRNA identification

CRISPR SpCas9 gRNA target sites were identified in the mouse *Grid1* gene (NC\_000080.6) using CRISPOR (Haeussler *et al.*, 2016). The seed sequence (GAACCCTAGCCCTGACGGCG) was chosen based on its relatively high specificity scores and the observation that it contains a Bgl I restriction enzyme site (GCCNNNNNGGC) that overlaps with the Cas9 cleavage site.

### Mouse *Grid1* genotyping

C57BL/6J mouse genomic DNA was isolated from tail biopsies or brain pieces containing microdissected dorsal raphe by digestion in DNA lysis buffer (50 mM KCl, 50 mM Tris-HCl (pH 8.0), 2.5 mM EDTA, 0.45% NP-40, 0.45% Tween-20, 0.5  $\mu$ g/mL proteinase K) for 3 hr at 55°C, and 1 hr at 65°C. Lysates were then used as templates to amplify a 654 basepair fragment including the 390F gRNA target site using Q5 HotStart Master mix (New England Biolabs). A portion of the finished PCR reaction was treated with Bgl I restriction enzyme (New England Biolabs) for 60 min and processed on an AAT1 fragment analyzer.

### Construction and packaging of AAV vectors

The AAV vector plasmid encoding SpCas9 (Swiech *et al.*, 2015) (pX551) expressed from the *Mecp2* promoter was a gift from Feng Zhang (Addgene plasmid # 60957, AAV-Cas9). The AAV packaging plasmid encoding a nuclear envelope-embedded eGFP reporter (Addgene 131682) was constructed by amplifying the KASH domain from (Addgene 60231, a gift of Feng Zhang) and fusing it (in-frame) to the end of coding region for eGFP in (Addgene 60058, pOTTC407) using ligation-independent cloning (AAV-empty, Figure 5—figure supplement 1A). gRNA was cloned into a mU6 expression cassette and then moved into an AAV backbone expressing a nuclear envelope-embedded (KASH-tagged) eGFP reporter (Addgene 131683) by PCR amplification and ligation-independent cloning (AAV-*Grid1*). Insert-containing clones were verified by sequencing and restriction fragment analysis prior to virus production. All AAV vectors were produced using triple transfection method as previously described (Howard and Harvey, 2017). All vectors were produced using serotype 1 capsid proteins and titered by droplet digital PCR.

### Stereotaxic intracranial microinjections

Mice were anesthetized with a cocktail of ketamine/xylazine, immobilized in a stereotaxic frame (David Kopf Instruments), and received one midline injection of a 1:1 (v/v) cocktail of viruses AAV-Cas9 and AAV-empty or AAV-*Grid1* for total volume 400 nL delivered over 4 min. The coordinates for injection were AP  $-4.4$ ; ML 1.19, 20° angle; DV  $-3.62$  mm, with respect to bregma. Prior to surgery, mice were injected subcutaneously with warm saline (0.5 mL) to replace fluid lost during surgery and given carprofen (5 mg/kg) post-surgery for pain relief. Mice recovered for >4 weeks to allow expression.

### Behavioral assays

Behavioral assays were conducted during the dark cycle, using 3 separate cohorts of AAV-Cas9 and AAV-empty or AAV-*Grid1*-injected mice as biological replicates, 30–55 d post-injection. To measure basal locomotion, mice were placed in locomotor boxes (VersaMax System, Omnitech Electronics, Inc) in a dark room for 1 hr, following prior habituation to the locomotor boxes for >2 d (1 hr/d). VersaMax Analyzer software was used to determine the total distance traveled, time spent moving, and

velocity of movement in the last 30 min of the session. The boxes were cleaned with 70% ethanol and allowed to dry between trials. The elevated-plus maze was used to assay anxiety-related behaviors (Walf and Frye, 2007). The apparatus (Med associates, Inc) was placed 30 cm above the floor and consisted of two plastic light gray open arms (30 × 5 cm) and two black enclosed arms (30 × 5 cm) extending from a central platform (5 × 5 × 5 cm) at 90 degrees. Following habituation to the brightly lit room, mice were placed individually in the center of the maze, facing an open arm. Video tracking EthoVision XT software (Noldus Information Technology) was used to track mouse location, total distance traveled, velocity of movement, body elongation, and entries and time spent into the open and enclosed arms for each 5 min trial. Duration of head-dips, grooming, and enclosed-arm rearing were scored manually from videos played a 0.5x speed. Rearing in the enclosed arm was often associated with pressing one or both forepaws to the wall. Stretched-attend postures was defined by body elongation (70% threshold) and movement velocity <1 cm/s. No mice fell or jumped from the maze and open-arm rearing was not observed. The maze was cleaned with 70% ethanol after every trial and allowed to dry before the next trial. Mice were excluded from analysis if there was limited or no expression in the dorsal raphe, or if expression spread rostrally to the ventral tegmental area or caudally to locus coeruleus.

### Immunohistochemistry and confocal microscopy

Following behavioral assays, mice were euthanized with Euthasol and transcardially perfused with PBS followed by ice-cold 4% paraformaldehyde in PBS (pH 7.4). Brains were fixed overnight at 4 C and then sliced coronally in 50 μm sections. Alternatively, mice were anesthetized with isoflurane and euthanized by decapitation. Brains were removed and slices were prepared as for brain slice electrophysiology (220 μm), then fixed in room temperature 4% paraformaldehyde in PBS for 1 hr. Slices were mounted with Fluoromount-G with DAPI (Invitrogen) aqueous mounting medium. Confocal images were collected on an Olympus microscope (4x, 0.16 NA) and processed using Fiji.

### Data analysis and visualization

Data were analyzed using Clampfit 10.7. Data are presented as representative traces, or in scatter plots where each point is an individual cell, and bar graphs with means ± SEM. In traces with electrical stimulation, stimulation artifacts have been blanked for clarity. Unless otherwise noted,  $n$  = number of distinct cells or mice as biological replicates. No sample was tested in the same experiment more than once (technical replication).  $E_{revs}$  were determined by linear regression for each cell. Recordings in which current did not cross 0 pA were omitted from analysis. To minimize space-clamp errors, analysis of current during voltage ramps was limited to −10 mV where the currents were typically less than 500 pA. Ramp currents were averaged in 2 mV bins (20 ms). Data sets with  $n > 30$  were tested for normality with a Shapiro-Wilk test. When possible (within-group comparisons), significant differences were determined for two group comparisons by paired t-tests, Wilcoxon matched-pairs signed rank test, and in more than two group comparisons by nonparametric repeated-measures ANOVA (Friedman test). Significant mean differences in between-group comparisons were determined for two group comparisons by Mann Whitney tests, and in more than two group comparisons by Kruskal-Wallis tests. ANOVAs were followed, when  $p < 0.05$  by Dunn's multiple comparisons post hoc test. Linear trends were analyzed using a mixed model ANOVA. A difference of  $p < 0.05$  was considered significant. For behavioral assays, Grubbs test was used to identify outliers. Basal locomotion and time spent in stretched-attend posture from one *Grid1* mouse each were found to be outliers and were excluded from group comparisons. Exact values are reported unless  $p < 0.0001$  or  $> 0.999$ . Statistical analysis was performed using GraphPad Prism 8 (GraphPad Software, Inc).

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### Author contributions

Stephanie C Gantz, Conceptualization, Data curation, Formal analysis, Supervision, Funding acquisition, Validation, Investigation, Visualization, Methodology, Writing - original draft, Project administration, Writing - review and editing; Khaled Moussawi, Formal analysis, Validation, Visualization, Methodology, Writing - review and editing; Holly S Hake, Data curation, Investigation, Writing - review and editing

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### Ethics

Animal experimentation: This study was conducted in accordance with the National Institutes of Health Guide for the Care and Use of Laboratory animals. The protocol was approved by the National Institute on Drug Abuse Animal Care and Use Committee.

### Decision letter and Author response

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## Additional files

### Supplementary files

- Transparent reporting form

### Data availability

All data analysed for this study are included in the manuscript.

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