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## Cerebellum-Specific Deletion of the GABA<sub>A</sub> Receptor $\delta$ Subunit Leads to Sex-Specific Disruption of Behavior

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### SUMMARY

Granule cells (GCs) of the cerebellar input layer express high-affinity  $\delta$  GABA<sub>A</sub> subunit-containing GABA<sub>A</sub> receptors ( $\delta$ GABA<sub>A</sub>Rs) that respond to ambient GABA levels and context-dependent neuromodulators like steroids. We find that GC-specific deletion of  $\delta$ GABA<sub>A</sub> (cerebellar [cb]  $\delta$  knockout [KO]) decreases tonic inhibition, makes GCs hyperexcitable, and in turn, leads to differential activation of cb output regions as well as many cortical and subcortical brain areas involved in cognition, anxiety-like behaviors, and the stress response. Cb  $\delta$  KO mice display deficits in many behaviors, but motor function is normal. Strikingly,  $\delta$ GABA<sub>A</sub> deletion alters maternal behavior as well as spontaneous, stress-related, and social behaviors specifically in females. Our findings establish that  $\delta$ GABA<sub>A</sub>Rs enable the cerebellum to control diverse behaviors not previously associated with the cerebellum in a sex-dependent manner. These insights may contribute to a better understanding of the mechanisms that underlie behavioral abnormalities in psychiatric and neurodevelopmental disorders that display a gender bias.

### Graphical Abstract

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#### AUTHOR CONTRIBUTIONS

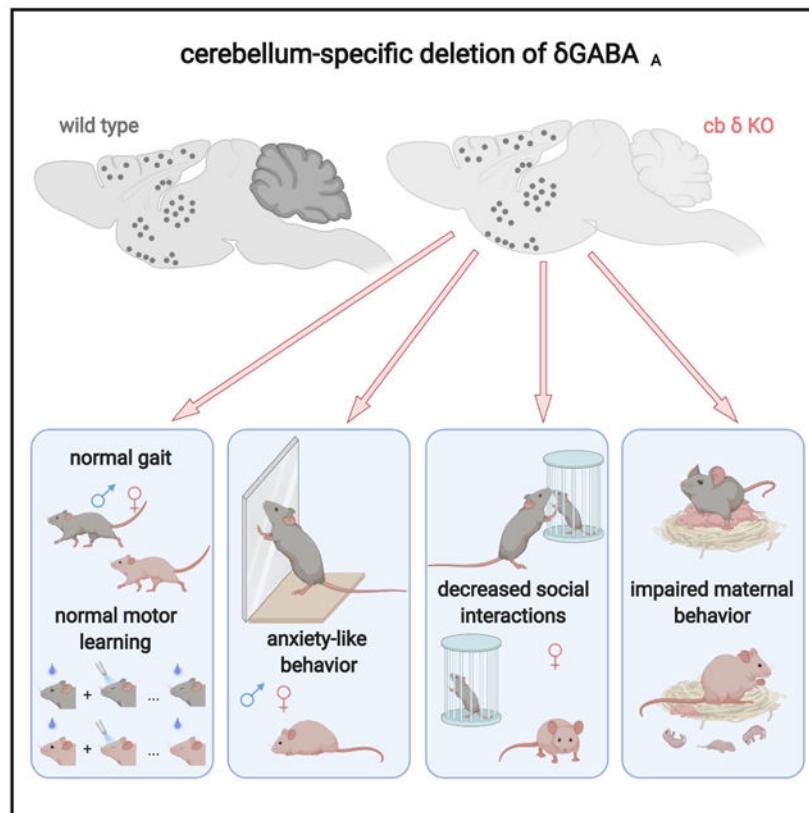
S.R. and W.G.R. conceived the project. S.R. performed electrophysiology recordings, immunohistochemistry (with help from I.F.), *in situ* hybridization (with help from M.E.-R.), imaging and associated analyses, behavioral testing (with help from J.M.K.), and stress experiments. C.G. performed gait analysis and eyeblink conditioning and associated analyses and prepared Figure 3. S.L.P. performed MoSeq analysis with help from W.F.G. and S.R.D. H.N. performed western blots with help from S.R. T.O. helped with data analysis. T.O. and S.L.P. prepared Figures S5 and S6. S.R. prepared all figures other than Figures 3 and S5. S.R. and W.G.R. wrote the manuscript with comments from all authors.

#### SUPPLEMENTAL INFORMATION

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#### DECLARATION OF INTERESTS

S.R.D. is on the scientific advisory board of RBNC Inc., which has licensed the MoSeq technology, and holds patents related to the use of MoSeq to characterize behavior for drug development.



## In Brief

Rudolph et al. show that deletion of the neuromodulator and hormone-sensitive  $\delta$ GABA<sub>A</sub> receptor subunit from cerebellar granule cells results in anxiety-like behaviors and female-specific deficits in social behavior and maternal care.  $\delta$ GABA<sub>A</sub> deletion is associated with hyperexcitability of the cerebellar input layer and altered activation of many stress-related brain regions.

## INTRODUCTION

Normal brain function requires GABAergic inhibition (Ferguson and Gao, 2018; Wood et al., 2017). GABA<sub>A</sub> receptor-mediated inhibition occurs on different timescales and can consist of rapid (phasic) and persistent (tonic) components (Brickley et al., 1996; Farrant and Nusser, 2005; Rossi et al., 2003). Tonic inhibition influences neuronal membrane properties, such as membrane potential and resistance, and, consequently, controls excitability and synaptic integration (Cavelier et al., 2005; Duguid et al., 2012; Mitchell and Silver, 2003; Semyanov et al., 2004). Typically, tonic inhibition is mediated by GABA<sub>A</sub> receptors containing  $\delta$ GABA<sub>A</sub> subunits ( $\delta$ GABA<sub>A</sub>Rs), which are high affinity, slowly desensitizing, extrasynaptically located, and sensitive to fluctuations in ambient GABA levels and neuromodulators (Farrant and Nusser, 2005; Glykys et al., 2008; Martenson et al., 2017; Mihalek et al., 1999; Spigelman et al., 2003; Stell and Mody, 2002; Stell et al., 2003; Vicini et al., 2002; Wei et al., 2003).  $\delta$ GABA<sub>A</sub>Rs are thus poised to respond to context-dependent signals that are essential for controlling behavior.  $\delta$ GABA<sub>A</sub>Rs are involved in

diverse physiological and pathophysiological processes, such as learning and memory; anxiety; stress; sleep; pain; seizures; psychiatric and neurodevelopmental disorders (Hines et al., 2012; Whissell et al., 2015), including attention deficit hyperactivity disorder (ADHD) and autism spectrum disorder (ASD) (Bridi et al., 2017; Martin et al., 2014; Zhang et al., 2017a); pregnancy and maternal behaviors (Maguire and Mody, 2008; Maguire et al., 2009); and estrous cycle-dependent fluctuations in mood and seizure susceptibility (Carver et al., 2014; Maguire et al., 2005; Smith et al., 2006).  $\delta$ GABA<sub>A</sub>Rs have therefore garnered significant interest as potential pharmacological targets for numerous disorders, including postpartum depression (Melón et al., 2018; Meltzer-Brody et al., 2018), epilepsy (Petersen et al., 1983), trauma, panic and anxiety disorders (Rasmusson et al., 2017), insomnia (Orser, 2006; Thakkar et al., 2008), and some syndromic forms of ASD, such as fragile X syndrome (FXS) (Braat and Kooy, 2015; Cogram et al., 2019; Modgil et al., 2019; Olmos-Serrano et al., 2011).

Global  $\delta$ GABA<sub>A</sub> knockout (global  $\delta$  KO) mice show diverse behavioral deficits (Maguire and Mody, 2008; Mihalek et al., 2001; Spigelman et al., 2002; Wiltgen et al., 2005), but the brain regions involved in these deficits are unknown.  $\delta$ GABA<sub>A</sub>Rs are present in many brain regions, however, most attention has focused on  $\delta$ GABA<sub>A</sub>s in the hippocampus, hypothalamus, amygdala, and cortex, regions that have classically been associated with fear and anxiety-related behaviors, stress, and cognition. In contrast, even though the cerebellum shows the highest expression levels of  $\delta$ GABA<sub>A</sub> (Jones et al., 1997; Pirker et al., 2000; Poulter et al., 1992; Wisden et al., 1992), it has been speculated that cerebellar  $\delta$ GABA<sub>A</sub>Rs contribute to motor function rather than non-motor behaviors (Whissell et al., 2015). However, many studies have demonstrated that the cerebellum controls cognitive, social, and emotional processes (Buckner, 2013; Schmammann et al., 2019; Stoodley and Schmammann, 2018; Strick et al., 2009). In addition, many studies have found that cerebellar disruption is associated with psychiatric and neurodevelopmental disorders (Becker and Stoodley, 2013; Fatemi et al., 2012; Kloth et al., 2015; Sathyanesan et al., 2019; Stoodley et al., 2017; Tsai et al., 2012, 2018). Thus, there is considerable overlap in behavioral deficits and disorders that involve the cerebellum and  $\delta$ GABA<sub>A</sub>Rs, suggesting that  $\delta$ GABA<sub>A</sub>Rs in cerebellar granule cells (GCs) could contribute to these behaviors.

In the cerebellar input layer, mossy fibers carry multimodal sensory information from diverse cortical, subcortical, and peripheral regions and excite GCs (Chabrol et al., 2015; Huang et al., 2013; Ishikawa et al., 2015; Witter and De Zeeuw, 2015). Golgi cells provide ambient GABA that tonically inhibits GCs by activating extrasynaptic GABA<sub>A</sub>Rs containing the  $\delta$  and  $\alpha 6$  subunits (Jechlinger et al., 1998).  $\delta$ GABA<sub>A</sub>Rs are therefore well situated to regulate the excitability of the input layer. However, in previous studies, removing the  $\alpha 6$ GABA<sub>A</sub> subunit eliminated tonic inhibition but left excitability unaltered because of compensatory potassium channel expression (Brickley et al., 2001). The behavioral effects of GC hyperexcitability thus remained elusive. How input layer hyperexcitability controls behavior is of particular interest, given the recent observations of dense GC activation during certain behaviors (Badura and De Zeeuw, 2017; Cayco-Gajic and Silver, 2019; Giovannucci et al., 2017; Wagner et al., 2017) that contradict classic models of the cerebellar computation that relied on sparse coding (Marr, 1969; Albus, 1971). Thus, it remains to be determined

whether elimination of  $\delta$ GABA<sub>A</sub> alters GC excitability and whether excitability changes affect behavior.

We examined the effects of GC-specific deletion of  $\delta$ GABA<sub>A</sub> on physiology and behavior. We found that tonic inhibition in GCs lacking  $\delta$ GABA<sub>A</sub> is reduced, leading to hyperexcitability. Despite profound excitability changes in the cerebellar input layer, motor function was unaffected in cerebellar (cb)  $\delta$  KO mice. Remarkably, cb  $\delta$  KO mice displayed deficits in diverse behaviors not previously associated with the cerebellum, and many were selectively altered in cb  $\delta$  KO females, including reduced sociability and abnormal maternal behavior. We also found that GC-selective deletion of  $\delta$ GABA<sub>A</sub> resulted in differential activation of many cortical and subcortical brain areas involved in cognition, anxiety-like behaviors, and the stress response. These findings establish that the molecular make-up of the cerebellum, sensitive to sex and stress-related signals, enables modulation of behavior in a sex-specific manner. These insights may contribute to a better understanding of the mechanisms that underlie behavioral abnormalities in psychiatric and neurodevelopmental disorders that display a sex bias.

## RESULTS

We first determined the overlap of Cre expression in the *Gabra6*-Cre mouse line and  $\delta$ GABA<sub>A</sub> using fluorescence *in situ* hybridization (FISH) in whole-brain sagittal sections of *Gabra6*-Cre  $\times$  flox *tdTomato* mice (Ai14 reporter line; Figure 1A, left panels). FISH-labeled *tdTomato* RNA indicates Cre expression in the GC layer and in the pons (Figure 1A, top), consistent with earlier descriptions (Fünfschilling and Reichardt, 2002). In addition, *TdTomato* transcripts were absent in the thalamus, dentate gyrus, and hypothalamus (Figure 1A, top). *Gabrd* was expressed in the neocortex, striatum, dentate gyrus, thalamus, hypothalamus, and cerebellum but not in the pons (Figure 1A, center; Figure S1). *TdTomato* and *Gabrd* FISH signals did not overlap in other regions of the hippocampus, colliculus, and brain stem nuclei (Figure S1A). Previous reports show that GCs of cochlear nuclei lack *Gabrd* (Campos et al., 2001). Thus, Cre and  $\delta$ GABA<sub>A</sub> expression overlap selectively in cb GCs (Figure 1A, merged image, bottom). To selectively eliminate  $\delta$ GABA<sub>A</sub>s from cb GCs (cb  $\delta$  KO mice), we bred *Gabra6*-Cre to floxed *Gabrd* mice (Lee and Maguire, 2013) (*Gabra6*-Cre+, *Gabrd* flox/flox).

Immunohistochemistry of parasagittal whole-brain sections of control mice (*Gabrd* flox/flox) revealed that GCs strongly express  $\delta$ GABA<sub>A</sub> subunits (Figure 1B, top left). Confocal images of the cerebellar GC layer are consistent with GC membranes (Figure 1B, top right). However, in cb  $\delta$  KO mice, the  $\delta$ GABA<sub>A</sub> signal was abolished almost entirely in the cerebellum (Figure 1B, center). An image of the background fluorescence in global  $\delta$  KO mice is shown for comparison (Figure 1B, bottom). A comparison of the images in Figure 1B suggests that, in cb  $\delta$  KO mice, the  $\delta$ GABA<sub>A</sub> subunit is eliminated from GCs but remains intact in other brain regions. Immunohistochemistry also indicates that  $\delta$ GABA<sub>A</sub> expression does not overlap with *tdTomato* in a sparsely labeled population of hippocampal interneurons and in brain stem nuclei (Figures S1B and S1C). Fluorescent western blot analysis revealed that  $\delta$ GABA<sub>A</sub> is expressed at high levels in the cerebellum of control mice but eliminated from cb  $\delta$  KO and global  $\delta$  KO animals (Figures 1C, top panel, and 1E).

$\delta$ GABA<sub>A</sub> is expressed at relatively low levels in the forebrain of control mice, which were reduced to even lower levels in  $\delta$  KO animals, whereas levels were slightly elevated in cb  $\delta$  KO mice (Figures 1D, top panel, and 1E). We conclude that, in *Gabra6-Cre 3 Gabrd* flox mice,  $\delta$ GABA<sub>A</sub> subunits are eliminated selectively from cerebellar GCs.

Deletion of a GABA<sub>A</sub>R subunit can affect expression of other subunits and, therefore, the relative contributions of phasic and tonic inhibition (Korpi et al., 2002; Nusser et al., 1999; Ogris et al., 2006; Peng et al., 2014). In cerebellar GCs, tonic currents are mediated by extrasynaptic  $\alpha 6/\delta$ -containing receptors, and phasic currents are mediated by  $\alpha 1/\beta 3/\gamma 2$ -containing subunits (Nusser et al., 1998). Besides a small reduction in  $\alpha 6$  levels that is only significant in global  $\delta$  KO mice, there were no significant differences in cb  $\alpha 1$ ,  $\beta 3$ , and  $\gamma 2$  subunit levels in control, cb  $\delta$  KO, and global  $\delta$  KO mice (Figures 1C and 1F), suggesting that phasic currents are likely not upregulated. We also found that spontaneous phasic inhibitory postsynaptic currents (spIPSCs) in GCs were similar in amplitude (control:  $32 \pm 5$  pA,  $n = 7$ ; cb  $\delta$  KO:  $26 \pm 2$  pA,  $n = 9$ ;  $p > 0.2$ ), rise time (control,  $0.8 \pm 0.1$ ; cb  $\delta$  KO,  $1.1 \pm 0.1$  ms;  $p > 0.1$ ), and decay time (control,  $4.5 \pm 0.8$  ms; cb  $\delta$  KO,  $4.8 \pm 0.4$  ms;  $p > 0.4$ ; Mann-Whitney test). This suggests that elimination of  $\delta$ GABA<sub>A</sub> does not have a major influence on other GABA<sub>A</sub>R subunits in the cerebellum.

We assessed tonic currents and the passive properties in control and cb  $\delta$  KO GCs using patch-clamp recordings in cerebellar brain slices (Figures 2A–2E). Initial tonic currents were larger in control compared with cb  $\delta$  KO GCs (Figures 2A and 2B; control:  $74 \pm 10$  pA,  $n = 23$  cells; cb  $\delta$  KO:  $21 \pm 4$  pA,  $n = 14$  cells;  $p < 0.0005$ , Kolmogorov-Smirnov [KS] test). Tonic currents were accompanied by a characteristic increase in GABA<sub>A</sub>R channel noise (Figure 2A, insets; Figure S2F). 4,5,6,7-Tetrahydroisoxazolo (5,4-*c*)pyridin-3(-ol) (THIP), a  $\delta$ GABA<sub>A</sub>-preferring GABA<sub>A</sub>R agonist (Chandra et al., 2006; Meera et al., 2011; Stórustovu and Ebert, 2006), increased current amplitude to a larger extent in control than in cb  $\delta$  KO GCs (Figure 2C; control:  $116 \pm 8$  pA,  $n = 23$  cells; cb  $\delta$  KO:  $38 \pm 8$  pA,  $n = 14$  cells;  $p < 0.0001$ , KS test). The remaining effect of THIP in cb  $\delta$  KOs is consistent with activation of non- $\delta$ GABA<sub>A</sub>-containing receptors (Meera et al., 2011). Blocking GABA<sub>A</sub>Rs eliminated tonic current and greatly reduced noise (Figures 2A–2C, black). A small but significant depolarization was evident in cb  $\delta$  KO GCs (Figure 2D; control:  $-61 \pm 1$  mV,  $n = 28$ ; cb  $\delta$  KO:  $-57 \pm 1$  mV,  $n = 31$ ;  $p < 0.05$ , KS test; Figures S2A–S2C). The input resistance ( $R_i$ ) was higher in cb  $\delta$  KO GCs compared with the control (Figure 2E; control:  $1.0 \pm 0.1$  G $\Omega$ ,  $n = 18$  cells; cb  $\delta$  KO:  $1.5 \pm 0.2$  G $\Omega$ ,  $n = 14$  cells;  $p < 0.02$ , KS test), consistent with reduced tonic current. Additionally, in control animals, THIP decreased  $R_i$ , whereas blocking GABA<sub>A</sub>Rs increased  $R_i$  (Figure S2D), but THIP and SR95531 did not alter  $R_i$  in cb  $\delta$  KO GCs (Figure S2E). Thus, decreased tonic inhibition in cb  $\delta$  KO GCs results in increased  $R_i$ . Tonic current amplitudes were larger in GCs of control females than of males (Figure S2G), whereas THIP evoked currents were of similar amplitude (Figure S2H), suggesting that, although  $\delta$ GABA<sub>A</sub>R density is similar in males and females, baseline activation differs.

Tonic currents regulate GC excitability, but it is unclear whether deletion of  $\delta$ GABA<sub>A</sub> leads to hyperexcitability or whether compensatory mechanisms counteract the loss of tonic inhibition (Brickley et al., 2001). We found that depolarizing current steps evoked more

action potentials in cb  $\delta$  KO GCs than in control GCs (Figure 2F, top). This was true for a range of amplitudes (Figure 2F, bottom; control,  $n = 14$ ; cb  $\delta$  KO,  $n = 14$ ;  $p < 0.0001$ , 2-way ANOVA), indicating that cb  $\delta$  KO GCs are more excitable than control GCs. The observation that THIP decreased excitability (Figure 2G, top left black trace), whereas inhibiting GABA<sub>A</sub>Rs increased excitability of control GCs (Figure 2G, top right black trace; bottom, summary input-output curve; control THIP,  $n = 19$ ; control SR95531,  $n = 17$ ;  $p < 0.0004$ , 2-way ANOVA; Figure 2G) confirmed the importance of tonic GABA<sub>A</sub> currents in regulating GC excitability. Conversely, neither activating nor inhibiting GABA<sub>A</sub>Rs had a significant effect on excitability in cb  $\delta$  KO GCs (Figure 2H; THIP,  $n = 17$ ; SR95531,  $n = 15$ ;  $p > 0.9$ , 2-way ANOVA). Tonic current amplitudes were similar across lobules (Figure S2I); thus, hyperexcitability of cb  $\delta$  KO GCs affects all areas of the cb cortex. These experiments indicate that GCs of cb  $\delta$  KO mice are hyperexcitable because of reduced tonic inhibition.

GC hyperexcitability in cb  $\delta$  KO mice was surprising because eliminating  $\alpha 6$ GABA<sub>A</sub> did not result in hyperexcitability (Brickley et al., 2001). This might be related to the fact that  $\alpha 6$ GABA<sub>A</sub> receptors are still present in GCs after GC-specific  $\delta$ GABA<sub>A</sub> deletion (Martenson et al., 2017), whereas global  $\alpha 6$ GABA<sub>A</sub> resulted in a strong reduction of  $\delta$ GABA<sub>A</sub>. Thus, cb  $\delta$  KO mice provide a unique opportunity to assess the behavioral consequences of input layer hyperexcitability in the cb cortex.

Cerebellar dysfunction is associated with ataxia and abnormal fine motor coordination (Manto et al., 2012). We therefore hypothesized that a considerable increase in excitability of the cerebellar input layer in cb  $\delta$  KO mice could lead to motor dysfunction, including abnormal gait and deficits in motor learning. However, similar to observations with global  $\delta$  KO mice (Wiltgen et al., 2005), we found that motor function was normal in cb  $\delta$  KO mice. To examine whether  $\delta$ GABA<sub>A</sub> deletion results in subtle effects on motor function, we analyzed the gait of cb  $\delta$  KO mice. We tracked walking mice and extracted the positions of the four paws, the base and tip of the tail, and the nose over time (see representative frames of the recorded video in Figure 3A). A velocity plot of different mouse body parts illustrates that control and cb  $\delta$  KO mice move similarly. Nose and tail advance at near-constant velocity, and the hind and front paws move together and touch the floor or are moving forward rapidly. (Figure 3B). There was no difference in any parameter describing gait (Figures 3C–3F, top). As expected, these parameters depend on velocity (Figures 3C–3F, bottom), and the velocity dependence of any of these parameters was the same in control and cb  $\delta$  KO animals. We also found that tail oscillations that accompany walking and tail elevation were unaltered in cb  $\delta$  KO mice (Figures 3G and 3H). Finally, we analyzed the fraction of the cycle in which there are a given number of paws on the floor. At low velocities, it is common that four paws are in contact with the floor. As velocity increases, progressively fewer paws contact the floor, to the point where no paws are in contact with the floor (Figures 3I and 3J). Unlike mice with gait abnormalities that tend to have more feet in contact with the floor at a given velocity (Machado et al., 2015), there was no difference between control and cb  $\delta$  KO mice.

The cerebellum has a well-established role in motor learning, and eyeblink conditioning is a learning paradigm that requires the cerebellum (Thompson, 1986). Pairing a conditioning



stimulus (CS, a weak illumination) that does not initially cause the eye to close with an unconditioned stimulus (US, air puff to the eye) that causes the eye to close by the fifth day of pairing resulted in the CS alone causing the eyelid to partially close in a fraction of trials and by the fifteenth day in eye closure in most trials. Delay conditioning experiments were performed in mice running on a rotating platform as the CS and US were presented (Albergaria et al., 2018). Cb  $\delta$  KO mice learn similarly as control animals (control, Figures 3K and 3L; cb  $\delta$  KO, Figures 3M and 3N). All control mice and cb  $\delta$  KO mice learned to respond to the CS, and control and cb  $\delta$  KO mice learned at approximately the same rate (Figure 3O; control,  $n = 8$ ; cb  $\delta$  KO,  $n = 8$ ). Average responses to the CS alone and to paired CS + US stimulation were very similar for control and cb  $\delta$  KO mice on days 5 and 15 (Figures 3L and 3N). Because cb deficits can affect motor learning (Kloth et al., 2015; Tsai, 2016), we also performed a 3-day accelerating rotarod paradigm. We found that there was no difference between control mice and cb  $\delta$  KO mice in initial performance (control:  $90 \pm 5$  s on rotarod,  $n = 36$ ; cb  $\delta$  KO:  $89 \pm 8$  s on rotarod,  $n = 20$ ;  $p > 0.9$ , Mann-Whitney test; Figure 3P) in females and males (Figure S3C), and motor learning was similar in control and cb  $\delta$  KO animals by the end of training (Figure 3P; control:  $136 \pm 6$  s on rotarod,  $n = 36$ ; cb  $\delta$  KO:  $133 \pm 7$  s on rotarod,  $n = 20$ ;  $p > 0.9$ , Mann-Whitney test) in males and females (Figures S3D and S3E). Thus, remarkably, despite hyperexcitability of the cb input layer in cb  $\delta$  KO mice, we did not detect any abnormalities in motor function.

### Anxiety-like Behaviors in cb $\delta$ KO Mice

$\delta$ GABA<sub>A</sub>Rs are involved in anxiety-related behaviors and stress (Gunn et al., 2011). The behavioral responses to stress are diverse and can include hypervigilance and changes in locomotion (Cabib et al., 1988; Sequeira-Cordero et al., 2019). Open field testing revealed hyperlocomotion in cb  $\delta$  KO mice (Figures S3A and S3B), which could indicate increased stress levels. To examine the effect of  $\delta$ GABA<sub>A</sub> deletion in the cerebellum on anxiety-related behaviors, we first conducted a light/dark test that has been used previously to test the effect of anxiolytic drugs in mice (Crawley, 1985; Lezak et al., 2017). During the test, the mouse was allowed to navigate a dark and a brightly lit compartment (Figure 4A). A mouse displaying increased anxiety-like behavior is thought to elevate avoidance of the bright compartment. Cb  $\delta$  KO mice spent less time in the bright compartment than control mice, as shown in median heatmaps (Figure 4B) and in the summaries of individual animals tested (Figure 4C; control:  $34\% \pm 3\%$ ,  $n = 32$ ; cb  $\delta$  KO mice:  $18\% \pm 3\%$ ,  $n = 27$ ;  $p < 0.0008$ , Mann-Whitney test). Both males and females displayed increased anxiety-like behavior (Figure S4). In addition, a higher fraction of cb  $\delta$  KO mice exclusively remained in the dark compartment during the observation period compared with control animals (6 of 27 and 1 of 32, respectively).

We also observed males and females in the open field. Decreased time spent in the center region of the enclosure is thought to indicate anxiety-like behavior (Figure 4D). Median occupation heatmaps (Figure 4E) and summaries of individual animals (Figure 4F) show that cb  $\delta$  KO mice spent less time than cb  $\delta$  KO in the center of the arena (cb  $\delta$  KO:  $7.3\% \pm 1\%$ ,  $n = 30$ ; control:  $10.3\% \pm 1\%$ ,  $n = 41$ ;  $p < 0.036$ , Mann-Whitney test). Interestingly, only cb  $\delta$  KO females spent less time in the center (control:  $10.7\% \pm 0.9\%$ ,  $n = 32$ ; cb  $\delta$  KO:  $7.7\% \pm 1.1\%$ ,  $n = 21$ ;  $p < 0.032$ ; Figure S4E) but not males (control:  $8.0\% \pm 1.1\%$ ,  $n = 9$ ; cb

$\delta$  KO:  $7.6\% \pm 1.6\%$ ,  $n = 9$ ;  $p > 0.9$ ; Figure S4F). The light/dark and open field tests suggest that especially female cb  $\delta$  KO mice display increased anxiety-like behavior compared with control mice.

### Sex-Specific Differences in Spontaneous Behavior in cb $\delta$ KO Mice Identified with MoSeq

We further used Motion Sequencing (MoSeq) (Wiltschko et al., 2015) to test whether cb  $\delta$  KO mice show abnormalities in spontaneous behavior that are difficult to detect with alternative approaches. MoSeq posits that spontaneous behavior consists of discrete and stereotyped three-dimensional motifs of behavior called “syllables” and allows identification and quantification of these syllables using unsupervised machine learning. MoSeq can detect subtle behavioral differences caused by genetic mutations in mouse models of disease by presentation of sensory cues and in response to optogenetic manipulation (Markowitz et al., 2018; Pisanello et al., 2017; Wiltschko et al., 2015).

Single mice of either sex were observed in a circular behavioral arena with a depth camera (Figure 4G, left). MoSeq identified over 70 distinct syllables (Figures S5 and S6). Three examples are shown in Figure 4G, right. A comparison of the most common syllables ( $>1\%$  usage) identified a subset of syllables whose usage differed between control and cb  $\delta$  KO females but not males (Figure 4H). Although many syllables were used equally in control and cb  $\delta$  KO females (Figures 4H and 4J; control,  $n = 8$ ; cb  $\delta$  KO,  $n = 8$  females), pausing syllables were used less in cb  $\delta$  KO females (Figures 4H and 4I; control,  $n = 8$ ; cb  $\delta$  KO,  $n = 8$  males). Conversely, usage of syllables corresponding to rapid movements, such as darting or turning, were elevated in cb  $\delta$  KO females (Figures 4H and 4K). MoSeq also identified less common behaviors that were more frequent in cb  $\delta$  KO females, including long bouts of grooming (Figure 4L) and high wall jumps, presumably to attempt escape (Figure 4M). These syllables were not differentially expressed in control and cb  $\delta$  KO males. Thus, although the syllabic repertoire of control and cb  $\delta$  KO animals is similar, syllable usage patterns differ between the genotypes, especially in females (Figure 4H, right). The behaviors observed in cb  $\delta$  KO females are consistent with maladaptation to stress, such as removal from the home cage and placement in a behavioral arena, and are reminiscent of complex behaviors displayed after chronic stress (Cabib et al., 1988; Füzesi et al., 2016). This observation is particularly interesting because the  $\delta$ GABA<sub>A</sub> subunit has been implicated in these behaviors (Gunn et al., 2011; Liu et al., 2017; Maguire et al., 2005; Shen et al., 2007; Zhang et al., 2017b), but the possibility that  $\delta$ GABA<sub>A</sub> in the cerebellum could contribute to anxiety-like and stress-related behaviors has not been demonstrated. The apparent milder phenotype observed in cb  $\delta$  KO males could be related to well-established sex differences in the stress response and circulating sex steroids, both of which modulate  $\delta$ GABA<sub>A</sub>Rs (Bangasser and Valentino, 2012; Goldstein et al., 2010; MacKenzie and Maguire, 2013; Tolin and Foa, 2006).

### Decreased Sociability and Abnormal Maternal Behavior in cb $\delta$ KO Females

Our findings suggest that loss of  $\delta$ GABA<sub>A</sub> in cerebellar GCs contributes to stress-related behaviors. Global loss of  $\delta$ GABA<sub>A</sub> function is associated with increased fear-related behavior and stress sensitivity (Maguire and Mody, 2007; Mody and Maguire, 2012; Wiltgen et al., 2005). Stress has been linked to diminished interest in social interactions (Beery and



Kaufer, 2015; File and Hyde, 1978), and anxiolytic agents acting on  $\delta$ GABA<sub>A</sub> have prosocial effects (File, 1980). In addition, cerebellar dysfunction can lead to social deficits in humans and in animal models (Badura et al., 2018; Carta et al., 2019; Schmahmann and Sherman, 1998; Tsai, 2016), and decreased tonic inhibition with the cb input layer is evident in several mouse models of psychiatric disorders (Bruinsma et al., 2015; Kim et al., 2017). We therefore examined whether loss of  $\delta$ GABA<sub>A</sub> in the input layer affects social behavior. We used a 3-chamber paradigm where mice were allowed to explore a chamber containing a sex-matched juvenile conspecific (social stimulus [S]) and a chamber containing a novel object (O). Median occupation heatmaps indicate that control mice preferred investigating the S to the O, whereas cb  $\delta$  KO mice show no preference for either stimulus (Figure 5A; control, left; cb  $\delta$  KO, right). Summary data of individual animals are shown in Figures 5B and 5C (control: 32%  $\pm$  1% in O, 50%  $\pm$  1% in S, n = 54, p < 0.0001; cb  $\delta$  KO: 37%  $\pm$  2% in O, 44%  $\pm$  2% in S, n = 29, p > 0.1; Mann-Whitney test). Control and cb  $\delta$  KO animals did not prefer either chamber in the absence of stimuli (Figures S7A–S7C). Control animals also investigated the conspecific for a longer duration than cb  $\delta$  KO mice (Figure S7D; control: 133  $\pm$  9 s, n = 54; cb  $\delta$  KO: 108  $\pm$  9 s, n = 29; p < 0.04, Mann-Whitney test). The number of entries to the O or social chambers were similar (Figure S7E), suggesting that control and cb  $\delta$  KO mice did not avoid either chamber and showed normal exploratory behavior. However, social deficits were pronounced in females and absent in males. Although control females strongly preferred the social over the O chamber, cb  $\delta$  KO mice had no preference (Figure 5D; summary in Figures 5E and 5F; control: 32%  $\pm$  2% in O, 50%  $\pm$  2% in S, n = 29, p < 0.0001; cb  $\delta$  KO: 39%  $\pm$  4% in O, 41%  $\pm$  3% in S, n = 15, p > 0.8). In contrast, there was no difference in social interest between control and cb  $\delta$  KO (Figures 5G–5I; control: 33%  $\pm$  2% in O, 48%  $\pm$  2% in S, n = 25, p < 0.002; cb  $\delta$  KO: 34%  $\pm$  3% in O, 48%  $\pm$  3% in S, n = 14, p < 0.04; Mann-Whitney test). Thus, deletion of cb  $\delta$ GABA<sub>A</sub> subunits in GCs leads to social deficits in females but not in males.

We tested whether the lack of social interest in cb  $\delta$  KO mice was due to an olfactory deficit by performing an odor discrimination task. We used non-social (water, coconut, raspberry, and banana) and social (male and female urine) odors. Control and cb  $\delta$  KO mice explored most odors for similar durations (Figure S7F). No major differences were observed between the sexes (Figure S7G). These experiments indicate that social deficits in cb  $\delta$  KO mice are not a consequence of an inability to detect social odors.

The neurosteroid sensitivity of  $\delta$  subunit-containing GABA<sub>A</sub>Rs could contribute to the sex dependence of social deficits in cb  $\delta$  KO mice. This raises the possibility that  $\delta$ GABA<sub>A</sub>-containing receptors in the cerebellum might also contribute to other steroid-sensitive behaviors in females, such as postpartum-related changes in maternal care (Maguire and Mody, 2008). Given the high  $\delta$ GABA<sub>A</sub> expression levels in the cerebellum, social deficits specifically in cb  $\delta$  KO females, and the established relationship between stress and poor maternal care (Hillner et al., 2012), we next assayed parental behavior in cb  $\delta$  KO females. We included heterozygous cb  $\delta$  females (cb  $\delta$  HET) because parental care is compromised in HET global  $\delta$  KO mice (Maguire and Mody, 2008). Newborn pups emit olfactory signals and ultrasonic vocalizations that can stimulate spontaneous parental behavior in virgin females, including licking and crouching over pups (Calamandrei and Keverne, 1994; Gandelman, 1973; Noirot, 1969; Thomas and Palmiter, 1997). We thus first tested whether

$\delta$ GABA<sub>A</sub> deletion in the cerebellum affects spontaneous parenting in virgin females. We developed a test where female virgins were allowed to interact with a nest of newborn (postnatal day 1 [P1]–P3) pups and inanimate pup-sized Os to control for novelty (Figure 6A). Median occupation heatmaps (Figure 6B) and summaries of individual experiments (Figures 6C and 6D; Figures S8A and S8B) show that, although control females show a preference for pups, cb  $\delta$  HET and KO females were less interested and also interacted with Os. Upon habituation with pups, female virgins are known to retrieve displaced pups to the nest (Noirot, 1969). We found that control virgins retrieved pups at higher rates than cb  $\delta$  KO and cb  $\delta$  HET virgins (Figure S8C; control, 43%; cb  $\delta$  HET, 29%; cb  $\delta$  KO, 5%;  $p < 0.009$ , chi-square test). These results suggest that virgin cb  $\delta$  KOs and cb  $\delta$  HET females are less likely to express spontaneous parent-like behavior. To determine whether cb  $\delta$ GABA<sub>A</sub>Rs are involved in spontaneous parenting, we took advantage of the behavioral effects of  $\delta$ GABA<sub>A</sub> deletion in cb  $\delta$  HET animals. We administered the  $\delta$  GABA<sub>A</sub>R-preferring agonist THIP via drinking water (STAR Methods) to control, cb  $\delta$  HET, and cb  $\delta$  KO females and assayed parental behavior. We found that THIP rescued behavior in cb  $\delta$  HET females (Figures 6E and 6F; cb  $\delta$  HET, 17%  $\pm$  4% time; cb  $\delta$  HET THIP, 28%  $\pm$  3% time;  $n = 9$ ,  $p < 0.004$ , Wilcoxon matched-pairs signed rank test; for time with Os, see Figures S8D and S8E), whereas it had no effect on controls (control, 30%  $\pm$  3% time; control THIP, 30%  $\pm$  3% time;  $n = 15$ ,  $p > 0.8$ ) and cb  $\delta$  KO females (cb  $\delta$  KO, 16%  $\pm$  4% time; cb  $\delta$  KO THIP, 14%  $\pm$  3% time;  $n = 8$ ,  $p > 0.6$ ). THIP also decreased retrieval time in cb  $\delta$  HETs (Figure S8F) but had no effect on baseline locomotion in cb  $\delta$  HET and cb  $\delta$  KO mice, indicating that THIP was provided at non-sedating concentrations (Figures S8G and S8H). These experiments establish that activation of remaining  $\delta$  GABA<sub>A</sub>Rs in cb  $\delta$  HET females can rapidly rescue parental behavior. The observation that  $\delta$ GABA<sub>A</sub> levels in cb  $\delta$  KO animals are not diminished in brain regions outside of the cerebellum (Figure 1D), together with the lack of effect of THIP on parental behavior in cb  $\delta$  KO females, suggests that the behavioral effect of THIP is due to activation of cerebellar  $\delta$ GABA<sub>A</sub>Rs.

To test whether cerebellar  $\delta$ GABA<sub>A</sub> deletion alters postpartum behavior, we first assayed a robust behavior expressed by postpartum mice: retrieval of pups to the nest (Figure 6H). We found that cb  $\delta$  KO dams delayed initiation of retrieval compared with controls (control: 16  $\pm$  2 s,  $n = 25$ ; cb  $\delta$  HET: 29  $\pm$  5,  $n = 12$ ; cb  $\delta$  KO: 51  $\pm$  9 s,  $n = 10$ ;  $p < 0.008$ , Kruskal-Wallis test; Figures 6H and 6I). Cb  $\delta$  KO dams also took longer to complete retrieval of three displaced pups (Figures S9A and S9B). Reproduction was normal in cb  $\delta$  KOs (Figures S9C and S9D). However, similar to previous reports of global  $\delta$  KO females (Maguire and Mody, 2008), we observed increased cannibalization of newborn pups in cb  $\delta$  KO females (see pups of representative litters in Figures S9E and S9F). Contrary to global  $\delta$  KO dams (Maguire and Mody, 2008), we found no differences in the Porsolt forced swim test (Figure S9G) or nest building (data not shown). Analogous to rescue of virgin behavior, we tested whether THIP could rescue parental behavior in cb  $\delta$  HET dams. We found that THIP administration during the peripartum period (STAR Methods) reduced the retrieval time in cb  $\delta$  HET dams (10  $\pm$  2 s,  $n = 6$ ) to similar durations as for control females (11  $\pm$  2 s,  $n = 6$ ), whereas simultaneously tested cb  $\delta$  HET females that received regular drinking water (87  $\pm$  12 s,  $n = 6$ ; Figures 6J and 6K; Figures S9H and S9I) took longer to retrieve pups. Cannibalization of pups did not occur in THIP-treated cb  $\delta$  HET dams (Figure S9J). These results suggest that

the maternal behaviors disrupted in global  $\delta$  KOs were recapitulated in cb  $\delta$  KO and cb  $\delta$  HET KO mice.

### Differential Activation of Many Brain Areas in cb $\delta$ KO Animals

To examine the consequences of GC hyperexcitability *in vivo*, we assessed neuronal activity at rest and after exposure to stress using whole-brain labeling of c-Fos, an immediate-early gene product that serves as an activity marker (Bullitt, 1990). Control and cb KO animals were left undisturbed or exposed to restraint stress (Figure 7A), a paradigm that has been shown to robustly increase c-Fos expression in many brain regions (de Medeiros et al., 2005; Maras et al., 2014). c-Fos levels in GCs were low in unstressed control and cb  $\delta$  KO animals (Figures 7B, top panels, and 7C; Figure S10). Restraint stress, however, increased c-Fos levels in GCs in control and cb  $\delta$  KO animals, but regional differences were apparent (Figures 7B and 7C; Figure S10). The largest increases were observed in lobules II/II, IV/V/VI, and VIII/IX of the vermis but also in crus 1 and 2, regions implicated in social behavior (Badura et al., 2018; Stoodley et al., 2017). It is unclear whether GC hyperexcitability would increase or diminish net PC output because GCs directly excite and disinaptically inhibit PCs. We did, however, find that restraint stress increased c-Fos levels in the fastigial nucleus (FN) to a larger extent in cb  $\delta$  KO animals than in control animals (Figures 7B and 7C) and that FN neurons project to brain regions involved in motor and non-motor control (Fujita et al., 2020). These observations suggest that cerebellum-specific deletion of  $\delta$ GABA<sub>A</sub>Rs can result in hyperactivation of GCs and cerebellar nuclei.

$\delta$ GABA<sub>A</sub> deletion also caused differential c-Fos activation in other target brain regions of the cerebellum; e.g., the thalamus (Fujita et al., 2020; Strick et al., 2009; Figure 7C; Figure S11; mediodorsal [MD] thalamus, ventral posteromedial [VPM] thalamus, and ventromedial [VM] thalamus); parabrachial nucleus (PBN; Hashimoto et al., 2018; Sadakane et al., 2000; Supple and Kapp, 1994); hippocampal CA1, CA2, and CA3 regions; dorsal caudate; basolateral amygdala (BLA); MD septum; periaqueductal gray (PAG); motor cortex (MoCtx; Figure 7B, third row); retrosplenial cortex (RspCtx; Figures 7B, fourth row, and 7C), limbic cortices (Figure 7C); and bed nucleus of the *stria terminalis* (BST; Figure S11). Conversely, decreased c-Fos levels were evident in the paraventricular nucleus of the hypothalamus (PVN; Figures 7B, bottom row, and 7C) of cb  $\delta$  KO animals. In many regions, stress increased c-Fos expression to a similar extent in control and cb  $\delta$  KO animals, including the lateral septum, lateral amygdala, parietal and auditory cortices, and lateral hypothalamus. Our results suggest that cerebellum-specific manipulation has widespread consequences for the activity of many downstream brain areas, contributing to diverse behavioral changes in cb  $\delta$  KO animals.

## DISCUSSION

Our findings provide numerous insights into the cerebellum and behavior. We show that specific deletion of  $\delta$ GABA<sub>A</sub> from cerebella GCs attenuates tonic inhibition and increases excitability of the input layer, resulting in differential activation of many downstream cortical and subcortical brain regions and behavioral deficits. Most importantly, we find that the cerebellum regulates behaviors that are relevant to psychiatric and neurodevelopmental

disorders in a sex-specific manner. This has important implications for the many disorders that display sex differences with regard to prevalence, severity, range of symptoms, and age of onset (Goldstein et al., 2013; Merikangas and Almasy, 2020; Werling and Geschwind, 2013). Even though risk genes for sex-biased disorders like ASD and schizophrenia are highly expressed in the cerebellum (Menashe et al., 2013; Willsey et al., 2013), the cerebellum generally was not thought to contribute to sex differences in behavior. We also find that the cerebellum unexpectedly regulates maternal behaviors, which is relevant to disorders like postpartum depression. We find that, despite profound increases in input layer excitability, motor function was normal. This points toward a surprising resilience and redundancy of motor circuits.

### **Hyperexcitability and Lack of Dynamic Regulation of Tonic Inhibition in GCs of *cb δ* KO Mice**

$\delta$ GABA<sub>A</sub> deletion (Figure 2) from GCs, unlike ablation of  $\alpha$ 6GABA<sub>A</sub> (Brickley et al., 2001), fails to sufficiently upregulate compensatory 2-pore K<sup>+</sup> channels to prevent hyperexcitability. Our c-Fos experiments further support the notion that GCs in *cb δ* KO animals are more active during stress, suggesting that GCs are hyperexcitable *in vivo* and more responsive to excitatory input. Another important consequence of  $\delta$ GABA<sub>A</sub> deletion is diminished responsiveness to contextual signals. During stress, neurosteroids normally decrease excitability of  $\delta$ GABA<sub>A</sub>R-expressing neurons and attenuate the stress response, but this mechanism is diminished in *cb δ* KO animals. Neuromodulators can also fine-tune ambient GABA levels and tonic inhibition of GCs by regulating the activity of input layer interneurons and PCs (Duguid et al., 2012; Farrant and Nusser, 2005; Guo et al., 2016; Mitchell and Silver, 2003). In addition, neurosteroids modulate  $\delta$ GABA<sub>A</sub>-containing receptors directly (Stell et al., 2003; Vicini et al., 2002) and thus regulate GC excitability during diverse physiological states, such as during the estrous cycle (Maguire et al., 2005; Wu et al., 2013), the postpartum period, and stress (Camille Melón and Maguire, 2016; Maguire and Mody, 2008; Maguire et al., 2009). For these reasons, the ability of GCs to dynamically adjust their excitability in an activity- and context-dependent manner is compromised in *cb δ* KOs.

### **Motor Function Is Normal in *cb δ* KO Mice**

Given the cerebellum's role in locomotion, we were surprised to find normal motor behavior in *cb δ* KO mice (Figure 3). However, there is precedence for the relative insensitivity of motor performance to manipulations of the input layer (Bruinsma et al., 2015; Wiltgen et al., 2005). Our c-Fos experiments suggest that stress increases the excitability of GCs throughout the cerebellar cortex, including motor (e.g., anterior vermal lobules) and non-motor regions (crus 1/2 and posterior vermal lobules). Together, these observations suggest that basic motor function is remarkably insensitive to manipulations of the input layer.

Several explanations could account for the lack of motor deficits in *cb δ* KO mice. Circuit-level compensation possibly overcomes GC hyperactivation in *cb δ* KO mice. This could involve weakening GC-to-PC synapses. It is also possible that molecular layer interneurons (MLIs) can compensate for GC hyperexcitability by increasing inhibition of PCs. Paradoxically, our c-Fos experiments show that stress results in more pronounced c-Fos

expression in the MoCtx and other motor areas in cb  $\delta$  KO mice, suggesting that motor circuits might adapt better to GC hyperexcitability than non-motor systems.

### Sex-Specific Behavioral Differences Revealed in cb $\delta$ KO Mice

Unbiased behavioral testing is well-suited to characterize changes in motor and non-motor behavior arising from cerebellar manipulation. We found that cb  $\delta$  KO females were more hyperactive (Figure 4), reminiscent of stimulant-treated mice (Wiltschko et al., 2020), ASD and ADHD mouse models (Angelakos et al., 2017; Dalla Vecchia et al., 2019; Schmeisser et al., 2012), and chronically stressed animals (Cabib et al., 1988; Füzesi et al., 2016; Strekalova et al., 2005). The observed sex differences in our mouse model could be due to differences in the neuroendocrine response to stress or circulating levels of sex steroids and their synthesis enzymes, all of which are related to sex differences in the stress response across species (Bangasser and Valentino, 2012; Bangasser and Wiersielis, 2018; Beery and Kaufers, 2015; Goldstein et al., 2010). These factors could lead to differential modulation of tonic currents and a greater sensitivity to  $\delta$ GABA<sub>A</sub> deletion in females (Schüle et al., 2014; Ströhle et al., 2002; Yoshizawa et al., 2017). Another possible explanation is the well-established influence of gonadal hormones on social behavior (Bell, 2018). Finally, protein deletion can affect gene expression in a sex-dependent manner, as in Angelman syndrome (Koyavski et al., 2019). Sex differences in functional compensation have been observed previously in the cerebellum (Mercer et al., 2016). Social deficits are consistent with the cerebellum's role in social behavior (e.g., in the context of ASD), especially the hyperactivation of crus 1 and 2 and posterior vermal lobules identified in our c-Fos study (Schmahmann, 2019; Tsai, 2016; Wang et al., 2014). What sets our findings apart from previous cerebellum-specific manipulations is that the social deficits were confined to females. With its dense expression of neurosteroid-sensitive  $\delta$ GABA<sub>A</sub>Rs, the cerebellum is poised to control behavior in a sex-specific manner. The sex specificity is intriguing because many psychiatric disorders show sex bias in humans. We speculate that  $\delta$ GABA<sub>A</sub>R-targeting treatments, such as the recently introduced postpartum depression drug brexanolone (Zulresso, Sage Therapeutics), might also be effective for treating other disorders. Clinical trials of drugs targeting  $\delta$ GABA<sub>A</sub>Rs in Angelman syndrome, a monogenic form of ASD associated with sex-dependent phenotypes (Koyavski et al., 2019), are already underway (L. Bird et al., 2019, AAN, abstract).

It is unlikely that social deficits in cb  $\delta$  KO mice reflect differences in primary sensory processing. We found no difference in odor recognition, and although our mouse line expresses Cre in the cochlear nuclei, there is no  $\delta$ GABA<sub>A</sub> expression in GCs of the ventral or dorsal cochlear nuclei (Campos et al., 2001), and hearing is normal in global  $\delta$  KO mice (Maison et al., 2006). In addition, c-Fos experiments showed that activation of the auditory cortex was similar in control and cb  $\delta$  KO mice (Figure 7).

### Increased Anxiety-like Behavior in cb $\delta$ KO Mice

Many clinical reports implicate the cerebellum in anxiety, phobia, generalized anxiety disorder, stress, and post-traumatic stress disorder (PTSD) (Caulfield et al., 2016; Moreno-Rius, 2018), but animal models linking the cerebellum and anxiety are rare (but see Hilber et al., 2004; Lorivel et al., 2014). Many previous studies investigating  $\delta$ GABA<sub>A</sub>Rs and anxiety

focused on brain regions classically associated with fear, such as the amygdala, hypothalamus, and hippocampus (Lee et al., 2014; Liu et al., 2017; Maguire et al., 2005). Our c-Fos experiments suggest, however, that the many brain regions involved in anxiety and stress-related behaviors, such as the BLA, PAG, PVN, BST, RspCtx, and limbic cortex, are differentially activated in cb  $\delta$  KO animals.

$\delta$ GABA<sub>A</sub> has an important role in anxiety and fear-related behaviors (MacKenzie and Maguire, 2013; Whissell et al., 2015), and endogenous compounds and drugs acting on  $\delta$ GABA<sub>A</sub>Rs affect anxiety (Eser et al., 2006). Similarly, neurosteroids and  $\delta$ GABA<sub>A</sub> expression levels are thought to contribute to anxiety-like behaviors during the ovarian cycle (Maguire et al., 2005; Smith et al., 2006) and puberty (Smith, 2013). Ethanol, targeting extrasynaptic  $\delta$ GABA<sub>A</sub>Rs, is thought to produce its effects by increasing tonic inhibition (Richardson and Rossi, 2017). Our study establishes the importance of  $\delta$ GABA<sub>A</sub>Rs in the cb input layer in regulating anxiety and stress-related behaviors.

### Abnormal Maternal Behavior in cb $\delta$ KO Females

There is considerable evidence that  $\delta$ GABA<sub>A</sub>Rs and neurosteroid signaling are involved in postpartum depression (Maguire and Mody, 2009). Although  $\delta$ GABA<sub>A</sub>Rs in dentate GCs (Maguire and Mody, 2008; Maguire et al., 2009) and in hippocampal parvalbumin-positive (PV+) interneurons (Ferando and Mody, 2013) have been implicated in postpartum depression, the brain regions regulating postpartum depression have not been identified directly in conditional  $\delta$  KO mice. Unexpectedly, we observed abnormal maternal care in cb  $\delta$  KO mice (Figure 6). Given the strong connection between anxiety, stress, and postpartum depression (Camille Melón and Maguire, 2016; Payne and Maguire, 2019), these are likely contributing factors to maternal care deficits in cb  $\delta$  KO females. c-Fos experiments show differential activation in hippocampal and hypothalamic regions, but further studies are needed to determine whether  $\delta$ GABA<sub>A</sub>Rs in other brain regions act synergistically to mediate maternal behaviors.

### Differential Activation of Many Brain Regions in cb $\delta$ KO Animals in Response to Acute Stress

Emerging evidence from animal and human studies suggests that the cerebellum plays a role in the stress response (Moreno-Rius, 2019), enabled by anatomical connectivity to stress-related brain structures and its molecular machinery. Assaying restraint stress-evoked c-Fos expression provides important insights into how elimination of  $\delta$ GABA<sub>A</sub> in GCs leads to differential activation of downstream brain regions that could account for deficits in social, anxiety-like, and parental behaviors. Stress-evoked increases in c-Fos expression in the cerebellum indicate activation of the anterior sensorimotor as well as posterior areas associated with cognition and emotion (Badura et al., 2018; Schmähmann, 2019; Stoodley et al., 2017; Strick et al., 2009). Although changes in PC activity cannot be monitored faithfully with c-Fos because of their spontaneous activity, we found increased c-Fos signal in the deep cerebellar nuclei (DCN), especially the FN, as well as in the PBN, suggesting altered PC output in cb  $\delta$  KO animals. The target regions have been associated previously with cognition, emotion, and autonomic control (Berntson and Torello, 1982; Fujita et al., 2020; Sadakane et al., 2000). Stress results in hyperactivation of sensorimotor cortices and



many stress-related brain structures that are thought to be modulated by the cerebellum; e.g., various regions of the cortex (Badura et al., 2018; Choe et al., 2018; Strick et al., 2009), PAG (Gonzalo-Ruiz et al., 1990; Koutsikou et al., 2014; Vaaga et al., 2020), hypothalamus (Dietrichs, 1984; Zhu et al., 2006), hippocampus (Iglói et al., 2015; Watson et al., 2019), striatum (Fujita et al., 2020), and other regions, like the BST and amygdala. In addition, hypoactivation of the PVN in cb  $\delta$  KO mice can indicate hypothalamic–pituitary–adrenal (HPA) axis dysregulation and chronic stress in cb  $\delta$  KO animals (Cohen et al., 2006; Kinlein et al., 2015; Whitaker and Gilpin, 2015). These observations are in line with recent tracing studies (Fujita et al., 2020; Hashimoto et al., 2018; Pisano et al., 2020) and suggest that altering sensory integration in the input layer can affect numerous circuits throughout the brain.

## Conclusions

Here we show that excitability of the cerebellar input layer can affect many behaviors and that the repertoire of cerebellum-dependent behaviors is far greater than appreciated previously. Our results substantiate the cerebellum's role in stress and anxiety-related and social behaviors and provide insights into how the molecular make-up of the cerebellum can allow sex-specific modulation of behavior. These findings are critical for a better understanding of psychiatric and neurodevelopmental disorders that show a sex bias. Thus, we speculate that manipulating excitability of the cerebellar input layer could relieve some symptoms associated with these disorders.

## STAR★METHODS

### RESOURCE AVAILABILITY

**Lead Contact**—Further information and requests for resources and reagents should be directed to and will be fulfilled by the Lead Contact, Wade Regehr (Wade\_Regehr@hms.harvard.edu).

**Materials Availability**—The study did not generate new unique reagents

**Data and Code Availability**—The datasets and code supporting the current study have not been deposited in a public repository but are available from the corresponding author on request. MoSeq code can be obtained from S.R.D. (srdatta@hms.harvard.edu)

### EXPERIMENTAL MODEL AND SUBJECT DETAILS

**Mice**—Animal procedures have been carried out in accordance with the NIH and Animal Care and Use committee (IACUC) guidelines, and protocols approved by the Harvard Medical Area Standing Committee on Animals. *Floxed* Gabrd mice (*Gabrd<sup>tm1.1Jmag/J</sup>*, Jackson Labs stock# 023836) and Gabrd knock-out mice (*Gabrd<sup>tm1Geh/J</sup>* Jackson labs stock# 003725) were obtained from Dr. Jamie Maguire (Tufts University). Gabra6-Cre (B6.D2-Tg(Gabra6-Cre)B1Lfr/Mmucd) mice were obtained from the MMRC. A reporter line expressing tdTomato in Cre-positive cells (*floxed* tdTomato line A14) was obtained from Jackson labs (stock# 007908). Animals were kept on a mixed background (129Sv/SvJ and B6/C57). Mice were housed under standard conditions in groups of 2–5 animals on a 12

h light-dark cycle with food and water available *ad libitum*. For parental behavior assays dams were singly housed. Adult animals of either sex 2–5 months of age were used for all experiments, including FISH, immunohistochemistry, electrophysiology, and behavioral testing. For behavioral experiments and physiology, littermate controls (Cre-negative, *Gabrd f/f* or Cre-positive  $\times +/+$ ), *cb  $\delta$  KO* (Cre+  $\times$  *Gabrd f/f*) and in select cases *cb  $\delta$  HET* (Cre+  $\times$  *f/f*) animals were used. No difference was observed between cre-negative, *Gabrd f/f* or Cre-positive  $\times +/+$  animals and data were thus pooled).

## METHOD DETAILS

**Fluorescence *In situ* hybridization (FISH)**—Animals were anesthetized with isoflurane and the brain was rapidly removed and frozen on dry ice before embedding in optimal cutting temperature (OCT) compound (Tissue-Tek). Tissue was cut on a cryostat (Microm HM500-CM) at a thickness of 20  $\mu$ m and mounted on glass slides (Superfrost Plus, VWR, 48311–703). Fluorescent *in situ* hybridization was performed according to the ACD-Bio RNAscope Multiplex Assay manual, (document Number 320513) with minor modifications. The samples were then fixed in 4% paraformaldehyde in phosphate-buffered saline (PBS) for 15 min at 4°C, then dehydrated with 50% ( $\times$  1), 70% ( $\times$  1), and 100% ( $\times$  2) ethanol washes for 5 min each. The slides were then air-dried and a barrier was drawn around the tissue section with an Immedge hydrophobic barrier pen (Vector Laboratories). The tissue was incubated in RNAscope protease III reagent (ACD-Bio 322337) at room temperature for 30 min, then rinsed twice in PBS for 5 min. Fluorophore-conjugated probes, *Gabrd-Mm-C3* probe (Cat# 459481-C3), *tdTomato-C2* probe (Cat# 317041) were incubated with the slide-mounted tissue sections at 40°C in a HybEZ II oven (ACD-Bio) for 2 hours and washed twice in RNAscope wash buffer reagent (ACD-Bio 310091). Fluorescence amplification steps were then applied as follows: incubate in AMP 1-FL for 30 minutes at 40°C (HybEZ oven), followed with 2 $\times$  wash with 1x wash buffer for 3 min at room temperature, the tissue was then incubated in AMP 2-FL for 15 minutes at 40°C, followed by 2 $\times$  wash, then incubated in AMP 3-FL for 30 min at 40°C, followed with 2x wash, and lastly was incubated in AMP 4-FL-A for 15 minutes at 40 degrees, then 2 $\times$  wash. Sections were then stained with DAPI and mounted with ProLong antifade reagent (Thermo Fisher Scientific P36930). In addition, control probes were used to ensure the quality of the *in situ* experiment. The positive control is a cocktail of housekeeping genes (*C1-Mm-Polr2a*, *C2-Mm PPIB* and *C3-Mm-UBC*). The negative control probe targets bacterial RNA (*C1*, *C2*, *C3-dapB*). The slides were imaged using whole slide scanning microscope (Olympus VS120) with a 20X air objective.

**Immunohistochemistry**—Mice were anesthetized with isoflurane and perfused with ice cold phosphate buffered saline (PBS, pH = 7.4, Sigma Cat# P-3813), followed by a solution containing 4% paraformaldehyde in PBS. The brain was removed and postfixed in the same solution at 4°C overnight. For slicing, the brain was embedded in 4% agar (Sea Plaque, Lonza, Cat# 50101) and then sliced in PBS using a vibratome (VT1000S, Leica) at a thickness of 50  $\mu$ m. Antigen retrieval was performed prior to immunostaining. Slices were permeabilized in 0.2% TritonX (Sigma Cat# T9284) in PBS for 30 min and then incubated in a solution containing 0.001% trypsin (Sigma Cat# T5266) and 0.001%  $\text{Ca}_2\text{Cl}$  in PBS for 1 minute. Slices were then rinsed 3 times for 5 minutes in PBS and then blocked in a

solution containing 4% normal goat serum (NGS), 0.1% TritonX in PBS for 1 hour. After blocking, slices were incubated in the same solution with the addition of primary antibody and 0.001% trypsin inhibitor (Sigma Cat# T6522) overnight at 4°C. Slices were then washed 3 times for 10 minutes and incubated in 4% NGS, 0.1% TritonX in PBS with addition of secondary antibody for 2 hours at room temperature. Slices were then washed 3 times for 5 minutes in PBS, mounted on glass slides (Superfrost Plus, VWR, Cat# 48311-703) and covered with mounting medium (ProLongDiamond, Thermo Fisher Scientific, Cat# P36961) and a glass coverslip. The mounting medium was allowed to cure for at least 24 hours before imaging.

**Imaging and image analysis**—Whole-brain images were taken on an Olympus VS120 slide scanner, and confocal stacks were acquired on an Olympus FV1000 confocal microscope. Images were processed using standard routines in Fiji (ImageJ). Some figures (Graphical Abstract and Figures 4A, 4D, 6A, 6G, and 7A) were prepared with BioRender ([Biorender.com](https://www.biorender.com)).

**Quantitative western blotting**—Western blotting was performed according to standard protocols. For quantitative assessment of protein levels in mice brain tissues, fluorescently tagged secondary antibodies were used. Prior to harvesting tissue mice were anesthetized with isoflurane and perfused transcardially with ice cold PBS. The brain was then rapidly removed from the skull and tissue was dissected in PBS and frozen at -80°C until further processing. Tissue was lysed in a buffer containing 150 mM NaCl, 25 mM HEPES, 4 mM EGTA and protease inhibitor (Sigma-Aldrich, Cat# P8340). After addition of SDS harvested tissue underwent ten freeze-thaw cycles (-80 to +55°C). After SDS-PAGE, gels were transferred onto nitrocellulose membranes and blocked with filtered 5% nonfat milk/5% goat serum in Tris-buffered saline for one hour at room temperature and incubated with primary antibodies with 5% BSA in Tris-buffered saline containing 0.1% Tween-20 (TBST) overnight at 4°C. Each membrane was incubated with primary antibodies against GABA<sub>A</sub> receptor subunits as follows: rabbit anti- $\alpha$ 1GABA<sub>A</sub>R (1:10 000; Abcam, Cat# ab33299, RRID: AB\_732498), rabbit anti-  $\alpha$ 6GABA<sub>A</sub>R (1:1000; GeneTex, Cat# GTX130947), mouse anti-  $\beta$ 3GABA<sub>A</sub>R (Neuromab, Cat# 75-149, RRID: AB\_2109585), rabbit anti-  $\gamma$ 2 GABA<sub>A</sub>R (1:1000; Synaptic Systems, Cat# 224 003, RRID: AB\_2263066), rabbit anti-  $\delta$ GABA<sub>A</sub>R (1:2000; custom made, gift from Dr. W. Sieghart and Dr. P. Scholze, Medical University of Vienna, Vienna, Austria). The following fluorescent secondary antibodies were used prior to washing with TBST: donkey anti-mouse IRDye 800CW IgG (1:10,000; LI-COR, Cat. No.: 926-32212, RRID: AB\_621847), donkey anti-rabbit IRDye 800CW IgG (1:10,000; LI-COR, Cat. No.:926-32213, RRID: AB\_621848), donkey anti-mouse IRDye 680RD IgG (1:10,000; LI-COR, Cat. No.:926-32222, RRID: AB\_621844), and donkey anti-rabbit IRDye 680RD IgG (1:10,000; LI-COR, Cat. No.:926-32223, RRID: AB\_621845). Visualization was carried out with the LI-COR Odyssey® fluorescent scanner and software (LI-COR Biosciences). Blots were imaged using an Odyssey Infrared Imaging System Scan, at a resolution of 42  $\mu$ m. Images were analyzed in NIH ImageJ Software.

**Slice preparation for electrophysiology**—Mice to be used for electrophysiological recordings were retrieved from the animal facility and allowed to acclimate for at least 8

hours before the experiment to reduce stress. Animals of either sex aged 2–4 months were anesthetized in their home cage by introduction of an isoflurane-soaked cloth and then perfused transcardially under continued isoflurane anesthesia with ice cold cutting solution containing (in mM) 110 CholineCl, 7 MgCl<sub>2</sub>, 2.5 KCl, 1.25 NaH<sub>2</sub>PO<sub>4</sub>, 0.5 CaCl<sub>2</sub>, 25 Glucose, 11.5 Na-ascorbate, 3 Na-pyruvate, 25 NaHCO<sub>3</sub>, 0.003 (R)-CPP, equilibrated with 95% O<sub>2</sub> and 5% CO<sub>2</sub>. The brain was rapidly dissected and the cerebellum was cut into 250–270 μm thick parasagittal slices in the same solution on a vibratome (VT1200S, Leica). Slices were then transferred to 34°C warm artificial cerebrospinal fluid (ACSF) containing (in mM) 125 NaCl, 26 NaHCO<sub>3</sub>, 1.25 NaH<sub>2</sub>PO<sub>4</sub>, 2.5 KCl, 1 MgCl<sub>2</sub>, 1.5 CaCl<sub>2</sub>, and 25 glucose, equilibrated with 95% O<sub>2</sub> and 5% CO<sub>2</sub> and incubated for 30 min. Slices were then stored at room temperature until recording for up to 6 hours.

**Electrophysiology**—Whole-cell recordings were obtained from visually identified granule cells using a 40× water-immersion objective on an upright microscope (Olympus BX51WI). Pipettes were pulled from BF150-86-10 borosilicate glass (Sutter Instrument Co., Novato, CA) at resistances of 4–5 MΩ on a Sutter P-97 horizontal puller. Electrophysiological recordings were performed at ~32°C. For voltage clamp recordings, the internal solution contained (in mM): 30 K-gluconate, 110 KCl, 10 HEPES, 0.5 EGTA, 3 MgATP, 0.5 Na<sub>3</sub>GTP, 5 phosphocreatine-tris<sub>2</sub>, 5 phosphocreatine-Na<sub>2</sub>, 5 QX314 chloride. The chloride reversal potential was ~0 mV. For current clamp recordings, the internal solution contained (in mM) 130 K-gluconate, 10 KCl, 10 HEPES, 0.5 EGTA, 3 MgATP, 0.5 Na<sub>3</sub>GTP, 5 phosphocreatine-tris<sub>2</sub>, 5 phosphocreatine-Na<sub>2</sub>. The chloride reversal potential was ~–65 mV, similar to what was reported for GCs (Brickley et al., 1996). Seal resistance for all GC recordings was > 5 GΩ. Electrophysiology data were acquired using a Multiclamp 700B amplifier (Axon Instruments), digitized at 50 kHz, filtered at 4 kHz, and controlled by software custom written in IGOR Pro (Lake Oswego, OR). The recording ACSF included (in μM): 2.5 (R)-CPP, 5 NBQX hydrochloride, 2 CGP, 1 strychnine to block glutamatergic receptors, GABA<sub>B</sub> receptors and glycine receptors, respectively. SR 95531 (100 μM) was used to block GABA<sub>A</sub> receptor-mediated currents. The δGABA<sub>A</sub>R preferring agonist 4,5,6,7-tetrahydroisoxazolo[5,4-c]pyridin-3-ol (THIP, 1 μM) was used to activate tonic GABAergic currents. All drugs were purchased from Abcam (Cambridge, MA) or Tocris (Bristol, UK). Analysis of electrophysiological data was performed with custom routines written in IgorPro (Wavementrics, Lake Oswego, OR) or in AxoGraphX.

**Behavioral testing**—All behavior testing was performed in adult mice (> 8 weeks old) of either sex with the experimenter blind to the genotype. Before behavioral testing, mice were transferred to the behavior room and allowed to acclimate for at least 30 min. Animals were housed on a 12 h light-dark cycle, and the experiments were carried out at the beginning of the dark cycle. All equipment used for behavioral testing was cleaned with 70% ethanol in between experiments. Most behavioral testing was videotaped at ~30 fps using a 720p USB Camera with IR LEDs (ELP, Ailipu Technology Co.) and the iSpy open source video surveillance software suite (<https://www.ispyconnect.com>). Offline analysis was automated using custom scripts written in MATLAB (Mathworks). Select behaviors were scored live (pup retrieval, olfactory testing), or analyses was carried out manually (grooming, forced swim).

**Gait analysis**—To assess baseline locomotion, gait analysis was run over the course of 8 consecutive days with 5 trials done for each animal per day. A custom video recording setup was used for evaluating gait patterns in mice. The setup consisted of an infrared illuminated transparent linear corridor (64.5(L) × 4 (W) × 6 (H) cm) atop a borosilicate glass floor. The corridor was flanked by two mirrors on the side angled at 48 degrees to project the images of the side views toward a video camera (Bonito CL-400B/C 2320 × 700 pixels at 200 frames per second, Allied Vision, Exton, PA.) situated beneath the glass floor. Three views were captured simultaneously at 200 fps as the animal walk down the linear corridor in a self-initiated manner. To annotate the locations of the body parts (nose, base of the tail, tip of the tail, left/right forepaws, and left/right hind paws), a convolutional neural network based on the stacked hour-glass-network (Newell et. al., 2016) was trained using PyTorch on 500 manually annotated sample frames. We used the neural network to annotate frame-by-frame which results in a time series for the location of all aforementioned body parts. A Hidden Markov Model (HMM) was applied on the time series to segment each video into individual gait cycle. For a single gait cycle, we measured the following parameters: cadence, the number of cycles per second in Hz; stride length, the maximum distance traveled by the forepaw within a single cycle; paw width, the lateral distance between the two diagonal supports at any given time; stance duration, the average amount of time that the paws are on the ground during; velocity, the average speed of the center of mass; and tail elevation, the absolute height of the tip of the tail from the floor. We used a linear mixed effect model of the form  $Y \sim \text{Intercept} + \text{genotype} + \text{sex} + \text{genotype}:\text{sex} + (1|\text{name})$  and ANOVA for statistical test in the non-covariate case. For the covariate analysis involving the velocity we fitted a model of the form  $Y \sim \text{Intercept} + \text{Velocity} + \text{genotype} + \text{sex} + \text{genotype}:\text{sex} + \text{velocity}:\text{genotype} + \text{velocity}:\text{sex} + (1|\text{name})$  and ANOVA for testing statistical significance.

**Rotarod**—To evaluate motor learning, we performed a rotarod assay over the course of three consecutive days, with 5 trials performed each day. Mice were placed on a rotating rod device (Rotamex-5 Rota-Rod, Columbus Instruments, controlled by Rotamex-5 software) running at 4 rpm baseline speed. After brief habituation, acceleration of the rod was initiated. The rod accelerated from an initial speed of 4 rpm to a maximum speed of 40 rpm in 10 s intervals. An animal fall was detected by infrared photo cells crossing the space above the rod and the time-to-fall was recorded. A fall was recorded either if the mouse fell off the rod, or loops around the rod without running. Animals were allowed to rest for 1 min between trials.

**Eyeblink conditioning**—Adult mice of either sex performed motor behavior experiments during the dark cycle when they are most active and alert. The setup was housed within a ventilated, anechoic, and sound-insulated behavior chamber. Prior to the experiment, animals were implanted with a head bracket and allowed to recover for four days of post-surgery. To accustom the animals to head-fixation and the treadmill, five days of habituation were performed prior to the first day of eyeblink training. During habituation, the animal was head-fixed atop of a motorized treadmill six inches in diameter rotating at 20 mm/s. The treadmill was kept at this speed throughout the habituation period. During training, a white LED flash was used as the conditioned stimulus (CS) and a 50 ms, 15 psi air puff directed at the opposite eye was delivered as the unconditioned stimulus (US). The air puff was

delivered via a 21-gauge blunt tip needle mounted on a manipulator to allow for individual adjustments. The pneumatic and electronics necessary for the control of the air puff was based on the design of Openspritzer (Forman et al., 2017). To record eyelid movement, we illuminated the behavior chamber using an IR lamp and recorded the eye with a high-speed camera (Mako U-029B, Allied Vision, Exton, PA.) and a macro lens (1/2" 4–12mm F/1.2, Tamron, Commack, NY.) at 300 fps. The vertical span of the eyelid opening was measured as a function over time using a custom MATLAB script. An inter-stimulus-interval of 500 ms was used. 100 trials of CS-US pairing and 10 trials of only the CS were presented to the animal per day for 15 days. The inter-trial interval was randomized between 4 to 12 s.

**Open field**—Animals were placed in an uncovered rectangular behavior arena (30.3 cm × 45.7 cm, 30.5 cm high) containing fresh bedding and observed for 10 min without intervention. Analysis was performed offline using MATLAB (position, velocity, path traveled).

**3-chamber assay**—Sociability was assayed with the 3-chamber task. The behavioral arena consisted of a clear rectangular Plexiglas box (40.5 cm wide, 60 cm long, 22 cm high) without a top cover and divided into 3 equally sized compartments by clear walls. Each divider contained a 10.2 cm × 5.4 cm rectangular opening to allow navigation between the compartments. The left and the right chamber contained inverted wire cups (10 cm in diameter). Before behavioral testing, mice were allowed to navigate the middle chamber for 5 minutes with openings to the adjacent chambers closed. During a 10-minute habituation session, the doors were then opened and mice were allowed to freely navigate all 3 chambers for 10 minutes in the absence of any stimuli. After habituation, the doors were closed again while the animal remained in the central chamber. A social stimulus (juvenile mouse aged 15–30 days of the same sex and strain) and a novel object (mouse-sized plastic toy, Schleich GmbH, Germany) were placed in opposite wire cups. The sides of social and non-social stimuli were randomly selected to control for preference to either side of the arena. After stimulus placement, the doors were opened and the animal was observed for another 10 minutes. Automated analysis was performed offline using MATLAB.

**Light/dark chamber**—A light-dark chamber assay was used to measure anxiety-like in adult mice. The experiment consisted of a Plexiglas arena with the same outer dimensions as used for the 3-chamber assay. The light-dark chamber arena consisted of two chambers, with one removable divider, including a door which remained open throughout. The light chamber was 40.5 cm × 40 cm chamber and brightly lit (> 600 lux) and uncovered. The dark chamber was 40.5 cm × 20 cm and covered with darkly tinted Plexiglass that allowed videotaping of the mouse with an IR camera. Light intensity inside the dark chamber was < 10 lux. At the beginning of the experiment, the animal was placed in the dark chamber and was allowed to freely navigate both chambers for 10 min.

**Motion Sequencing**—Motion Sequencing (MoSeq)-based behavioral analysis was performed as in Wiltschko et al., 2015 and Markowitz et al., 2018. In brief, MoSeq uses unsupervised machine learning techniques to identify the number and content of behavioral syllables out of which mice compose their behavior; identifying these syllables allows each



video frame of a mouse behavioral experiment to be assigned a label identifying which syllable is being expressed at any moment in time. Behavioral phenotypes that distinguish wild-type and mutant mice can be identified by comparing differences in how often individual syllables are used in a given experiment. Here, individual mice (n = 8 males and 8 females for each genotype) were imaged for 1–3 60-minute-long sessions using a Kinect2 depth sensor while behaving in a circular open field under red light illumination. These 3d imaging data were submitted to the MoSeq pipeline, which includes mouse extraction, denoising, and alignment steps before computational modeling. As has been done previously (Wiltchko et al., 2015), the kappa parameter (which sets the timescale at which syllables are identified) was specified by matching the distribution of syllable durations to a model-free changepoint distribution.

Separate models were trained for males and females. All mice of a given gender were jointly modeled with a shared transition. MoSeq identified 61 syllables in females and 64 syllables in males that make up 90% of frames that comprise each dataset. For convenience, in Figure 5, we depict genotype-driven differences in the top 28 used syllables with each accounting for at least 1% of the data in each genotype and gender. Together, these comprise 47% and 56% of the frames in the female and male datasets respectively. Significant differences in syllable expression between genotypes were determined using the Mann-Whitney U test, with the Benjamini and Hochberg correction for multiple comparisons at an alpha level of 0.05.

Because males and females were modeled independently, identified syllables were matched across genotypes but not across genders. In order to register syllable identity across genders, for each female syllable depicted in Figure 5, the best-matching male syllable was identified by computing pairwise Pearson's correlation distances between that syllable and every available male syllable. Distances were computed over sets of scalar values derived from raw depth footage associated with a particular syllable such as 2-d, 3-d and angular velocity, mouse length, width, and height.

Identified syllables represent gross behavioral differences like running versus rearing, but also distinguish closely related behaviors like slow and fast running or low and high rears. In order to depict the extent to which behavior differs between wild-type and knock-out mice, pairwise syllable distances were computed between the 10 most upregulated syllables within each cohort as well as between the 10 most differentially upregulated syllables across cohorts and depicted as probability density distributions in Figure 5. If knock-out and wild-type mice exhibit gross and consistent behavioral differences, pairwise syllable distances, on average, will be greater across cohorts relative to within-cohort comparisons. Alternatively, if both knock-out and wild-type mice express similar behaviors, within and across-cohort pairwise distance distributions should be more similar.

**Pup interaction assay and spontaneous retrieval**—Naive virgin females were placed in a behavioral arena (30 cm × 46 cm, 31 cm high) without a top containing fresh standard bedding and were allowed to explore the arena freely during a 10-minute habituation period. After habituation, 3 pups of the same mouse strain aged P1-P3 were placed on one side, and 3 pup-sized novel plastic objects were placed on the opposite side of

the arena. Interactions of the female with the pups and novel objects were recorded for 10 minutes (as defined as a 2.5 cm radius around the nest or the center of the toys). After 10 minutes one pup was removed from the nest and placed in the center of the behavioral arena. If retrieval of the pup had not occurred within the observation period the trial was counted as a failure to retrieve and was terminated. Pups were promptly returned to their mothers after the experiment. Analysis was performed offline using MATLAB.

**Postpartum pup retrieval and cannibalization**—The dam was allowed to remain in her home cage and 3 pups were removed from the nest and placed in the center and 2 corners of the cage opposing the nest (see Figure 5A for schematic). The time to retrieve each pup to the nest was scored once a day over a time period of 3 consecutive days, starting on P0. Care was taken not to disturb the dam and her pups unnecessarily during this period. Pups were counted on P0-P3 and cannibalization was noted. Cannibalization mostly occurred on P0 and P1 and rarely thereafter.

**Pharmacological rescue of parental behavior**—For the rescue of pup interaction virgin females were either tested under control (regular drinking water) or test conditions (after administration of THIP for 24–48 h, THIP was dissolved in the drinking water at a dose of 5 mg/kg similar to a previously described paradigm (Maguire and Mody, 2008). The dose had only minor effects on locomotion, suggesting it was non-sedating. Administration with the drinking water precluded excessive handling and resulting stress in the animals. The pup interaction assay was otherwise conducted and analyzed as described above. Each female was tested under control and test conditions, and the order was randomly assigned to account for learning effects (ref). If the animal was tested first under test conditions, the animal was allowed a resting period of at least 72 h to ensure clearance of THIP from the body. Drinking water consumption was recorded before each behavior experiment (control water consumption  $8.0 \text{ mL} \pm 0.9 \text{ ml/per mouse/24 h}$  THIP water consumption  $8.9 \text{ mL} \pm 1.9 \text{ ml/mouse/24 h}$ ,  $p > 0.6$ , Student's t test)

For the rescue of postpartum maternal behavior timed-pregnant dams were randomly assigned to either the control or THIP group. THIP was administered with the drinking water 72 – 48 h before birth until 48 h after birth. Only first-time pregnancies were scored. Postpartum retrieval and pup cannibalization were recorded from P0 – P3 as described above.

**Forced swim**—Mice were placed in a glass cylinder (height, diameter) filled with water at room temperature and observed for a 6-min period. Animals were then removed from the beaker, towel dried, and returned to their home cage. Videos were scored offline. Time spent mobile (swimming, actively struggling) and immobile (floating, with front paws and at least one hind paw immobile) were scored.

**Olfactory testing**—To assay odor discrimination we performed a simple olfaction task. Animals were introduced to a clean, covered cage without bedding, food or water. Three neutral, non-social odors (coconut, raspberry, banana), 2 social scents (female and male urine), as well as a water sample were pipetted onto a clean cotton swab and introduced consecutively to the cage at random order. Mice were allowed to explore each scent for 5

min, and time spent investigating (sniffing, biting or chewing) was recorded live by the investigator.

**Whole brain c-Fos expression analysis**—Control and cb 6 KO littermates were single-housed and either exposed to an acute stressor for 30 min (restraint in a DecapiCone, Braintree Scientific, #MDC-200), or remained in their home cage until sacrifice. After restraint, the animals were allowed to recover in their home cages for 60 min, after which they were anesthetized with isoflurane in their home cage. The thoracic cavity was opened and blood was withdrawn from the left ventricle of the heart and transferred to a heparin-coated vial (BD microtainer #365965) on ice. Blood plasma was reserved for further analysis. After blood collection, the animals were perfused transcardially with PBS and 4% PFA. Brains were removed and stored in 4% PFA at 4 degrees overnight. Brains were then sliced on a vibratome (Leica VT 1000S) at 40  $\mu$ m thickness. Immunostaining for c-Fos was performed as described above (dilution of c-Fos primary antibody 1:1000 for 48 h at 4°C, #2250, Cell Signaling, Alexa 647 or Alexa 488 secondary antibody, 1:1000 dilution for 24 h at 4°C). Slices were then washed, mounted, and coverslipped. Images were taken on an Olympus VS120 slide scanner, and confocal stacks were acquired on an Olympus FV1000 confocal microscope. Raw images were processed for quantification of c-Fos expression in ImageJ. The Allen Brain Institute reference atlas ([allenbrain.org](http://allenbrain.org)) was used to identify brain regions across sections. Regions of interest were thresholded to create a mask outlining c-Fos positive nuclei, and particles consistent with nuclear shape (diameter, circularity) were counted. Heatmaps represents values of c-Fos particles per region normalized to control conditions (no stress), and the ratio (cb KO stress/control stress)/control stress. C-Fos expression was quantified in tissue of two animals per condition (total of 8 animals).

## QUANTIFICATION AND STATISTICAL ANALYSIS

Electrophysiology data analysis was performed in Igor Pro (Wavemetrics), AxographX (Axograph) and Prism (Graphpad). The numbers of cells recorded are indicated in the figure legends and in the text. To determine significance in a dataset the Whitney-Mann test, Wilcoxon signed rank test or Kruskal-Wallis test (with Dunn's post-test) were used, as indicated. For select datasets, a one-sample t test (Western blot analysis), or one-way ANOVA (gait analysis, with Dunnett's multiple comparison post-test), were performed. Behavioral analysis was performed in MATLAB (Mathworks) using custom written scripts, and Prism. Unpaired Student's t test was used to determine statistical significance. The number of animals used for each dataset is indicated in the text or figure legend.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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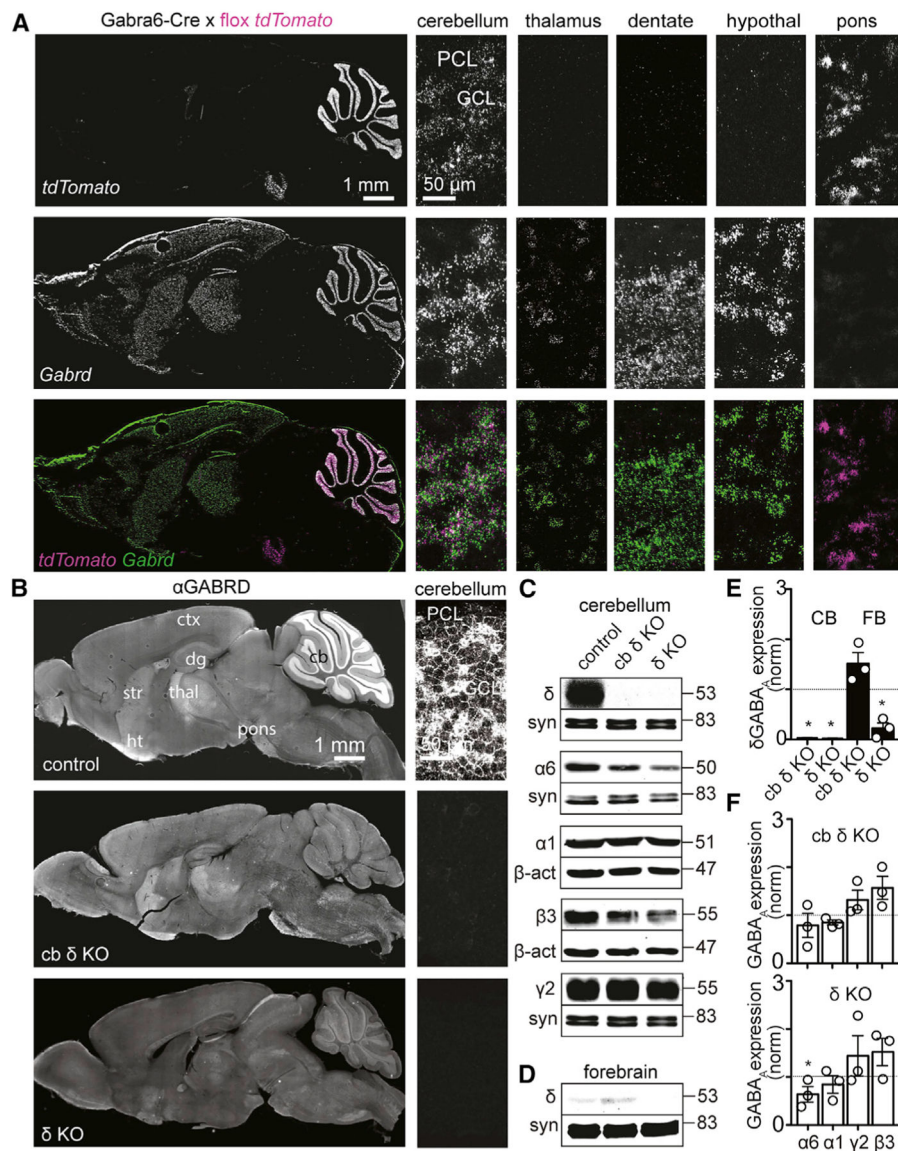


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**Highlights**

- $\delta$ GABA<sub>A</sub> deletion in cerebellar granule cells causes input layer hyperexcitability
- Hyperexcitability leads to increased anxiety- and stress-related behaviors
- Females lacking cerebellar  $\delta$ GABA<sub>A</sub> show deficits in social and maternal behavior
- Hyperactivation of granule cells during stress alters activity in many brain regions



**Figure 1. Specific Deletion of the  $\delta$ GABA<sub>A</sub>R in cb GCs**

(A) FISH labeling of *tdTomato* transcripts in the *Gabra6-Cre* mouse line crossed to the Ai14 reporter line (top), *Gabrd* transcripts (center, *Gabrd*), and merged images (bottom). Left: sagittal section of the whole brain. Right: confocal images of the cerebellum, thalamus, dentate gyrus, hypothalamus, and pons. *tdTomato* and *Gabrd* transcripts are co-expressed exclusively in cb GCs.

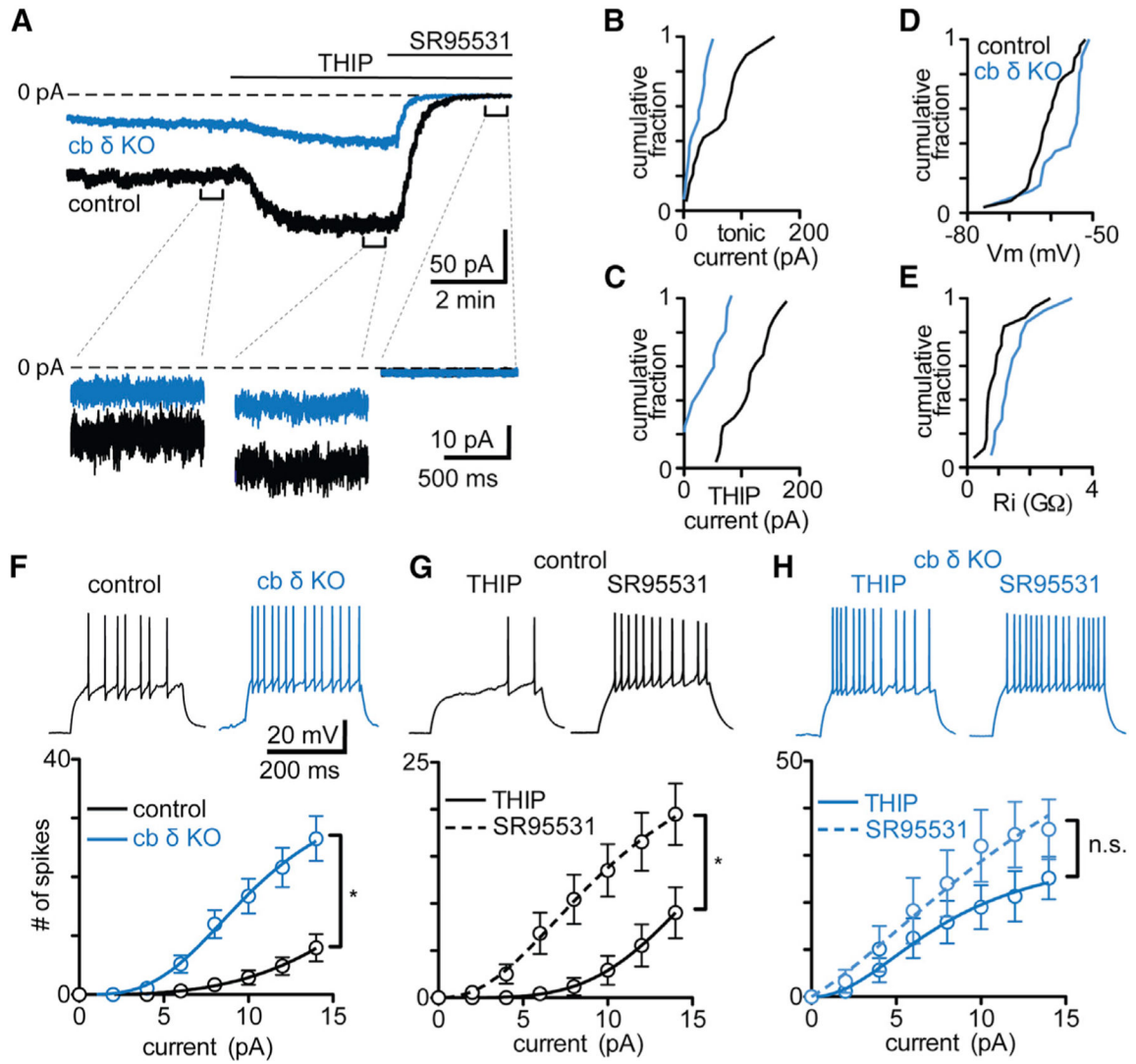
(B) Immunostaining against  $\delta$ GABA<sub>A</sub> in sagittal sections of the brain (left) and confocal images of the cerebellum (right) in control (top), cb  $\delta$  KO (center), and global  $\delta$  KO tissue (bottom).

(C) Western blot of the cerebellum against the  $\delta$ GABA<sub>A</sub>,  $\alpha 6$ GABA<sub>A</sub>,  $\alpha 1$ GABA<sub>A</sub>,  $\beta 3$ GABA<sub>A</sub>, and  $\gamma 2$ GABA<sub>A</sub> subunits; loading controls were synapsin (syn) or  $\beta$ -actin (bottom lanes).

(D) Western blot of the forebrain against  $\delta$ GABA<sub>A</sub>R.

(E) Quantification of  $\delta$ GABA<sub>A</sub> expression levels in the cerebellum (white bars) and forebrain (black bars) in cb  $\delta$  KO (left) and global  $\delta$  KO mice.  $\delta$ GABA<sub>A</sub> expression is abolished in the cerebellum of cb  $\delta$  KO and  $\delta$  KO mice and in the forebrain of  $\delta$  KO mice but not in the forebrain of cb  $\delta$  KO mice (one-sample t test of samples normalized to control, n = 3 animals, p < 0.05). Error bars represent SEM.

(F) Quantification of  $\alpha$ 6GABA<sub>A</sub>,  $\alpha$ 1GABA<sub>A</sub>,  $\beta$ 3GABA<sub>A</sub>, and  $\gamma$ 2GABA<sub>A</sub> expression levels in the cerebellum (white bars) of cb  $\delta$  KO mice (center) and global  $\delta$  KO mice (bottom).  $\alpha$ 6GABA<sub>A</sub> expression was decreased significantly in the cerebellum of  $\delta$  KO mice (p < 0.05). All other receptors did not show significant changes in expression levels (one-sample t test of samples normalized to control, n = 3 animals, p > 0.05). Error bars represent SEM.



**Figure 2. Hyperexcitability and Decreased Tonic Inhibition in GCs Lacking the GABA Receptor  $\delta$  Subunit**

(A) Top: example traces of whole-cell voltage-clamp recordings of tonic currents measured during a baseline period, in the presence of the  $\delta$ GABA<sub>A</sub> subunit containing-preferring GABA<sub>A</sub>R agonist THIP, and in the presence of the non-selective GABA<sub>A</sub>R antagonist SR95531 for a control (black) and a *cb δ KO* GC (blue). Bottom: current traces on an expanded timescale show varying noise levels depending on the recording conditions. SR95531 strongly reduces noise levels compared with the control and THIP.

(B) Cumulative histograms of initial tonic currents in control (black line) and *cb δ KO* GCs (blue line).

(C) Cumulative histograms in the presence of THIP.

(D) Cumulative histograms of membrane potential ( $V_m$ ).

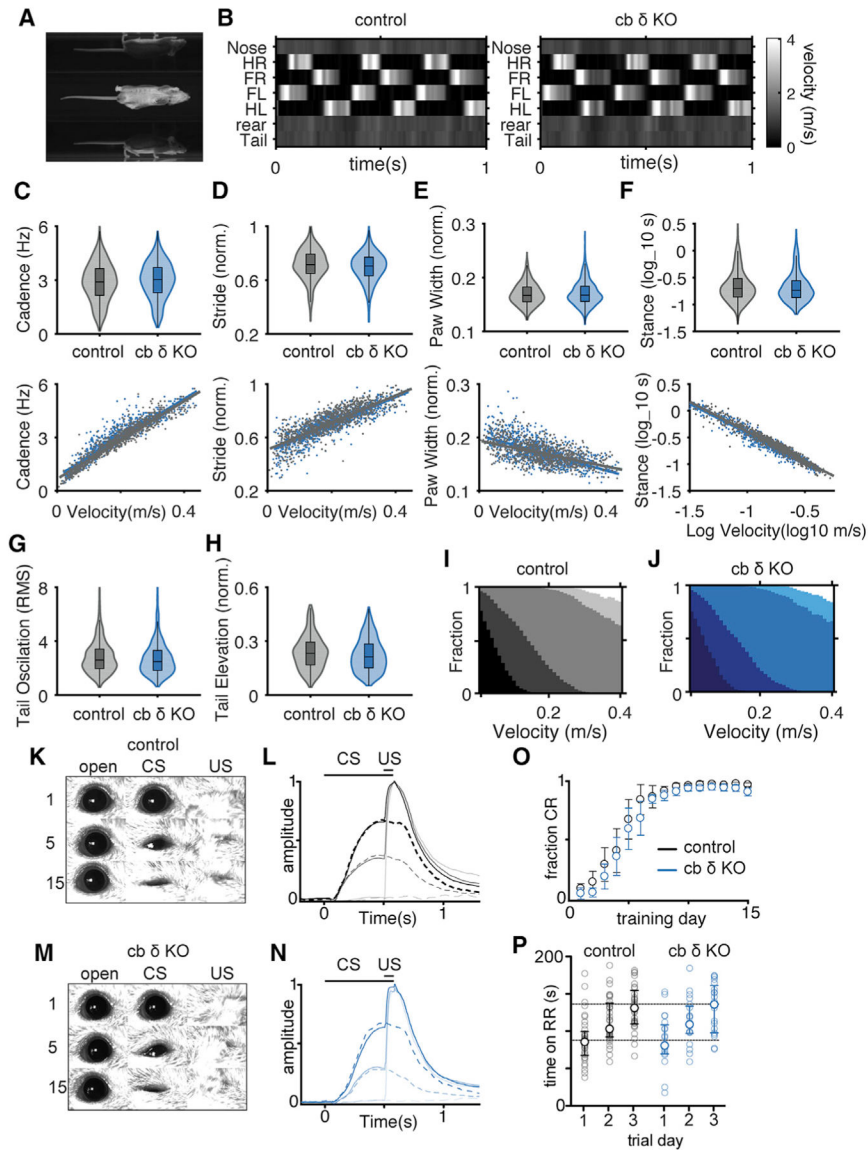
(E) Cumulative histogram of input resistance ( $R_i$ ).

(F) Top: example traces of whole-cell current clamp recordings in response to a 10-pA depolarizing current injection in *cb δ KO* GCs (blue) and in control GCs (black). Bottom:

summary data showing the input-output relationships for a range of depolarizing current steps and the resulting number of action potentials. Error bars represent SEM.

(G) As in (F) but for control animals in the presence of THIP and with GABA<sub>A</sub>Rs blocked. Error bars represent SEM.

(H) As in (G) but for cb  $\delta$  KO mice. Error bars represent SEM.



**Figure 3. Normal Motor Function and cb Learning in Cerebellar  $\delta$  KO Mice**  
 (A) Sample frame of a walking mouse viewed from the left, right, and bottom.  
 (B) Velocity of different body parts over time in control and cb  $\delta$  KO mice.  
 (C–F) Top: box-and-whisker plots overlaid on the density plots for cadence (C), stride (D), paw width (E), stance (F), tail oscillation (G), and tail elevation (H), measured stride to stride and pooled across all subjects, with control shown in gray and cb  $\delta$  KO in blue. Bottom: scatterplots and linear regression of cadence, stride, paw width, and stance against velocity.  
 (I and J) Fraction of time during a single gait cycle for a given velocity that had four (darkest) to zero (white) supports on the floor.  
 (K) Representative images of eyeblink responses in a control animal sampled at the beginning (day 1, top), in the middle (day 5, center), and at the end of training (day 15,



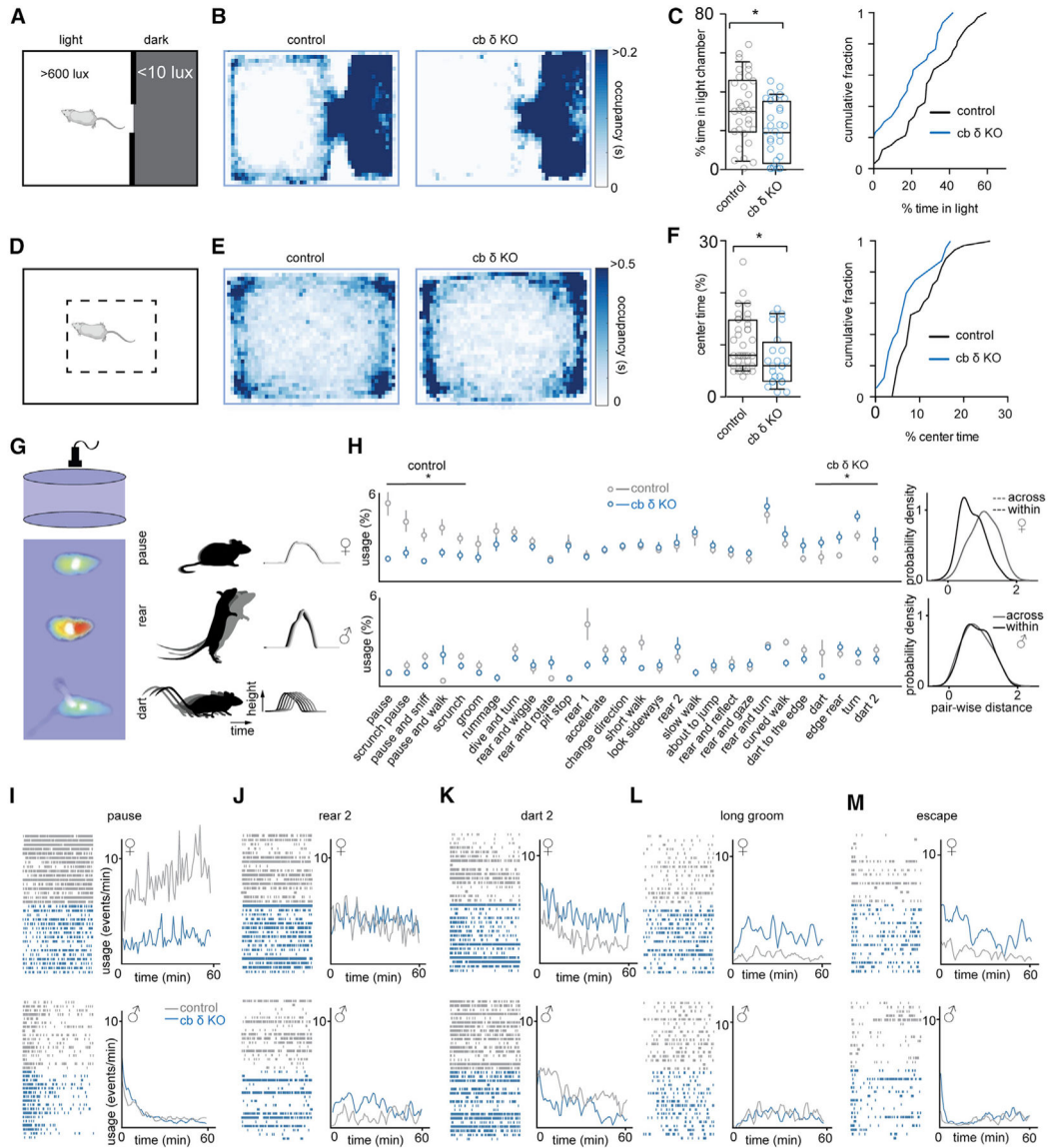
bottom). The baseline, conditioned response (CR) and air puff response (UR) are presented in sequence column-wise. Initially (day 1), the eye remains fully open during presentation of the CS and closes fully only with presentation of the US. With continued learning (day 5), the mouse begins to close its eye in response to the CS until near-full closure at the end of the training period (day 15). The US always elicits complete closure of the eye.

(L) Average normalized eye closure from early (day 1, light gray), middle (day 5, gray), and end of training (day 15, dark gray).  $t = 0$  marks the presentation of the CS and 0.5 s presentation of the US. Solid lines represent paired presentation of CS and US at different learning stages, and dashed lines represent CS presentation only.

(M and N) Analogous to (K) and (L) but in  $cb \delta$  KO animals.

(O) Fraction of trials with a CR on each training day.  $Cb \delta$  KO animals (blue circles) learn at a similar rate as controls (white circles) in an eyeblink task (control,  $n = 8$ ;  $cb \delta$  KO,  $n = 8$ ). Data are presented as mean and SEM.

(P) Motor learning, as assayed in a 3-day accelerating rotarod paradigm, is similar in control (black) and  $cb \delta$  KO animals (blue; control,  $n = 36$  animals;  $cb \delta$  KO, 20 animals;  $p > 0.9$ , Mann-Whitney test). Gray circles represent individual animals.



**Figure 4. Anxiety-like Behavior and Sex-Specific Behavioral Abnormalities in *cb δ* KO Mice**

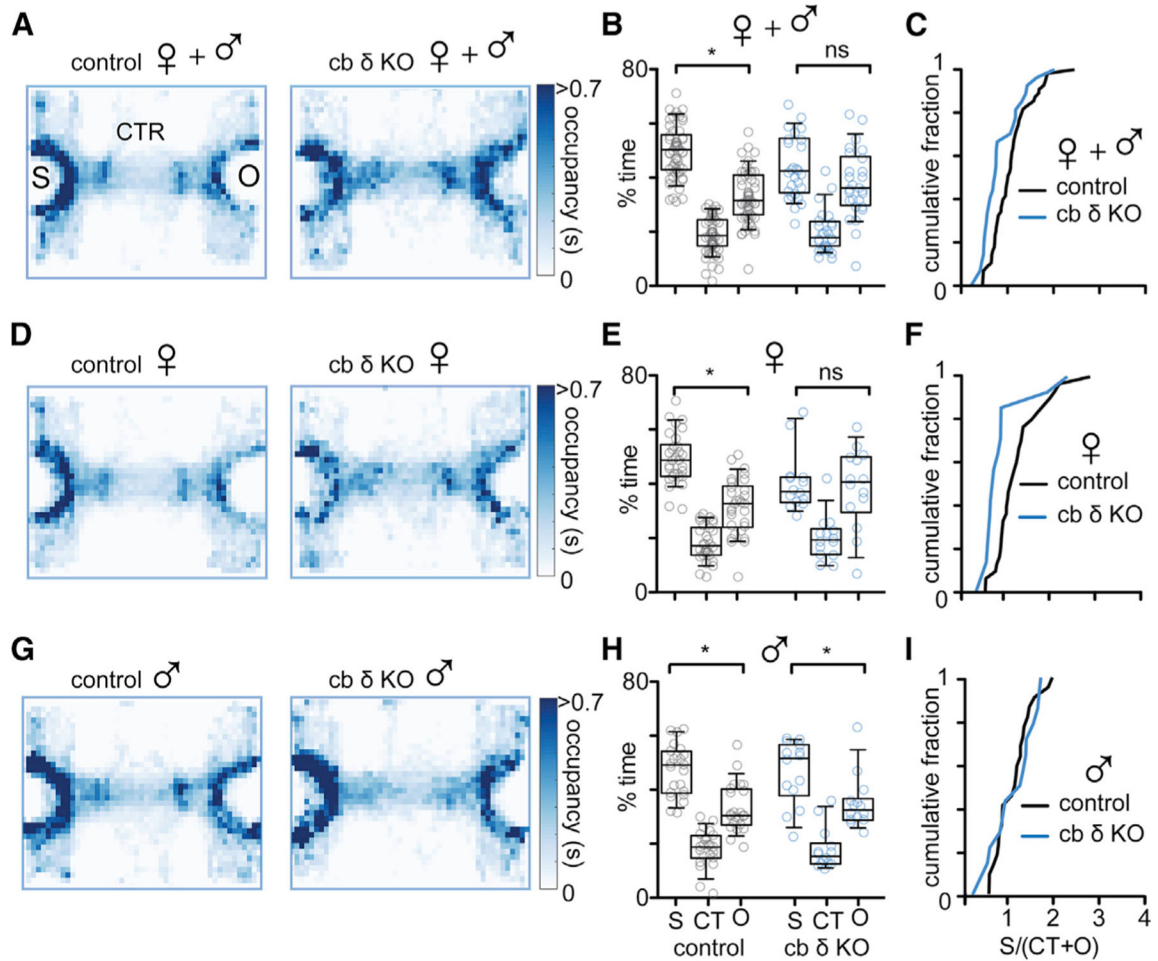
(A) Schematic of the light/dark chamber. Mice are placed in a behavioral arena divided into a dark compartment (<10 lux) and a lit compartment (>600 lux). A door connects the two chambers, allowing navigation between both compartments.

(B) Median occupancy plots indicate that *cb δ* KO animals (right) spend less time in the lit compartment than control animals (left).

(C) Summary data of the percentage of time spent in the light compartment. Left: boxes denote interquartile range and median; whiskers represent 10–90 percentile. Circles show individual control (n = 32, gray circles) and *cb δ* KO (n = 27, blue circles) animals (p < 0.009, Mann-Whitney test). Right: cumulative fraction of time spent in the light compartment (control, black line; *cb δ* KO, blue line).

(D) Schematic of the open field behavior arena. A dashed rectangle denotes the center of the area.

- (E) Median occupancy plots of control (left) and cb  $\delta$  KO animals (right).
- (F) Left: summary data of the percentage of time spent in the center of the behavioral arena. Circles show individual control (n = 36, gray circles) and cb  $\delta$  KO (n = 24, blue circles) animals ( $p < 0.02$ , Mann-Whitney test). Right: cumulative probability graph of center time. Boxes denote interquartile range and median; whiskers represent 10–90 percentile.
- (G) 3D imaging of mice in the open field with a depth camera, followed by MoSeq-based segmentation of the behavior data, reveals behavioral syllables (STAR Methods); example behavioral syllables and the associated inferred positions of the spine in time and space are also shown (center and right).
- (H) Left: usage plot for the most frequently used syllables in females (top). A subset of behavioral syllables is differentially expressed in control and cb  $\delta$  KO females. Male syllables were matched to corresponding female syllables (bottom). Control and cb  $\delta$  KO males showed fewer differences in behavior. Right: pairwise distance comparison across control and cb  $\delta$  KO syllables and within control and cb  $\delta$  KO syllables reveals greater distance across control and cb  $\delta$  KO syllables in females and similar distances across and within male syllables.
- (I) Example syllables performed more frequently by control females. Top left: raster plot of syllable occurrence during the observation period. Right, syllable usage over time. The observed behavior associated with each syllable is indicated. Bottom: raster plot and usage over time of matching male syllables.
- (J) Like (H) but depicting common example syllables that are equally prevalent in control and cb KO mice.
- (K) Like (H) but depicting common example syllables performed more frequently by cb KO females.
- (L and M) Like (H) but depicting rarely used syllables that are more frequently used by cb KO females.



**Figure 5. Sexually Dimorphic Effect of Cerebellum-Specific  $\delta$ GABA<sub>A</sub> Deletion on Social Behavior**

(A) Median occupancy plots indicate a clear preference of control (left) animals for the S over the O, whereas cb  $\delta$  KO (right) animals show no preference for either stimulus.

(B) Summary data for the percentage of time spent in the O, center, and S compartment of the behavior chamber during the observation period. On average, control animals display a strong preference for the S (gray circles,  $p < 0.0001$ ), whereas cb  $\delta$  KO animals show no preference (blue circles). Boxes denote interquartile range and median; whiskers represent 10–90 percentile.

(C) Cumulative probability plot of the ratio of time spent with the S and the sum of the time spent in the center (CTR) and O chambers. Sociability ratio  $S/(O+C)$ ; control, black trace; cb  $\delta$  KO, blue trace.

(D–F) Same as (A)–(C) but females only.

(D) Median occupancy plots of control (left) and cb  $\delta$  KO females (right). Control females show a strong preference for S (left), whereas cb  $\delta$  KO (right) females do not show a preference for either stimulus.

(E) Summary data of the percentage of time spent in the S, CTR, and O compartments. Boxes denote interquartile range and median; whiskers represent 10–90 percentile.

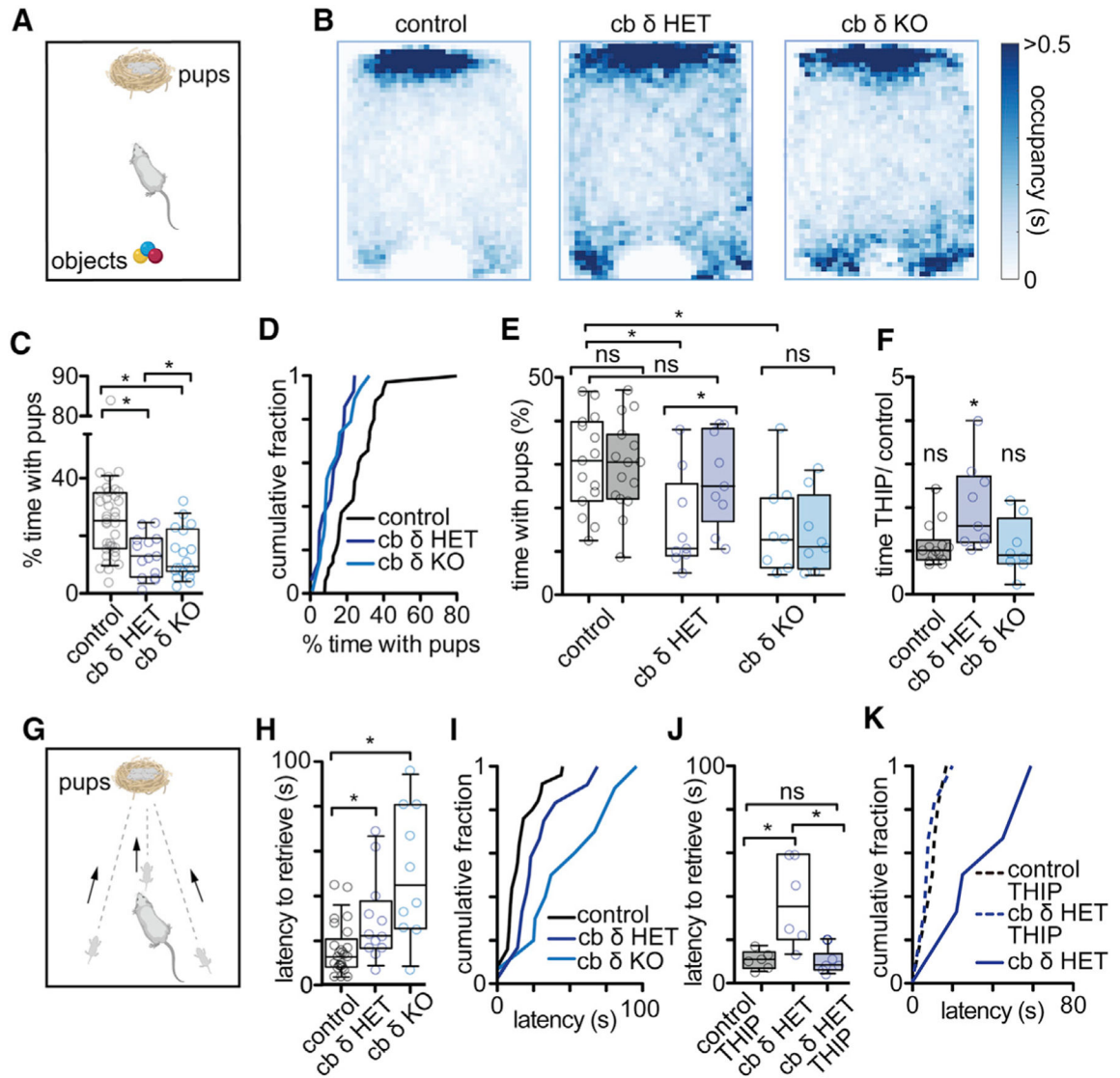
(F) Cumulative probability plot of sociability ratios in females. Boxes denote interquartile range and median; whiskers represent 10–90 percentile.

(G)–(I) Same as (A)–(D) but males only.

(G) Median occupancy plots of control (left) and cb  $\delta$  KO males (right). Control and cb  $\delta$  KO males prefer S over the O.

(H) Summary data of the percentage of time spent in the S, CTR, and O compartments.

(I) Cumulative probability plot of the sociability ratios in males.



**Figure 6. Altered Maternal Behavior in *cb δ* KO Females**

(A) Schematic of the behavioral paradigm. A virgin female is placed in a behavioral arena that contains a nest with three pups (top) and three pup-sized novel Os (bottom).

(B) Median occupancy plots of control, *cb δ* HET, and *cb δ* KO females. Control females preferentially remain in close proximity of the nest, whereas *cb δ* HET and *cb δ* KO mice disperse around the nest and explore Os.

(C) Summary plot of time spent with pups during the observation period. Circles show individual mice tested (control: gray,  $n = 37$ ; *cb δ* KO het: dark blue,  $n = 12$ ; *cb δ* KO: blue,  $n = 19$ ; Kruskal-Wallis test with Dunn’s post-test,  $p < 0.003$ ). Boxes denote interquartile range and median; whiskers represent 10–90 percentile.

(D) Cumulative probability plot of time spent with pups ( $p < 0.02$  control compared with KS *cb δ* HET,  $p < 0.03$  control compared with KS *cb δ* KO test,  $p > 0.9$  *cb δ* HET compared with *cb δ* KO).



(E) Pharmacological rescue of behavior in cb  $\delta$  HET virgin females. The time animals spent with pups was measured under control conditions and after administration of THIP. THIP increased the time that cb  $\delta$  HET females spent with pups but not had no effect on the time in cb  $\delta$  KO and control females spent with pups, control and control THIP (gray circles, shaded gray,  $n = 15$ ;  $p > 0.8$ ) cb $\delta$ HETandcb $\delta$ HET THIP (dark blue circles, shaded dark blue,  $n = 9$ ;  $p < 0.004$ ), and cb  $\delta$  KO and cb  $\delta$  KO THIP (light blue circles, shaded light blue,  $n = 8$ ;  $p > 0.6$ ); Wilcoxon matched-pairs signed rank test. Boxes denote interquartile range and median; whiskers represent 10–90 percentile.

(F) Ratio of time spent with pups before and during THIP administration, control (shaded gray,  $n = 15$ ,  $p > 0.9$ ), cb  $\delta$  HET (shaded dark blue,  $n = 9$ ,  $p < 0.004$ ), and cb  $\delta$  KO (shaded blue,  $n = 8$ ,  $p > 0.9$ ); Wilcoxon signed-rank test.

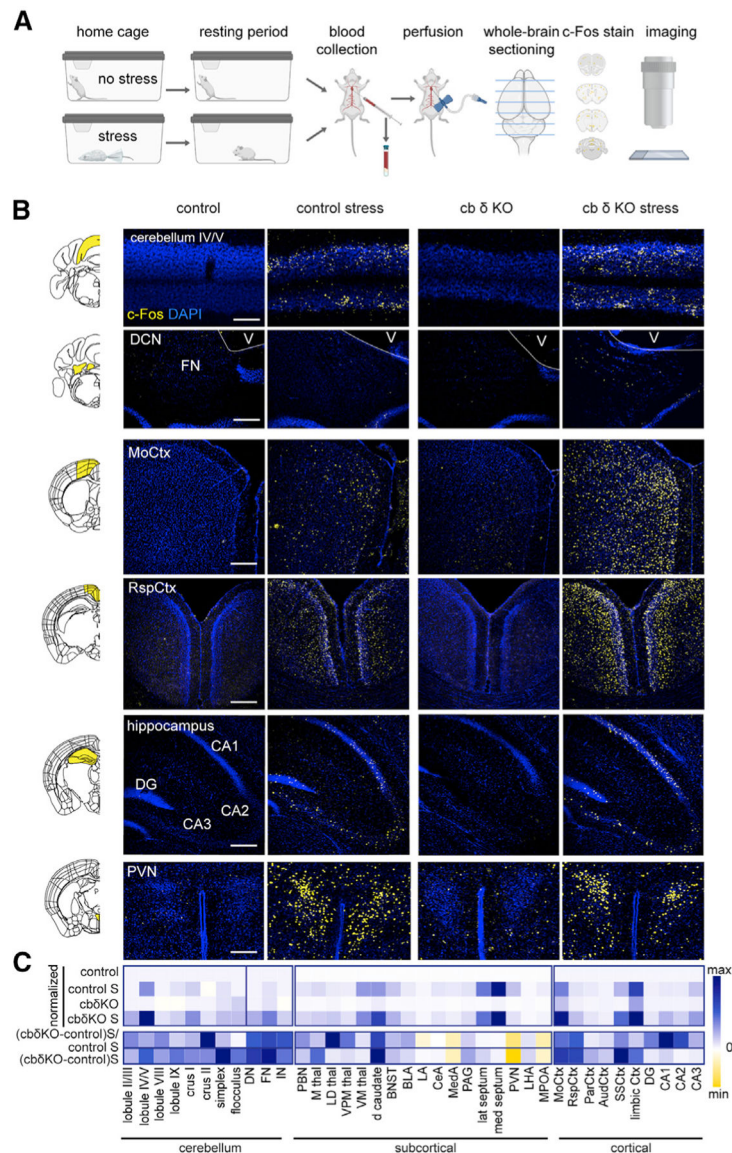
(G) Schematic of the retrieval paradigm. In the home cage, 3 pups are removed from the nest and placed in the opposite 2 corners and in the CTR of the cage.

(H) Summary plot of latency to retrieve the first pup. Circles show individual mice tested (control: gray,  $n = 25$ ; cb  $\delta$  KO het: dark blue,  $n = 12$ ; cb  $\delta$  KO: blue,  $n = 10$ ; Kruskal-Wallis test with Dunn's post-test,  $p < 0.008$ ). Boxes denote interquartile range and median; whiskers represent 10–90 percentile.

(I) Cumulative probability plot of retrieval times (KS test,  $p < 0.008$ ).

(J) Pharmacological rescue of retrieval latency in cb  $\delta$  HET postpartum females. Boxes denote interquartile range and median; whiskers represent 10–90 percentile.

(K) Cumulative probability plot of latency to retrieve in the presence and absence of THIP. Control ( $n = 6$ ) and cb HET ( $n = 6$ ),  $p < 0.02$ ; cb HET and cb HET THIP ( $n = 6$ ),  $p < 0.04$ ; control and cb HET THIP,  $p > 0.9$ ; KS test.



**Figure 7. Differential Activation of Diverse Brain Regions in cb  $\delta$  KO Animals in Response to Stress**

(A) Schematic of the experimental paradigm. Animals were subjected to 30 min of restraint stress or remained undisturbed in their home cage. After discontinuation of restraint, animals were allowed to recover for 60 min. Animals were then anesthetized, and trunk blood was taken from the left atrium of the heart before transcardial perfusion with a fixative. The whole brain was then sectioned, immunostained for c-Fos, and imaged for further analysis. (B) Left: reference atlas images of coronal brain sections (<https://mouse.brain-map.org/static/atlas>). The brain regions shown in the corresponding confocal images (right) are highlighted in yellow. Right: representative confocal images of unstressed and stressed control and cb  $\delta$  KO animals (blue, DAPI; yellow, c-Fos). Example brain regions include the cb cortex (lobule IV/V), deep cb nuclei (DCN); fastigial nucleus [FN], fourth ventricle [V]), motor cortex (MoCtx), retrosplenial cortex (RspCtx), hippocampus (CA1, CA2, and CA3 regions), and the paraventricular nucleus (PVN) of the hypothalamus.

(C) Top 4 panels: heatmap of normalized c-Fos expression. Acute restraint stress increased c-Fos expression in many cb, subcortical, and cortical brain areas. Bottom panel: difference and normalized difference heatmap of c-Fos expression in stressed cb  $\delta$  KO animals and stressed controls.

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## KEY RESOURCES TABLE

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Chemicals, Peptides, and Recombinant Proteins		
NBQX disodium salt	Abcam Cambridge, MA	Cat# ab120046
(R)-CPP	Abcam Cambridge, MA	Cat# ab120159
SR95531 (gabazine)	Abcam Cambridge, MA	Cat# ab120042
Strychnine HCl	Sigma Aldrich	Cat# S-8753
QX314 chloride	Abcam Cambridge, MA	Cat# ab120118
CGP 55845 hydrochloride	Abcam Cambridge, MA	Cat# ab120337
THIP hydrochloride (gaboxadol)	Abcam Cambridge, MA	Cat# ab120426
RNAscope fluorescent kit	ACD, Cambridge, MA	Cat# 320850
RNAscope Gabrd-Mm-C3 probe	ACD, Cambridge, MA	Cat# 459481-C3
RNAscope tdTomato- probe	ACD, Cambridge, MA	Cat# 317041
Anti-GABA <sub>A</sub> $\delta$ Antibody, rabbit	Alomone labs, Jerusalem, Israel	Cat# AGA-014; RRID: AB_2340938
Anti-rabbit Alexa 647	Abcam, Cambridge, MA	Cat# 150083; RRID: AB_2714032
Anti-rabbit Alexa 488	Abcam Cambridge, MA	Cat# 150081; RRID: AB_2734747
rabbit anti-GABA <sub>A</sub> receptor- $\alpha$ 1	Abcam	Cat# ab33299; RRID: AB_732498
rabbit anti-GABA <sub>A</sub> receptor- $\alpha$ 6	GeneTex	Cat# GTX130947
mouse anti-GABA <sub>A</sub> receptor- $\beta$ 3	Neuromab	Cat# 75-149; RRID: AB_2109585
rabbit anti-GABA receptor- $\gamma$ 2	Synaptic Systems	Cat# 224 003; RRID: AB_2263066
rabbit anti-GABA <sub>A</sub> receptor- $\delta$ 1:2000	gift from Dr. W. Sieghart and Dr. Petra Scholze, Medical University of Vienna, Austria	N/A
rabbit anti c-fos (9F6)	Cell Signaling Technology	Cat# 2250S; RRID: AB_2247211
Experimental models: Organisms/ Strains		
<i>floxed</i> Gabrd mice ( <i>Gabrd<sup>tm1.1Jmag/J</sup></i> )	Dr. Jamie Maguire, Tufts University Jackson Labs	Stock# 023836 (Lee et al., 2014)
Gabrd knock out mice ( <i>Gabrd<sup>tm1Geh/J</sup></i> )	Dr. Gregg Homanics, University of Pittsburgh Jackson labs	Stock# 003725 (Mihalek et al., 1999)
Gabra6-cre mice B6.D2-Tg(Gabra6-cre) B1Lfr/Mmucd	MMRC	RRID: MMRRC_015966-UCD (Fünfschilling and Reichardt, 2002)
Floxed tdTomato reporter line (Ai14, Gt(ROSA)26Sor <sup>tm14(CAG-tdTomato)Hze/J</sup> )	Jackson Labs	Stock# 007908 (Madisen et al., 2010)
Software and Algorithms		
Igor Pro 6	Wavemetrics	<a href="https://www.wavemetrics.com/">https://www.wavemetrics.com/</a>
MafPC	Courtesy of M.A. Xu-Friedman	<a href="https://www.xufriedman.org/mafpc">https://www.xufriedman.org/mafpc</a>
MATLAB (R2017a)	MathWorks	<a href="https://www.mathworks.com/">https://www.mathworks.com/</a>
Fiji (ImageJ)	NIH	<a href="https://fiji.sc/">https://fiji.sc/</a>
AxographX 1.7.0	Axograph	<a href="https://axograph.com/">https://axograph.com/</a>
Prism 6	Graphpad	<a href="https://www.graphpad.com/">https://www.graphpad.com/</a>
Adobe Illustrator	Adobe	<a href="https://www.adobe.com/products/illustrator.html">https://www.adobe.com/products/illustrator.html</a>

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Moseq2	Sandeep Datta Laboratory	<a href="https://github.com/orgs/dattalab/teams/moseq2-users">https://github.com/orgs/dattalab/teams/moseq2-users</a>
Python	Python	<a href="https://www.python.org/">https://www.python.org/</a>
Biorender	Biorender	<a href="https://biorender.com/">https://biorender.com/</a>

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