Cerebellar climbing fibers impact experience-dependent plasticity in the mouse primary somatosensory cortex

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Summary

In the cerebellum, climbing fibers (CFs) provide instructive signals for supervised learning at parallel fiber to Purkinje cell synapses. It has not been tested so far whether CF signaling may also influence plasticity in other brain areas. Here, we show that optogenetic CF activation suppresses potentiation of whisker responses in L2/3 pyramidal cells in primary somatosensory cortex (S1) of awake mice that is observed after repeated whisker stimulation. Using two-photon imaging and chemogenetics, we find that CFs control plasticity by modulating SST- and VIP-positive interneurons in S1 cortex. Transsynaptic labeling identifies zona incerta (ZI) to thalamic posterior medial nucleus projections as a pathway for cerebellar output reaching S1 cortex. Chemogenetic inhibition of PV-positive neurons in the ZI prevents CF co-activation effects, identifying the ZI as a critical relay. Our findings demonstrate that CFs impact sensory signal processing and plasticity in S1 cortex and thus may convey instructive signals.

Keywords:

Calcium, Instructive signal, Parvalbumin-positive interneuron, Posterior medial thalamic nucleus, Purkinje cell, Pyramidal cell, Somatostatin-position interneuron, Supervised learning, Vasoactive intestinal polypeptide-positive interneuron, Zona incerta

Introduction

Sensory experience and learning are associated with plasticity of multiple brain regions.
Neocortical plasticity canonically depends on the activity of its direct input structures[1]. For
example, plasticity in the barrel field of the primary somatosensory cortex (S1 cortex) of rodents
is evoked upon repeated whisker stimulation. Receptive field maps may reorganize after whisker
clipping[2-4] or a history of rhythmic whisker stimulation[5-7] that necessitates adaptive plasticity of
synaptic weights, e.g. via long-term potentiation (LTP) and/or long-term depression (LTD) at
synaptic inputs onto pyramidal neurons[8, 9]. The high degree of plasticity in S1 somatosensory
circuits enables adaptation to altered inputs[10]. Though experience-dependent plasticity in
neocortex is highly responsive to input features, modulating influences from other brain regions
may play a role.

The cerebellum is increasingly assigned functions that were previously attributed to the
neocortex[11, 12]. It projects via the thalamus to motor cortex and non-motor areas, including the
prefrontal cortex[13, 14], sensory, and associative areas[15-17]. A plausible possibility is therefore that
cerebellar output impacts cortical signaling. It has indeed been shown that modulation of
cerebellar outputs influences activity in the frontal cortex[18], coherence of neuronal oscillations
between neocortical areas[19-21], and plasticity in parietal-motor cortex connections[22]. However,
the cellular mechanisms of these interactions have not been described.

Here, we ask the specific question of whether cerebellar climbing fiber (CF) signals have the
mechanistic capacity to impact neocortical plasticity. Our motivation to study CF signaling is
rooted in the importance of their activity for normal cerebellar function, where CF-controlled
plasticity represents a classic example of supervised learning in the brain[1]. Depending on
dendritic architecture, Purkinje cells (PCs) receive input from one or more climbing fibers
(CFs)[23] that originate in the contralateral inferior olive (IO). As originally predicted by Marr in
his *Theory of Cerebellar Cortex*, CF signaling supervises plasticity at parallel fiber (PF) – PC
synapses, providing *contexts* for learning[24]. It was later experimentally demonstrated that PF and
CF co-stimulation initiates LTD at PF synapses[25-28], and that the CF – via evoked calcium
transients – tightly controls plasticity at these synapses[29-34] and provides instructive signals for
cerebellum-controlled learning[35-38]. CFs provide instructive error signals[39], convey sensory input
across modalities[40-44], and may even signal the absence of expected sensory signals[45]. CFs may
also carry reward or reward prediction-related signals[46, 47]. Contexts in which CFs are recruited
continue to be identified, underscoring the importance of establishing whether they may
influence neocortical activity and plasticity.

To determine whether CF signaling has the capacity to impact plasticity of S1 neurons, we use
the mouse brain to give us experimental access at the cellular level. Our observation that CF
signaling regulates plasticity of L2/3 pyramidal cells in S1 cortex shows that activity in the
olivo-cerebellar system has distinct consequences for input processing and plasticity in S1
cortex, and significantly expands what is known about the basic interaction between these
regions. This mechanism could be used to provide signals to neocortex that are instructive in
nature, depending on the context in which CFs are recruited, similar to those signals that play a
role in cerebellar supervised learning.

Activity-dependent plasticity in S1 cortex is regulated by optogenetic CF activation

To study the effects of CF activity on S1 plasticity, we expressed GCaMP6f in neurons in S1
cortex and used two-photon microscopy in awake mice to measure responses to air puff
stimulation of the whisker field in layer 2/3 (L2/3) neurons (Figure 1A, D). For optogenetic CF
activation, channelrhodopsin-2 (ChR2) was expressed in neurons of the inferior olive (IO),
which give rise to CFs terminating in the contralateral cerebellar cortex (Figure 1B, E; Figure
S1A, B). An LED was used to deliver 470nm light pulses over an optical window centered on
crus I/II, an area of the cerebellar cortex responsive to stimulation of the ipsilateral whisker
field[40]. The efficacy of optogenetic activation was tested by expressing GCaMP6f in PCs of crus
I/II and measuring calcium transients evoked by LED light pulses, which we varied in frequency
and duration to achieve optogenetically evoked signals resembling spontaneous CF transients
(Figure 1C; Figure S1F). Single light pulses lasting 50ms evoked CF transients which were not
significantly different in amplitude or peak latency from spontaneous events recorded in the
same PCs (Figure S1H, I)[48]. For plasticity induction in S1 cortex, we used a multi-whisker
stimulation protocol ('rhythmic whisker stimulation'; RWS) similar to those previously used to
study sensory-driven synaptic plasticity *in vivo*[7, 9, 49, 50]. Here, RWS consisted of an application of
100ms air puffs (8psi) to the contralateral whisker field for 5min at 8Hz, a natural whisking
frequency for mice that are sampling the environment[51]. RWS stimulation resulted in an increase
in whisker-evoked calcium responses ($\Delta F/F$ over the range of 0-700ms post-stimulus onset) that
we observed in morphologically identified L2/3 pyramidal neurons (PNs; Figure 1F) for the
duration of the recording (60min post-RWS; Figure 1G-I). The somatic calcium response
amplitudes measured with GCaMP6 correlate with neuronal spike rates[52]. L2/3 pyramidal
neurons respond to stimulation of the corresponding principal whisker but may also respond to
stimulation of surround whiskers[53]. Therefore, the plasticity observed here in response to multi-
whisker stimulation may reflect both a potentiation of synapses conveying already existing
whisker input as well as an expansion of the neuron's receptive field to include additional
surround whiskers. Given increases in the activity of excitatory neurons may be coupled with
decreased activity of local inhibitory neurons in this form of plasticity[54], we also assessed the
responses of neurons that were not identified as pyramidal cells (i.e. putative interneurons; INs)
in the same mice. Indeed, INs collectively responded to RWS stimulation with a significant
depression in the amplitude of evoked calcium events (Figure 1J-L; Figure S2F, G). As
parvalbumin (PV)-positive interneurons, somatostatin (SST)-positive interneurons, and
vasoactive intestinal polypeptide (VIP)-positive interneurons make up ~40%, ~30%, and ~12%
of the neocortical interneuron population[55, 56], respectively, this plasticity effect is likely
comprised of a substantial portion of SST and PV interneurons. Using transgenic mice to tag
SST and PV interneurons with tdTomato, we demonstrate RWS causes a significant depression
in SST and PV interneurons (Figure S2H, I) – consistent with our observations in putative
interneurons.

To determine whether CF co-activation with whisker stimulation influences this form of
experience-dependent plasticity, 50ms light pulses were delivered at 1Hz during the 5min period
of repeated whisker stimulation (RWS+CF), which is the rate of spontaneous CF discharge[39].
Each light pulse was delivered with a delay of 45ms with respect to the onset of whisker stimuli,
mimicking the natural latency of CF responses to whisker stimulation[40]. RWS+CF blocked the
potentiation of L2/3 pyramidal cell responses in S1 cortex and depression was observed instead
(Figure 1G-I; Figure S2D, E). The depression of IN responses observed after RWS was absent
upon optogenetic CF co-activation (Figure 1J-L, Figure S2F, G), and recordings in mice with
tdTomato-tagged SST and PV interneurons further demonstrated the absence of inhibitory

interneuron depression (Figure S2H, I). Importantly, both the S1 plasticity and optogenetic CF
activation effects were observed when trials with active movement of the mouse were included
(Figure 1; Figures S2+3) or when analysis was restricted to trials during which the mice were
resting both before stimulus onset and throughout the entire response period (Figure S3). Note
movements of the vibrissae were absent during rest.

To test to robustness of these effects, we reproduced these findings using an alternate
methodology that is well suited to detect receptive fields, including barrels in S1 cortex. Intrinsic
optical imaging[57] was used to measure plasticity effects at low spatial resolution in anesthetized
mice to facilitate measurement of single whisker responses. We slipped a glass pipette over an
individual untrimmed whisker and moved it back and forth at 10Hz for a period of 5min. In the
barrel field of contralateral S1 cortex, we confirmed the activation of the corresponding barrel
and recorded responses to this passive sensory experience for 20min before and 30min after
repeated stimulation. We observed an increase in the responsive area beyond the barrel that was
not seen in the absence of repeated stimulation (Figure S4). When ChR2-expressing CFs in crus
I/II of the cerebellum were optogenetically activated at 1Hz for 5min during the period of
repeated stimulation, this plasticity was not observed in S1 cortex (Figure S4). Thus, intrinsic
optical imaging demonstrates receptive field plasticity of barrels can be regulated by cerebellar
signaling.

Taken together, our findings demonstrate that whisker-induced response potentiation in S1
cortex results from L2/3 pyramidal cell plasticity. Local inhibitory interneurons show the
opposite direction of plasticity in the same animals. The observation that this IN plasticity is of
lower amplitude and does not perfectly mirror pyramidal cell plasticity suggests that the former
supports the latter and may help to control it, but that pyramidal cell LTP may take place even in
the absence of LTD in IN neurons. CF activity regulates this form of whisker map plasticity.
Note that 'LTP' and 'LTD' here reflect target neuron responsiveness – as measured by neural
calcium signals – and are not specifically shown to rest on synaptic gain changes.

CF activation modulates inhibitory interneurons in S1 cortex

Local inhibitory interneurons regulate the activity and plasticity of neocortical pyramidal
neurons, phenomena studied extensively in the rodent barrel cortex[56]. Thus, to identify the
potential mediators of the gating effect of optogenetic CF activation on S1 plasticity, we next
determined how the basic whisker response properties of specific neuron types in the S1 circuit
are altered during CF co-activation (Figure 2A). To begin, we demonstrated that these basic
responses are not significantly changed in the population of examined L2/3 pyramidal neurons
upon optogenetic CF activation (Figure 2B; 0-700ms analysis window). However, we found that
in 11/23 mice, an overall response decrease resulted from CF co-activation, while in 5/23 mice
an increase was seen (using a 10% threshold). Significance in the observed mild response
reduction emerged in the analysis across all neurons when a late analysis window was selected
(650-850ms, selected as an early component of the post-peak response). These findings suggest
that CF co-activation causes a range of response changes in the population of L2/3 pyramidal
neurons, but that an overall suppressive effect becomes apparent in the late response phase.

In the same mice, the basic responses of INs to whisker stimulation were increased upon CF co-
activation (Figure 2C). Like the plasticity effects measured in the IN populations (Figure 1J-L),
these basic response increases are likely comprised of a majority of PV and SST interneurons.

Using tdTomato-tagged interneurons to identify SST and PV interneurons (Figure 2D-F), we
showed that basic responses to whisker stimulation were indeed enhanced in SST and PV
interneurons upon CF co-activation (Figure 2G-I). VIP interneurons – which primarily inhibit
other interneurons including SSTs and PVs[56] – showed no consistent change in their basic
responses to whisker stimulation upon CF co-activation in the default analysis window (0-
700ms; Figure 2G). However, like pyramidal neurons (Figure 2B), VIPs demonstrated
significant suppression in the late analysis window of 650-850ms (Figure 2G).

Opposing roles of SST and VIP interneurons in CF-mediated control of S1 plasticity
As SST interneurons are activated by CF co-activation, we tested whether they assume a key role
in the CF-mediated regulation of S1 plasticity. To do so, we next used chemogenetic approaches
for activity manipulation. We focused on SST interneurons because they contact the dendrites of
L2/3 pyramidal cells (Figure 3A, E), where inhibition can impact local synaptic plasticity
processes. We first expressed the excitatory hM3D(Gq) DREADD in neocortical SST
interneurons (Figure 3B). Administration of the DREADD agonist deschloroclozapine (DCZ) by
intraperitoneal injection caused an increase in SST interneuron responses (Figure 3C). Under
these conditions, RWS did not potentiate L2/3 pyramidal cell responses and calcium signals were
depressed instead (Figure 3D), mimicking the results of optogenetic CF co-activation (Figure
1G-I). In contrast, inhibitory hM4D(Gi) DREADD expression in SST interneurons (Figure 3F)
and acute administration of DCZ caused a significant reduction in SST interneuron responses
(Figure 3G). This rescued potentiation of L2/3 pyramidal cell responses when RWS was paired
with CF co-activation (Figure 3H). These findings confirm that SST interneurons are positioned
to control plasticity of L2/3 pyramidal neurons[9] and suggest that this mechanism is recruited by
CF co-activation.

SST interneurons receive relatively weak input from cortical and subcortical regions providing
input to S1 but are strongly inhibited by VIP interneurons (Figure 4A, E)[58, 59]. VIP interneurons
may suppress SST activity during RWS, thus enabling plasticity[9, 60]. We have shown above that
optogenetic CF activation reduces whisker-evoked responses in a subset of VIP interneurons
(Figure 2G). To examine whether this activity regulation is critical for RWS+CF-induced
suppression of L2/3 pyramidal cell plasticity, we used chemogenetic approaches to manipulate
the activity of VIP interneurons. We first observed that expression of inhibitory hM4D (Gi)
DREADDs in VIP interneurons (Figure 4B) and acute DCZ administration significantly reduced
activity in VIP interneurons (Figure 4C) and prevented S1 potentiation upon RWS stimulation
alone (Figure 4D). This manipulation was similar to the effect of RWS+CF (Figure 1G-I) or SST
activation during RWS (Figure 3D). Conversely, expression of excitatory hM3D(Gq) DREADDs
in VIP interneurons (Figure 4F) and acute DCZ administration enhanced VIP interneuron
responses (Figure 4G) and rescued S1 plasticity upon paired RWS and optogenetic CF activation
(RWS+CF; Figure 4H), mimicking the effect of RWS (Figure 1G-I) or suppression of SST
interneuron activity during RWS+CF (Figure 3H).

To control for the continued presence of DCZ, identical experiments for each condition were
performed without the administration of RWS (Figure S5A, D) or RWS+CF (Figure S5G, J, M).
When comparing control and plasticity experiments, both the SST activation and VIP
inactivation experiments successfully show SST activation is sufficient to block RWS-mediated
PN potentiation (Figure S5C), and VIP activity is required for RWS-mediated PN potentiation

(Figure S5F). The VIP activation experiments also successfully showed VIP activation rescues
PN potentiation after RWS+CF (Figure S5O), and that this effect and was not due to sustained
increases in responsivity of pyramidal neurons caused by the presence of DCZ in these
conditions (Figure S5N). We additionally tested the potential contribution of PV interneurons to
the CF-mediated control of S1 plasticity. Expression of inhibitory hM4D (Gi) DREADDs in PV
interneurons indeed rescued PN potentiation after RWS+CF when compared to controls (Figure
S5L), a robust effect given that PV interneurons are usually basket cells with strong recurrent
connectivity to pyramidal neurons[56]. To support our interpretation that VIP interneurons control
a disinhibitory network in S1 cortex, we further analyzed the IN populations (which consist of
putative SSTs and PVs) in the VIP chemogenetic manipulation experiments (Figure 5). VIP
inactivation during RWS, which blocked PN potentiation (Figure 4D), indeed caused a
corresponding increase in IN activity (Figure 5C) to mimic the effects of RWS+CF (Fig. 1G-L)
even without CF activity. VIP activation during RWS+CF, which rescued PN potentiation
(Figure 4H), caused a corresponding decrease in IN activity (Figure 5F) to mimic the effects of
RWS (Figure 1G-L) even when CFs were active. These findings confirm that VIP interneurons
orchestrate the inhibitory network in S1 cortex and show that this mechanism is recruited by
optogenetic CF activation. Taken together, the findings presented thus far show that the critical
circuit for RWS-driven plasticity is located in S1 cortex itself and identify VIP and SST
interneurons as effectors that are sufficient to mediate the consequences of CF activation. The
notion of sufficiency does not exclude potential effects of plasticity processes elsewhere that
might well modulate effector activation in this context and others not tested here.

Zona incerta to the thalamic posterior medial nucleus (POm) as a multi-synaptic pathway
from the cerebellar nuclei to S1 cortex

Cerebellar climbing fibers must exert their effects on the neocortex via the deep cerebellar nuclei
(DCN), beyond which there may be several potential pathways to the primary somatosensory
cortex. The barrel cortex in mice receives whisker sensory information from two thalamic
sources, the ventral posterior medial thalamic nucleus (VPM), and the posterior medial thalamic
nucleus (POm). Whereas the VPM projects densely to the middle layers of S1 cortex, POm
inputs terminate largely in L5 as well as L1, where, in addition to pyramidal cells, they synapse
on VIP interneurons[58, 59]. POm-to-VIP inputs have previously been shown to gate RWS-
mediated plasticity of PNs in barrel cortex[50]. However, because the DCN send only weak direct
projections to POm (and VPM), we focused on other output regions of the cerebellum that could
influence the activity of sensory thalamic nuclei: the thalamic reticular nucleus (TRN) and zona
incerta (ZI). The ZI represents a more plausible pathway through which the CF could exert its
effects on S1 given its robust innervation by the cerebellar nuclei[17, 61, 62] and projections to the
POm[63]. To rule out a pathway to S1 through the thalamic reticular nucleus, whose GABAergic
neurons primarily inhibit VPM[64], we used retrograde fluorogold labeling from the somatosensory
thalamus (which included VPM and POm) and anterograde labeling from the contralateral DCN.
We found no overlap in the TRN (Figure S6). Next, we determined the primary output pathways
of ZI cells that receive cerebellar input by using a dual-injection approach which leveraged the
transsynaptic transport properties of AAV1[65]: AAV1-Flp recombinase (pAAV-EF1a-Flpo) was
delivered to the deep cerebellar nuclei, followed by a Flp-dependent label (Frt-ChR2-EYFP) to
contralateral ZI. Abundant EYFP-expressing neurons were found in the ZI (Figure 6A, B; Figure
S7), with their axon terminals visible in both POm (Figure 6A, B; Figure S7) and S1 cortex

(Figure 6A, B). These results suggest that the ZI-POm pathway, and not the TRN-VPM pathway,
constitutes a critical thalamic route for cerebellar signals reaching S1 cortex (Figure S6C).

To assess whether there is a monosynaptic connection from the cerebellar nuclei to neurons in
the ZI that project to POm, we induced ChR2 expression in the cerebellar nuclei and performed
in vitro whole-cell patch-clamp recordings from POm-projecting ZI neurons (i.e. cells that were
retrogradely labeled by fluoro-Ruby injected into POm) and tested them for ChR2-evoked
responses (Figure 6D, Figure S8). Optogenetic activation of ChR2+ terminals at 10Hz elicited
EPSPs that persisted in the presence of bath-applied TTX (1 µM; Figure 6E, F), despite
significant amplitude reduction. This TTX-insensitive component represents the ChR2-evoked
EPSP at the synapses between cerebellar nuclei neurons and ZI neurons and demonstrates the
monosynaptic nature of this pathway[66].

It has previously been demonstrated that most POm-projecting neurons in ZI are PV neurons[67].
To test whether these neurons convey cerebellar instructive signals to S1 cortex, we
chemogenetically inhibited PV neurons in ZI during basic whisker and whisker plus optogenetic
CF activation conditions and measured responses in L2/3 pyramidal neurons in S1 cortex.
Expression of inhibitory hM4D (Gi) DREADDs in ZI-residing PV neurons (Figure 6G) and DCZ
application prevented the suppressive effect of CF co-activation on pyramidal cell responses
(Figure 6H, I). These data demonstrate that thalamically projecting PV neurons in ZI convey CF
signals to S1 cortex.

Discussion

Multiple functions have been assigned to cerebellar CFs. As in our recordings, optogenetic CF
co-activation suppresses S1 pyramidal neuron plasticity in an experimental paradigm lacking
behavioral reward, this behaviorally neutral signal acts like an error signal would. However, it
remains possible that CF recruitment and regulation of S1 plasticity may happen in a reward /
reward omission context as well. CF activation might be driven by reward / reward omission in
sensory-association training. This might be the reason why sensory stimulation in such training
context drives cortical plasticity less well than passive sensory stimulation alone[68]. Plasticity in
S1 cortex results from an update in sensory input – here exposure to a new sensory experience,
prolonged rhythmic whisker activation at 8Hz – that necessitates circuit adaptation. CF signaling
may get recruited when an obvious disruption or violation of a sensory prediction occurs, or
simply when no stability in sensory input is reached. The absence of *sameness* may be detected
by inferior olive neurons as an error in the sensory environment, and subsequently CFs prevent
the otherwise unfolding potentiation. Such function provides an example for a "sensory role" of
the olivo-cerebellum[69,70], in this case by controlling adaptive plasticity in a primary sensory
cortex, without immediate effects on the motor domain.

In rodents, direct trigemino-olivary connections convey whisker-related information to the
olive[71]. Passive whisker activation evokes CF-mediated complex spike firing in PCs in ipsilateral
crus I/II[40]. These evoked spikes occur on top of a spontaneous complex spike firing rate of
~1Hz[39]. What distinguishes CF activity that suppresses plasticity in the cerebral cortex from
these more common activity patterns? First, prolonged low-frequency air puff stimulation to
sensitive skin areas on the snout or wrist of rodents evokes potentiation, and not depression, of
PC responses[72, 73]. This finding is in line with the current understanding of CF signaling: novelty
activates it[39]; when stimuli are repeatedly presented (producing/inhabiting a state of *sameness*),

evoked complex spike firing weakens. Thus, RWS stimulation likely will not produce continuous
elevated CF activation. For this to happen under natural conditions, novel input features have to
emerge. Second, the synchronization of CF responses might be critical. Complex spike
synchrony is absent or weak during spontaneous activity and might increase with movement
initiation[74, 75] or sensory stimulation[76]. Synchrony in a sufficiently high number of PCs is needed
to evoke inhibition and subsequent rebound excitation in target neurons in the cerebellar nuclei[77,
78]. Parallel fibers (PFs) likely cannot generate synchronous activity in enough PCs due to the
spatial arrangement of PF input[79]. A threshold level of synchronization will be reached with
optogenetic CF activation but may also naturally be reached with specific qualities of the sensory
input signal, e.g. those that make it an error signal. It is also conceivable that synchronization to
a threshold level is driven by an error signal that does not originate from whisker stimulation but
has another sensory or non-sensory origin.

Whisker-related sensory information in rodents is conveyed by the trigeminal ganglion via
thalamic nuclei VPM and POm to the barrel cortex; information from specific whiskers is
projected via VPM to cortical layer 4[80]. The observation that cerebellar signals impact S1 cortex
via ZI and POm fits the description of a more modulatory pathway that does not interfere with
thalamic input in layer 4, but with synaptic signal integration in the more superficial as well as
deeper layers. Both two-photon imaging and chemogenetics identify SST interneurons in S1
cortex as the ultimate effectors of the activation of this modulatory pathway (note that a similar
mechanism might underlie a phenomenon clinically known as CBI or 'cerebellar-brain-
inhibition', where electrical or magnetic stimulation over the cerebellum in human subjects
causes a reduction in excitability of the motor cortex[81, 82]). SST interneurons, in turn, are
regulated by VIP interneurons that receive activation from POm[59]. We identified PV neurons in
ZI as neurons that are critically involved in this pathway as well, likely reducing the POm-
mediated activation of VIP interneurons in S1 cortex. Our findings demonstrate that cerebellar
output may recruit PV neurons in ZI sufficiently well to ultimately cause a removal of VIP
interneuron-mediated inhibition of SST interneurons and consequently enhance dendritic
inhibition of L2/3 pyramidal neurons. This activation sequence likely requires an initial
activation of excitatory neurons in the cerebellar nuclei by output from the cerebellar cortex,
which may result from rebound excitation following synchronous PC activity[78]. Another context
where PV neurons in the ZI become activated – via glutamatergic input from S1 cortex – is self-
grooming in the orofacial area[83]. It is plausible that under such conditions of predicted sensory
input, whisker-related L2/3 pyramidal neuron plasticity is suppressed as well. While our findings
identify the cerebellar nuclei to ZI to POm projection as critical to conveying CF signals to S1
cortex, it is likely that more projections that are not tested here can provide links from the
cerebellum to S1 cortex. This is suggested by the distinct multi-channel output from the
cerebellar nuclei[84]. For example, we cannot exclude a participation of direct projections from the
cerebellar nuclei to VPM or POm[85] (note, however, the prominence of projections that reach the
ZI and originate in the cerebellar nuclei, as shown in Figure 6A-C; Figure S7). Moreover, it
remains possible that cerebellar output reaches S1 cortex via M1 cortex, as both cortices show
strong reciprocal connections[86] and S1-projecting pyramidal neurons in M1 target VIP
interneurons in S1 cortex[87].

Our data show that activity of cerebellar CFs has an impact on S1 cortex, preventing plasticity of
L2/3 pyramidal neurons via the recruitment of local SST interneurons. In our experiments, CFs
were artificially (optogenetically) stimulated, and definitive conclusions about CF engagement
and its consequences under natural conditions remain open. However, our study demonstrates

both the capacity for such impact on cortical signaling and a pathway that enables this impact.
These are requirements for CF signals to act as instructive signals in supervised learning[88]. A
condition for such an effect likely is a synchronization of CF responses in groups of PCs[77, 78], a
scenario that provides a qualifying distinction from other CF activity patterns[74-76]. The observed
external influence enables regulatory control that might safeguard the cerebral cortex from
maladaptation, albeit at the cost of exposing it to pathological dysfunction of the olivo-cerebellar
system, e.g. sensory over-responsivity that has been described in syndromic autism[48].

Materials and Methods:

Animals

All animal experiments were approved and conducted in accordance with the regulations and
guidelines of the Institutional Animal Care and Use Committee of the University of Chicago
(IACUC 72496). Mice were housed on a 12-hour light/dark cycle and fed a standard rodent diet.
Animals of either sex were used in all experiments and no sex-dependent differences were
observed in any reported measures. Strains included wildtype mice (C57BL/6J; Jax: 000664),
PVCre mice (B6.129P2-Pvalbtm1(cre)Arbr/J; Jax: 017320), SSTCre mice (STOCK
Ssttm2.1(cre)Zjh/J; Jax: 013044), VIPCre mice (B6J.Cg-Viptm1(cre)Zjh/AreckJ; Jax: 031628),
and Ai9 mice (B6.Cg-Gt(ROSA)26Sortm9(CAG-tdTomato)Hze/J; Jax: 007909). Homozygous
SSTCre, PVCre, and VIPCre animals were bred with wild type animals to generate
heterozygotes. To generate SST and VIP reporter lines (SSTCre^{tdTom} and VIPCre^{tdTom}),
homozygous SSTCre and VIPCre mice were bred with homozygous Ai9 mice. Only
heterozygous animals were used for experiments given evidence that endogenous somatostatin
(SST) and vasoactive intestinal polypeptide (VIP) expression may be reduced in homozygous
SSTCre and VIPCre animals, respectively[89, 90].

Stereotaxic injections

For electrophysiological experiments. Stereotaxic virus injections for electrophysiological
experiments were performed as previously described using C57BL/6J mice between postnatal
day 23-28[91]. For optogenetics experiments in which cerebellar inputs to zona incerta were
measured, a 250 nL (40 nL/min) infusion of AAV5-hSyn-hChR2(H134R)-EYFP (UNC Vector
Core) was made into the right deep cerebellar nuclei (-1.5 ML; -5.8 AP; -3.2 DV[92]) using a 1 mL
Hamilton syringe (catalog: 65458-02) with a flat needle tip. Next, a small infusion (40 nL total,
at 7nL/min) of fluoro-Ruby (ThermoFisher, catalog: D1817) was made into left posterior medial
thalamus using a 0.5 mL Hamilton syringe (catalog: 65457-01). Following a two-week
incubation, animals were sacrificed for recordings in acute slices (see *Slice electrophysiology*).

For tracing experiments. For anterograde transsynaptic labeling experiments, a dual-injection
approach was utilized to specifically label the outputs of neurons in the zona incerta that receive
input from the contralateral cerebellar nuclei. Flpo-recombinase (AAV1-EF1a-Flpo, item#:
55637, Addgene) was injected into left deep cerebellar nuclei as above (+1.5 ML; -5.8 AP; -3.2
DV), followed by an infusion of AAV1-Flp-on ChR2-EYFP (AAV-hSyn-Coff/Fon-
hChR2(H134R)-EYFP-WPRE, UNC Vector core) into right zona incerta (-1.8 ML; -2.0 AP; -4.0
DV). Following a 5-week incubation – the period required for anterograde transport of the

AAV1 virus injected into zona incerta – animals were transcardially perfused for histology (see *Tissue preparation for fluorescence microscopy and immunohistochemistry*).

To visualize potential overlap between cerebellar terminals and cells in thalamic nuclei (i.e. the thalamic reticular nucleus) that project to somatosensory thalamus (ventral posteromedial nucleus and posterior medial thalamus), an orthograde virus (AAV5-hSyn-hChR2(H134R)-EYFP (UNC Vector core)) was injected into the right deep cerebellar nuclei (-1.5 ML; -5.8 AP; -3.2 DV) and fluorogold (2% w/v in sterile saline, Fluorochrome) was injected into the left somatosensory thalamus (primarily posterior medial thalamus). Following a 2-week incubation, animals were transcardially perfused for histology (see *Tissue preparation for fluorescence microscopy and immunohistochemistry*).

For two-photon calcium imaging experiments. Viral injections were performed to express channelrhodopsin-2 in cerebellar climbing fibers; fluorescent calcium indicators in cerebellar Purkinje cells or primary somatosensory (S1; barrel) cortex neurons; tdTomato, activating DREADDs, or inactivating DREADDs in S1 cortex; and inactivating DREADDs in interneurons in zona incerta. With respect to the inferior olive injection site (conferring viral expression in climbing fibers terminating in the contralateral cerebellar hemisphere), injections into cerebellar cortex were performed contralaterally, and injections into zona incerta and somatosensory cortex were performed ipsilaterally.

To express channelrhodopsin-2 in cerebellar climbing fibers, ChR2-EYFP (AAV9-CaMKIIa-hChR2(H134R)-EYFP; Addgene #26969-AAV9) or ChR2-mCherry (pAAV9-CaMKIIa-hChR2(H134R)-mCherry; Addgene #26975-AAV9) was injected into the left inferior olive as follows: after surgical preparation (see *Two-photon calcium imaging: Cranial window surgeries*), tissues attached to the occipital bone were surgically detached, followed by opening of the posterior atlanto-occipital membrane to expose the dorsal medulla. A 5μL Hamilton Neuros syringe (catalog: 65460-03) with a 33-gauge beveled needle tip (catalog: 65461-02) was inserted into the left medulla at an angle of 58-59° to the perpendicular line (Supplementary Data 2a), 0.75 mm laterally and 1.70 mm deep from the surface of the medulla as previously described[93]. A total volume of 400 nL was infused at a rate of 80 nL/min, and the syringe was held in place for five minutes before removal. ChR2-EYFP or ChR2-mCherry was chosen based on the excitation and emission spectra of other fluorophores expressed in the same animal to maximize signal separation when conducting post-hoc confocal imaging.

Stereotaxic virus injections for imaging of Purkinje cell dendrites were administered as previously described using P80-120 adult C57BL/6J mice[73]. A virus mixture was prepared with 0.5% L7-Cre (AAV1.sL7.Cre.HA.WPRE.hGH.pA, Princeton Neuroscience Institute Viral Core Facility, kindly provided by the lab of Dr. Samuel Wang at Princeton University), and 20% Cre-dependent GCaMP6f (AAV1.CAG.Flex.GCaMP6f.WPRE.SV40; Addgene, #100835) in saline solution. A glass pipette with a tip diameter of ~300 μm was prepared by a puller (model P-97, Sutter Instrument Co.) and used to inject a total volume of 1800 nL into two separate injection sites: 900 nL were injected into right medial Crus I (-1.8 ML; -7 AP; -1.8 DV), followed by a 900 nL injection into right medial Crus II (-2.5 ML; -7 AP; -2.2 DV). The pipette carrying the AAV virus was held in place for five minutes before withdrawal.

For DREADD suppression of PV interneurons in zona incerta, a 1 mL Hamilton syringe was used to infuse Cre-dependent hm4D(Gi) (AAV5-hSyn-DIO-hM4D(Gi)-mCherry; Addgene #44362-AAV5) into the left zona incerta (1.8 ML; -2.0 AP; -4.0 DV) of PVCre animals.

To express GCaMP6f in primary somatosensory (barrel) cortex neurons in C57BL/6J,
SSTCre^tdTom, and VIPCre^tdTom animals, a 5μL Hamilton Neuros syringe with a 33-gauge beveled
needle tip was used to infuse a total volume of 800 nL of GCaMP6f
(AAV1.Syn.GCaMP6f.WPRE.SV40; Addgene #100837-AAV) at two separate injection sites:
400 nL were injected at -1.2 AP; 2.5 ML and 400 nL at -1.5 AP; 3 ML. At each injection site,
200 nL were first injected 300 μm below the pial surface, followed by 200 nL at 150 μm below
the pial surface. The injections were performed at a rate of 100nL/minute, and the syringe was
held in place for five minutes before removal. To express GCaMP6f in barrel cortex neurons
with co-expression of tdTomato in parvalbumin (PV)-, SST-, and VIP-expressing interneurons in
PVCre, SSTCre, or VIPCre animals, respectively, GCaMP6f (Addgene #100837-AAV1) and
Cre-dependent tdTomato (AAV9-FLEX-tdTomato; Addgene #28306-AAV9) were co-injected
using the same technique. To optimize co-expression of the viruses, GCaMP6f with tdTomato
was mixed at a 3:1 volume ratio. For DREADD activation or suppression of VIP, SST, or PV
interneurons, Cre-dependent hM3D(Gq) (AAV5-hSyn-DIO-hM3D(Gq)-mCherry; Addgene
#44361-AAV5) or Cre-dependent hM4D(Gi) (AAV5-hSyn-DIO-hM4D(Gi)-mCherry; Addgene
#44362-AAV5) were used in place of tdTomato, respectively.

For all two-photon imaging experiments, recordings began after a 4-week incubation period after
the final virus injection (see *Two-photon calcium imaging: cranial window surgeries for detail*).

Tissue preparation for fluorescence microscopy and immunohistochemistry

After completion of two-photon experiments, selected animals (all those used in DREADD and
cerebellar imaging experiments, and a subset of animals in all other experimental cohorts) were
sacrificed for post-hoc characterization of viral expression. Animals were transcardially perfused
with ice-cold 4% paraformaldehyde in phosphate-buffered saline (25mL), pH 7.4. The brain was
extracted and postfixed in 4% paraformaldehyde overnight at 4°C before being transferred to a
cold 30% sucrose (in phosphate-buffered saline) solution for >48 hours. Brains were then
cryosectioned at 50 μm thickness, with an approximately equal number of brains sectioned
coronally or sagittally per experimental cohort. For tracing experiments, animals were perfused
with ice-cold phosphate-buffered saline (25mL) prior to perfusion with paraformaldehyde and
cryosectioned coronally at 150 μm thickness. Slices were stored in phosphate-buffered saline
prior to immunohistochemistry or confocal imaging.

For post-hoc visualization of Purkinje cells in mice expressing ChR2-EYFP in climbing fibers
(i.e. in animals without GCaMP6f expression in Purkinje cells), calbindin immunohistochemical
staining was performed. Tissue slices were washed in a 0.01M phosphate-buffered saline
solution containing 200 mM glycine for two hours at 4°C, then incubated in a 0.01M phosphate-
buffered saline solution containing 10 mM sodium citrate and 0.1% Tween-20 at ~60°C for 30
minutes using a heated water bath. After cooling to room temperature, the slices were washed
three times in a 0.01M phosphate-buffered saline solution containing 0.5% Tween-20 (PBS-
Tween) at room temperature for a period of 15 minutes per wash. Tissue was permeabilized with
200 mM glycine in PBS-Tween for 15 minutes at room temperature. Blocking was done for two
hours at room temperature with a PBS-Tween solution containing 5% bovine serum albumin and
10% normal donkey serum, followed by overnight incubation (12-15 hours) at 4°C in a PBS-
Tween solution containing 1% normal donkey serum and guinea pig anti-calbindin primary
antibody (1:500; Synaptic Systems Cat# 214 004, RRID:AB_10550535). After washing the

tissue three times for fifteen minutes in PBS-Tween at 4°C, slices were incubated for two hours
at 4°C in PBS-Tween containing 1% normal donkey serum and donkey anti-guinea pig Cy3
(555) secondary antibody (1:200; Jackson ImmunoResearch Labs Cat# 706-165-148,
RRID:AB_2340460). Finally, the slices were washed three times in PBS-Tween (ten minutes per
wash) before mounting with Vectashield (Vector Laboratories, Inc.). The mounted slices were
allowed to set overnight before imaging.

To confirm the channelrhodopsin-2 expression observed in cerebellar cortex was indeed
expressed in climbing fiber terminals, a subset of tissue was additionally labeled with VGluT2.
VGluT2 (rabbit anti-VGluT2, 1:500; Thermo Fisher Scientific Cat# 42-7800,
RRID:AB_2533537) was added to the primary antibody solution and donkey anti-rabbit AF647
(Jackson ImmunoResearch Labs Cat# 711-605-152, RRID:AB_2492288) was added to the
secondary antibody solution.

Confocal imaging

Image acquisition. Z-stack confocal images were taken at 5x (Olympus MPlan N 0.1NA, air),
10x (Zeiss Achroplan 0.25NA, air), 20x (Olympus UMPlanFL N 0.5NA, water), 40x (Zeiss EC
Plan-Neofluar 1.3NA, oil immersion), and 63x (Zeiss W Plan-Apochromat, water)
magnifications with a Zeiss LSM 900 Axio Examiner.Z1 scope. A subset of images (Figure 1e
and Figure S1B) were taken with a Zeiss LSM 5 Exciter, Axioskop 2 (using the 20x Olympus
UMPlanFL and 40x Zeiss EC Plan-Neofluar objectives, respectively). The images in Figure 6d
(shown with channel separation in Figure S8b) and Figure S6 were taken with a second Zeiss
LSM 900 confocal microscope. To facilitate direct comparison, images shown in series (as in
Figure 6c and Figure S7f) were taken with identical hardware and software configurations,
available in the image metadata.

Post-hoc image processing. To stitch multiple images (as in Figures 1B, 1D, 2D-F (right), 4B,
4F, 6B, 6C, S1A (top), S1E, S6A, S6B (top and bottom left), and S7C-E), a series of pairwise
stiches were performed using Fiji (NIH). Figure 1B and Figure S1A (top) each consisted of 4
images (each 5x magnification, 1024 x 1024 pixels). Figure 1D consisted of 12 images (each 5x
magnification, 512 x 512 pixels). Figure 2D (right) consisted of 10 images (each 10x
magnification, 1024 x 1024 pixels). Figures 2E and 2F (right) each consisted of 8 images (each
10x magnification, 512 x 512 pixels). Figures 4B and 4F each consisted of 2 images (each 20x
magnification, 512 x 512 pixels). Figure 6B (also shown as the first image in Figure S7D), and
all images in S7D and S7E each consisted of 6 images (each 5x magnification, 512 x 512 pixels).
Images in Figure 6C each consisted of 9 images (each 10x magnification, 512 x 512 pixels).
Figure S1E consisted of 8 images (each 5x magnification, 1024 x 1024 pixels). S7C consisted of
10 images (each 40x magnification, 512 x 512 pixels). Note a subset of the final stitched images
displayed in figures are cropped to show relevant detail.

Channel colors set during image acquisition followed conventional emission wavelength
assignments (i.e. red to indicate tdTomato, green to indicate GCaMP6f, etc.). To illustrate
interneuron subtypes and continuity across experiments (e.g., cyan to indicate ChR2 expression
across all optogenetic stimulation experiments), channel colors were specified post-imaging
using Fiji by separating channels and selecting the appropriate color when merging, ignoring
source LUTs. When necessary, post-hoc brightness and contrast adjustments were applied
linearly to an entire image – *after* stitching, if applicable – using Fiji. To facilitate direct

comparison, identical linear brightness/contrast adjustments were made across all images shown
in series (as in Figure 6C, Figure S7D, and Figure S7E).

To measure terminal density across the sections shown in Figure 6c, Fiji was used to rotate each
image such that the axis separating ventral from dorsal zona incerta was horizontal. A 400 μm x
1 mm region of interest (ROI) was drawn around the zona incerta, and each row of pixels within
this ROI was separated to facilitate calculation of the integrated density across each row of
pixels. A similar method was used to quantify the terminal density of cerebellar inputs to zona
incerta as in Figure S8. Briefly, images of sections matched for their rostro-caudal location were
rotated as before. Then, five line-ROIs oriented vertically, each being 200 μm wide, were
positioned side-by-side across the mediolateral axis of the zona incerta, and the spatial pattern of
fluorescence along these lines was measured by calculating the mean grey value along their
length (which was roughly 400 μm from the dorsal boundary to the ventral boundary). These
were averaged for each animal used. Shading represents SEM for both measurements.

Slice electrophysiology

Acute slice preparation and whole cell recordings. For recordings in acute slice, injected
animals (see *Stereotaxic injections: For electrophysiological experiments*) were deeply
anesthetized using the isoflurane drop method in a bell jar (with a raised platform for the animal)
and immediately transcardially perfused with 8-10 mL of ice cold oxygenated (95% O_2, 5%CO_2)
artificial cerebrospinal fluid, which contained the following (in mM): 125 NaCl, 25 $NaHCO_3$, 3
KCl, 1.25 NaH_2PO_4, 1 $MgCl_2$, 2 $CaCl_2$, and 25 glucose. The brain was then extracted, glue-
mounted on a vibratome (Leica) platform and blocked for coronal slices using an agarose (5%)
cube and sliced in the same solution (ice-cold). Slices were cut to 365 μm thickness. The brain
slices were then transferred to 32-34 °C oxygenated artificial cerebrospinal fluid that was
allowed to return to room temperature over the course of one hour, which constituted the slice
recovery period.

Slices containing the zona incerta and terminals from the cerebellar nuclei inputs were visualized
using differential interference contrast with an Axioskop 2FS microscope (Carl Zeiss).
Fluorescence from ChR2-EYFP expression and from retrograde fluoro-Ruby labeling was
confirmed using the 5x air objective and guided recording locations in zona incerta. Recordings
were made with a Multiclamp 700B amplifier and pCLAMP software (Molecular Devices).
Recording glass pipettes with 7–9 MΩ resistance were filled with an internal solution as follows
(in mM): 117 K-gluconate, 13 KCl, 1 $MgCl_2$, 0.07 $CaCl_2$, 10 Hepes, 0.1 EGTA, 2 Na_2-ATP, 0.4
Na-GTP, pH 7.3, 290 mOsm. Incertal cells whose morphology was clearly revealed by strong
red fluorescence, indicating retrograde labeling from posterior medial thalamus, were targeted
for recordings. Bath application of TTX (1 μM) was performed in a subset of recorded cells to
verify the monosynaptic nature of the cerebellar input to zona incerta.

Optogenetic stimulation was delivered using a 355 nm laser (DPSS: 3505–100), controlled with
galvanometer mirrors (Cambridge Technology) focused on the slice through a 5x air objective
using custom software in MATLAB (MathWorks). Focal photostimulation of the ChR2-
expressing synaptic terminals in zona incerta was performed at 10Hz (4 pulses of 1-ms duration
at 100 ms interstimulus interval).

Electrophysiological data analysis. Electrophysiological data were collected using custom MATLAB software and analyzed using Graphpad Prism (v7.0). The amplitude of excitatory responses (i.e., the first response in the stimulus train) to stimulation pulses was measured by subtracting the average value for 20 ms before the delivery of a pulse (baseline) from the maximum value of the peak in current clamp at resting membrane potential. Latency of the optogenetic response was calculated by measuring the delay in milliseconds from the onset of the first optogenetic stimulus to the moment the resulting response reached >10% of its maximum amplitude.

Two-photon calcium imaging

Cranial window surgeries for imaging of Purkinje cell dendrites. To minimize overall recovery burden, surgeries were performed in two stages: (1) inferior olive virus injection and headframe installation, and (2) cerebellar virus injection and cerebellar cranial window installation as previously described[73]. In the first stage, mice (P60-90) were deeply anesthetized using 1-2% isoflurane and clamped by ear bars at the external acoustic foramen. The line between the clamping point and maxilla was set parallel to the horizontal plane. After trimming the fur on top of the skull with clippers, the surgical site was prepared by applying betadine and 70% ethanol three times in an alternating fashion. Meloxicam (2 mg/kg), extended-release buprenorphine (0.1 mg/kg), and 0.5 mL saline were administered subcutaneously, and depth of anesthesia was confirmed via tactile stimulation of the toe before making a 20-25 mm incision in the skin to reveal the occipital bone and attached tissues. After opening of the posterior atlanto-occipital membrane and channelrhodopsin-2 virus injection into the left inferior olive (see *Stereotaxic injections: For two-photon calcium imaging experiments*), tissues were reattached to the occipital bone using instant adhesive, ensuring adequate space for later implantation of the cranial window over the right cerebellar cortex. A custom titanium headframe (H. E. Parmer Company, Nashville, TN, USA) was secured with dental cement (Stoelting Co., Wood Dale, IL, USA), completing the first stage.

After monitoring recovery for a period of 4 days, mice underwent the second surgery. A circular craniotomy over right cerebellar cortex with a diameter of 4mm, centered at -2.7ML; -6.9 AP, was performed using a dental drill. The dura was carefully removed to expose lobules Crus I and anterior Crus II before administering GCaMP6f injections (see *Stereotaxic injections: For two-photon calcium imaging experiments*). Following injection, a two-layer glass window was installed using C&B Metabond dental cement (Patterson Dental Company, Saint Paul, MN, USA), completing the second stage. The glass windows consisted of a 4 mm glass window (Tower Optical Corp, Boynton Beach, FL, # 4540-0495) adhered to a 5 mm glass window (Warner Instruments, Holliston, MA, USA, # CS-5R) using ultraviolet light-activated glue (Norland Optical Adhesive 71; Norland Products Inc., Jamesburg, NJ). Experiments were conducted 4.5 weeks after inferior olive injection and 4 weeks after GCaMP6f injection.

Cranial window surgeries for imaging of barrel cortex neurons. Surgeries were again separated into two stages: (1) inferior olive virus injection, headframe installation, and cerebellar cranial window installation; and (2) S1 cranial window installation and virus injection. First, channelrhodopsin-2 was injected into the left inferior olive as above (see *Cranial window surgeries for imaging of Purkinje cell dendrites* and *Stereotaxic injections: For two-photon calcium imaging*). After reattaching tissues to the occipital bone using instant adhesive, a dental

drill was used to perform a circular craniotomy over right cerebellar cortex (lobules Crus I and
II), followed by removal of the dura and installation of the glass window as above. Finally, the
custom titanium headframe was secured using dental cement, completing the first stage.

After monitoring recovery for a period of 4 days, mice underwent the second surgery. After
surgically detaching a small portion of the left temporalis muscle attached to the parietal bone, a
second circular craniotomy over left primary somatosensory cortex with a diameter of 4 mm
(centered at -1.5 AP; 3 ML) was performed. The dura was removed to expose the neocortical
surface before administering the appropriate viral injections (see *Stereotaxic injections: For two-*
photon calcium imaging experiments). Following injection, a glass window was installed over
the craniotomy using dental cement. To ensure optical isolation during recordings, a custom
sheath was used to couple a Thorlabs patch cable (used for LED stimulation of cerebellar
climbing fibers; see *Optogenetic and tactile stimulation*) to the cerebellar window. This sheath
was adhered over the cerebellar window using Metabond dental cement, completing the second
stage. For experiments involving DREADD manipulation of neurons in zona incerta, viral
injections into left zona incerta were administered three days prior to the first surgery. Thus,
experiments were conducted approximately 5 weeks after zona incerta injection, 4.5 weeks after
inferior olive injection, and 4 weeks after primary somatosensory cortex injections.

Habituation. After completion of surgeries, postoperative care was provided for 5-7 days. Once
animals exhibited exploratory behavior without signs of pain or distress, habituation began with
handling of mice for 15 minutes and transportation from the animal facility to the laboratory.
After two days, anxiety during handling typically diminished, and the handling period increased
to 30 minutes in the vicinity of the recording area. Following two days of handling, the mice
were introduced to a free-running treadmill and head restraint via clamping of the head frame on
left and right sides. The duration of time spent on the treadmill increased from 10 minutes to two
hours per day depending on the mouse's comfort. The relative time spent running on the
treadmill typically decreased after two days. To habituate mice to air puff stimulation during this
time, receptive field mapping of the field of view was conducted by delivering 8psi air puffs
through a capillary tube while observing epifluorescence signals and calcium responses in real
time. Air puffs were delivered at irregular frequencies, with at least 30 seconds between stimuli,
to avoid influencing subsequent plasticity experiments. In total, mice were habituated for a
period of 1-2 weeks until they showed no aversive responses to air puff stimulation, spent the
majority of time in the recording environment at rest, and reduced movement initiation
(whisking/running) with air puff stimulation. Animals used for DREADD manipulation
experiments were familiarized with the DCZ injection procedure by receiving DMSO/saline
injections during habituation for two days prior to recording.

Optogenetic and tactile stimulation. To optogenetically stimulate cerebellar climbing fibers,
the tip of a Ø200 μm core rotary joint patch cable (Thorlabs catalog: RJPSL2) was inserted into
the sheath attached to the cranial window over the right cerebellar cortex (see *Two-photon*
calcium imaging: Cranial window surgeries). For experiments in which Purkinje cell responses
to optogenetic climbing fiber stimulation were also measured, the patch cable was inserted into a
custom light shield that fit over both the objective and the cable. The tip of the patch cable was
centered over Crus II in all optogenetic stimulation experiments. Blue light pulses were delivered
with a 470 nm Fiber-Coupled LED (Thorlabs catalog: M470F3). To deliver whisker stimuli, a
0.86 mm diameter glass capillary tube was positioned over the right C-row whiskers 5 mm from
the right whisker pad at an angle of 45° in the X and Y planes (adjusted slightly during receptive

field mapping for each animal). Air puffs were delivered at 8 psi (Picospritzer III, Parker
Hannifin). For all imaging sessions, calcium activity was measured in trials lasting 20 seconds,
with a stimulus delivered once per trial. Trials began at random intervals, with an interval of at
least 5 seconds between trials. When testing Purkinje cell responses to optogenetic climbing fiber
stimulation, 470 nm pulses were delivered with the following durations: 1 x 20ms, 1 x 50ms, 3 x
15 ms at 8 Hz, and (3 x 15 ms at 8 Hz)*2 with a 250 ms interval between epochs. The 1 x 50 ms
light pulse was selected for all subsequent experiments given this stimulation duration evoked
responses resembling spontaneous events in the same Purkinje cells (Figure S1). To test the basic
transmission of the climbing fiber signal to S1 (i.e., the immediate influence of optogenetic
stimulation of climbing fibers on whisker responses shown in Figure 2), 100 ms air puff test
pulses were delivered with or without optogenetic co-stimulation of climbing fibers. For paired
test pulse stimuli, the 50 ms blue light stimulus was delivered with a delay of 45 ms with respect
to the onset of the whisker stimulus. This delay was chosen to mimic the natural latency of
climbing fiber responses to whisker stimulation, which peaks approximately 50 ms after stimulus
onset[40]. For plasticity experiments, whisker test pulses were delivered during a baseline period of
at least 30 minutes, followed by a 5-minute period of rhythmic whisker stimulation (RWS) at 8
Hz – the speed at which mice naturally sample objects[51]. For plasticity experiments with
optogenetic co-activation of climbing fibers (RWS+CF), the same protocol was followed with
the addition of 50 ms blue light pulses at 1 Hz during plasticity induction. The 8 Hz whisker
stimulation and 1 Hz CF stimulation occurred at a consistent time relative to one another, with a
blue light pulse always occurring at a 45 ms delay relative to the onset of a whisker stimulus (the
same delay used for the paired test pulses). After RWS or RWS+CF, whisker test pulses were
delivered for a period of at least 40 minutes. To coordinate precise stimulus delivery, a Cygnus
Digital Stimulator (catalog: PG4000A, Cygnus Technology, Inc.) was interfaced with an
Arduino Uno to trigger both the LED driver and picospritzer.

Cerebellar imaging protocol. Calcium imaging of the genetically encoded indicator GCaMP6f
was conducted in Crus I/II of the right hemisphere of awake, head-fixed mice using a laser
scanning two-photon microscope (Neurolabware, Los Angeles, CA, USA) and Scanbox software
(Scanbox, Los Angeles, CA). Calcium images were obtained at a frame rate of 30.98 fps with a
pixel dimension of 512 x 796, using an 8KHz resonant scanning mirror with bidirectional
scanning. A Mai Tai® DeepSee (Spectra-Physics, Milpitas, CA, USA) laser source was used to
excite GCaMP6f at 920 nm. The fluorescence emission was collected through a 16x water-
immersion objective (Nikon LWD 0.8NA, 3 mm WD) using a GaAsP PMT (Hamamatsu
Photonics, Shizuoka, Japan). A 2.0 or 2.8x digital zoom was applied during imaging, generating
a field of view of 760 x 613 or 551 x 444 μm, respectively. To minimize background noise
originating from ambient light, a custom light shield was fitted around the brain window and
objective. This shield had a small opening such that the Thorlabs patch cable used for LED
stimulation of climbing fibers could be positioned over the cerebellar window. The laser power
was set to 1% with a PMT gain between 0.79 and 0.81, allowing for prolonged recording while
minimizing phototoxicity.

S1 imaging protocol. Calcium imaging of GCaMP6f was conducted in barrel cortex of the left
hemisphere using the same hardware as above. Images were obtained at a frame rate of 30.98 Hz
with a pixel dimension of 512 x 796. A 2.4x digital zoom was applied during imaging,
generating a field of view of 638 x 515 μm. Two of 11 RWS+CF experiments and one of 10
RWS experiments were conducted using a 2.0x digital zoom (generating a field of view of 760 x
613 μm) and were added to the experimental cohorts, as results were not significantly different

across any reported measures. A custom light shield was fitted around the brain window and
objective to reduce background noise originating from ambient light. Note further isolation of the
LED light used for optogenetic stimulation of climbing fibers in these recordings was provided
by the sheath attached to the cerebellar window (see *Cranial window surgeries for imaging of*
barrel cortex neurons). While observing the epifluorescence signal, small vessels on the putative
surface of the brain were put into focus in the z plane and matched to a field of view selected
during receptive field mapping (see *Habituation*). After adjusting the focus 250 μm below the
putative pial surface, small adjustments were made in the x-y planes while observing calcium
signal in real time to maximize responsivity to whisker stimulation and overlap between
GCaMP6f and tdTomato or mCherry in the field of view. GCaMP6f was excited at 920 nm and
tdTomato or mCherry (for DREADD experiments) at 1040 nm, with emitted fluorescence
collected using two GaAsP PMTs. The laser power was set to 2%, and adjustments to the PMT
gain were made due to slight variance in the level of GCaMP6f and tdTomato/mCherry signal
across animals. PMT gain in the green channel was 0.73 (SD: 0.07) while PMT gain in the red
channel was 0.83 (SD: 0.14), which were not significantly different across experimental groups.
Responsivity to whisker stimulation was determined by intermittently delivering 8 psi air puffs at
random intervals, with a minimum of 30 seconds between stimuli, to avoid influencing plasticity
experiments. If more than ten minutes were spent determining an optimal field of view on the
day of recording, the experiment was aborted and instead conducted the following day. Mice
were occasionally used for multiple recording configurations but were sacrificed immediately
after performing plasticity experiments, and selected animals were used for histological
characterization (see *Tissue preparation for fluorescence microscopy and*
immunohistochemistry). Note DREADD expression was histologically verified in all animals to
ensure expression was restricted to neocortex.

DREADD experimental protocol. To activate the hM4D(Gi) and hM3D(Gq) receptors, we
used deschloroclozapine dihydrochloride (DCZ, MedChemExpress), which was chosen over
clozapine N-oxide (CNO) as an agonist due to its heightened selectivity for hM4D(Gi) and
hM3D(Gq) receptors over endogenous receptors, a more rapid onset than CNO, and significantly
increased potency. DCZ was dissolved in DMSO at a 0.02 mg/mL concentration and stored at
−80 °C. On the day of recording, DCZ aliquots were thawed to room temperature and diluted to
0.01 mg/mL with DMSO/saline. Two-photon calcium imaging experiments began with
recordings of baseline responses to whisker stimulation (with and without optogenetic
stimulation of climbing fibers in a subset of experiments), followed by an injection of DCZ (0.1
mg/kg) intraperitoneally. For DREADD inhibition of PV neurons, a dose of 0.02 mg/kg was
used instead, as the 0.1 mg/kg dose caused significant increases in neocortical excitability (note
PV animals that had received a dose of 0.1 mg/kg were not used for any subsequent experiments
or in any of the datasets shown here). Injections were carefully administered to ensure the field
of view would not be disturbed; experiments were aborted if this occurred, as responses in each
period could no longer be directly compared. Activation of receptors was allowed to take effect
for fifteen minutes prior to resumption of recording, after which a second period of baseline
activity was recorded. This period lasted between fifteen and thirty minutes, such that a period of
fifteen minutes of stable activity was recorded prior to plasticity induction. Plasticity induction
protocols were followed as above (see *Optogenetic and tactile stimulation*), and whisker test
pulses were delivered for at least 40 minutes after plasticity induction. To ensure observed
plasticity phenomena were not the result of prolonged changes in activity related to hM4D(Gi) or
hM4D(Gq) activation, control experiments lasting the same duration as the plasticity

experiments were conducted similarly, except without induction of plasticity (see Figure S5). To
ensure DREADD expression was localized to the neocortex, viral spread was histologically
characterized in all animals (see *Tissue preparation for fluorescence microscopy and*
immunohistochemistry).

Characterization of neurons. Calcium images taken in the green (PMT0) and red (PMT1)
channels were first merged, trial-averaged, and concatenated using Fiji. The average image was
then calculated, followed by identification of all cells expressing tdTomato or mCherry
(applicable to all recordings except the zona incerta manipulation experiments). ROIs were
manually drawn around each cell co-expressing GCaMP6f and tdTomato or mCherry. All
images across a session were then concatenated and trial-averaged before calculating the
maximum image. ROIs were manually drawn around all putative pyramidal neurons, which were
identified with the following criteria: a triangular or pyramidal soma, the presence of a single
apical dendrite extending in a similar orientation to other pyramidal cells in the same imaging
session, and the presence of basal dendrites smaller in diameter than the apical dendrite. In a
subset of experiments in which apical dendrites could not be clearly distinguished in the imaging
plane, z-stack images were taken at the end of the imaging session, averaged, and referenced to
visualize apical dendrites. After identification of fluorescently tagged interneurons and
pyramidal neurons, relevant putative interneurons were morphologically identified and compiled
(e.g., putative VIP and SST neurons in PV-tagged mice). Putative interneurons were identified
by the presence of an elongated soma and bipolar dendritic arbors (likely VIP), round somas
smaller in diameter than pyramidal cells, and neurons with multiple perisomatic dendrites
(usually highly branched; likely PV). The morphology of untagged interneurons was visually
compared to the morphology of tagged interneurons in relevant mice (e.g., morphologically
identified interneurons in PVCre mice were compared to fluorescently tagged interneurons in
SSTCre and VIPCre mice). The average (11.07μm, SD 1.64μm) and maximum (18.14μm, SD
2.74μm) soma diameter for morphologically identified pyramidal neurons (n = 3358) were
significantly higher than in morphologically identified interneurons in the same mice (n = 3522;
average: 9.90μm, SD 1.96; maximum: 16.37μm, SD: 3.39μm; $p < 0.001$ for both values,
unpaired t-test) and tagged VIP, SST, and PV neurons (n = 543; average: 8.96μm, SD 1.87μm;
maximum: 14.98μm, SD 3.51μm; $p < 0.001$ for both values, unpaired t-test).

<u>**Intrinsic Signal Imaging**</u>

Image acquisition. Mice underwent the same surgical procedures described in *Two-Photon*
calcium imaging: Cranial window surgeries for imaging of barrel cortex neurons. To facilitate
manipulation of single whiskers, mice were lightly anesthetized with 10 mg/kg ketamine and
placed on a stereotaxic frame. Using a custom GUI written in MATLAB, small vessels were
visualized under a green LED (525 nm wavelength) with a high-speed camera (Teledyne
DALSA Genie Nano GigE). Focus in the z-plane was adjusted approximately 250 μm below the
putative pial surface before taking a reference image of the vasculature. For measurement of
intrinsic signals, the field of view was illuminated with red light (625 nm). An Arduino Uno and
MATLAB were used to trigger trial onset, the device used to mechanically deflect whiskers (a
glass capillary tube attached to a rotary motor), a blue LED to stimulate climbing fibers, and
recording of intrinsic signals at 30 Hz. Trials consisted of a 1-second baseline period, followed
by a 1-second test pulse in which whiskers contralateral to the optical window were
mechanically deflected at 8 Hz (RWS). This was repeated with an inter-trial interval of 18

seconds for 30 minutes prior to plasticity induction, after which whiskers were stimulated at 8 Hz
for 5 minutes. Test pulses were delivered for 40 minutes after plasticity induction. For RWS+CF
and control experiments, 50 ms blue light pulses were delivered at 1 Hz during plasticity
induction with or without 8 Hz co-stimulation of whiskers, respectively.

Image processing. Evoked signals were calculated similarly to Vasquez *et al* [94]. Briefly, images
were smoothed with a Gaussian filter and spatially downsampled by a factor of four. For each
trial, an average baseline reflectance image (R0) was generated by calculating the average across
the 1-second (30 frame) period before stimulus onset. Post-stimulus reflectance images were
temporally averaged across 200 ms bins for 600 ms total, starting 400 ms after stimulus onset.
The change in reflectance, ΔR/R, was calculated by subtracting R0 from each post-stimulus
image and dividing the result by R0. Each of the three bins (400-600 ms, 600-800 ms, and 800
ms-1 s post stimulus onset) were averaged across a minimum of 30 trials and finally summed to
yield a single total stimulus-evoked ΔR/R image. The reference image taken with green light
illumination was used to draw a region of interest around the cranial window. Pixels outside this
ROI mask were set to zero. Binary images were generated by calculating z-scores of the ΔR/R
image and thresholding values below a z-score of -1.5 (note negative changes in reflectance
indicate an increase in the presence of deoxyhemoglobin caused by increased oxygen
consumption by active neurons). Binarized images were overlaid onto reference images. To
calculate the areal extent of single whisker-evoked responses, clusters of active pixels were
identified by applying a median filter with a 5 x 5-pixel neighborhood size, and the number of
pixels above threshold was calculated. To reduce the impact of residual noise (i.e. detected pixels
outside the area reliably activated by a single whisker), values were normalized by the area of the
cranial window.

<u>**Two-photon image processing**</u>

Processing of cerebellar data. Processing of Purkinje cell calcium traces was conducted as
previously described[23, 73]. Individual trials from each session were concatenated along the z-axis,
representing a time series across the entire recording session, to ensure consistency among trials
after motion correction. The concatenated file underwent motion correction using a custom
MATLAB script (MATLAB R2017b, MathWorks, Natick, MA, USA) based on whole frame
cross-correlation (provided by the lab of Mark Sheffield, University of Chicago). Cellular
regions of interest were manually selected in Fiji based on the average field of view across trials.
Trials were then separated for further processing. In a subset of trials, optogenetic stimulus
artifacts were visible in a portion of the field of view in 1-3 nonconsecutive frames, depending
on the stimulus duration. These frames were manually identified and excluded using Fiji before
further processing.

Using custom MATLAB script, ΔF/F values were calculated from the raw trace using the
following equation: (Ft − F0)/F0, where Ft represented the raw calcium intensity of the time
series, and the baseline fluorescence (F0) was set as the 20th percentile of the fluorescence trace
from each trial. ΔF/F values were then subjected to low-pass filtering using a five-frame moving
window smoothing function. Finally, traces were normalized to a pre-stimulus baseline period by
subtracting the average ΔF/F value from the 5-frame period prior to stimulus onset from the
entire trace. To visualize population average responses to optogenetic stimulation of climbing
fibers, calcium signals from each ROI were first trial-averaged, followed by concatenation and

averaging of the signal across cells and mice. To calculate maximum amplitude of responses in
the evoked period, the maximum in a window 0-700 ms after stimulus onset was calculated for
each ROI's trial-averaged trace, then concatenated across cells and mice. Area under the curve
(AUC) was calculated similarly by integrating the ΔF/F values over the same window. To
identify spontaneous events, frame indices in which the first and second derivatives of the ΔF/F
signal reached 0.04 were identified (F^{idx}). Peaks were detected using a peak prominence
threshold of 0.1 ΔF/F, with spontaneous events defined as peaks occurring within 200 ms of F^{idx}.
The onset of each spontaneous event was determined as the first frame in which the first
derivative of the ΔF/F dropped below 0.05 (working backward from frame index of the detected
peak). A 1-second window following event onset was extracted. Finally, after averaging all
detected spontaneous events within ROIs, spontaneous events were compiled across neurons to
generate the population average trace.

Processing of S1 data. Motion correction and ROI curation were performed as above (see
Processing of cerebellar data and *Two-photon calcium imaging: Characterization of neurons*).
Residual optogenetic stimulus artifacts were observed in a single frame in a subset of trials,
typically at or immediately after stimulus offset. These artifacts were detected using the
findpeaks function in MATLAB and corrected by replacing the artifact frame's value with a
linear interpolation of the two adjacent frames before further processing. Background subtraction
was performed by calculating the bottom 1st percentile of the raw fluorescence trace collected
from an ROI that did not contain cell bodies or neuropil, then subtracting this value from all
trials. Smoothing, expression of raw signals as ΔF/F, calculation of baseline noise (σ), and
calculation of the signal-to-noise ratio (SNR) were performed as in Ayaz *et. al.*[95]. Smoothing
was performed with a 51-point 1st-order Savitsky-Golay filter, followed by calculation of the
relative percent change of fluorescence ((Ft − F0)/F0) using the 1st percentile of the smoothed
trace as F0. Given background subtraction prior to calculation of F0 may generate negative F0
values and subsequent signal inversion, ROIs with F0 < 0 were rejected from further analyses
(44/7423 cells). To calculate σ for each neuron, a 5-second (155 frame) sliding window was first
used to calculate the standard deviation of the fluorescence change during each 5-second period
within a session. After compiling all values, the 1st percentile value was taken as σ. For each
neuron, the SNR was defined as the 95th percentile of ΔF/F signals recorded across all trials
divided by the baseline noise.

Evoked event detection. To detect whisker–evoked events, we first applied OASIS
deconvolution to the ΔF/F traces using an event detection threshold of 3σ to generate a binary
trace containing event times. An autoregressive model with order p=1 was used to ensure the
number of events ("spikes") for each calcium transient varied with the amplitude of the calcium
transient as previously observed[52]. Next, for each frame, the sum of all events was calculated and
divided by the number of trials to determine the proportion of trials with an event in a given
frame (P^{event}). P^{event} was averaged across frames, and consecutive frames with P^{event} 3 standard
deviations above the mean were selected. This generated an evoked event window of 387 ms
(Supplementary Data 2B). Finally, for each neuron, the calcium traces for all trials containing an
event within the evoked event window were compiled for further analysis. To visualize
population average traces for all experiments, evoked calcium signals for each neuron were first
averaged across trials, followed by averaging across neurons. For visualizations between the
same neural populations in different conditions, only those cells that responded to both
conditions, i.e. having an evoked event in each condition, were included (following the definition

of "Persistent cells" used in Williams *et. al.*[60]). Shaded regions in population average traces
represent SEM.

Calculation of amplitude and area under the curve in S1 recordings. For all calculations of
AUC in S1 recordings, the ΔF/F signal was integrated over a 700 ms window after stimulus
onset unless otherwise specified (details below).

To quantify plasticity phenomena caused by RWS or RWS+CF, all evoked events measured
within 50 minutes prior to plasticity induction ("Pre") or 60 minutes after plasticity induction
("Post") were extracted for each cell and trial-averaged before calculating AUC. Only the
neurons with evoked events both pre- and post-plasticity induction were included in subsequent
analyses. AUC were averaged both across all neurons or within animal to perform statistics
across neurons or mice (see *Statistics and quantifications*). To visualize the duration of plasticity
effects, evoked events were binned into 10-minute epochs and trial-averaged within cells,
followed by calculation of the AUC and averaging across neurons. To verify the robustness of
plasticity effects, AUC and time course measurements were also analyzed across all cells and
trials (i.e., those that did not have stimulus-evoked events; Figure S2) and in the absence of
animal movement (Figure S3; see *Determination of active and rest trials*). To quantify plasticity
phenomena in DREADD manipulation experiments, response amplitude was calculated in
addition to AUC before and after plasticity induction. Response amplitude was taken as the
maximum ΔF/F value in the same 700 ms window used for AUC calculations.

To quantify the immediate effect of climbing fiber activation on basic whisker responses in S1
cortex, AUC was calculated similarly for whisker stimulation (W) or whisker and CF co-
activation (W+CF) trials. An additional measurement was taken 650-850 ms after stimulus onset
for pyramidal neurons and VIP interneurons, as the effect of CF modulation was present in the
later phase of the calcium response. Note that while W and W+CF test pulses were delivered
prior to DCZ administration in DREADD experiments to verify functional opsin expression,
these mice were excluded from this analysis due to an insufficient number of trials in each
animal.

To verify that hM4D(Gi) and hM3D(Gq) expression in SST neurons indeed suppressed or
increased their activity, respectively, AUC was calculated before and after DCZ administration
in a window 1750 ms after stimulus onset. This larger window was used to capture the
significantly increased duration of whisker-evoked calcium responses in SST neurons expressing
hM3D(Gq). To verify that hM4D(Gi) and hM3D(Gq) expression in VIP neurons indeed
suppressed or increased their activity, respectively, AUC was calculated before and after DCZ
administration in a window 650-850 ms after stimulus onset. Given VIP neurons are modulated
by climbing fiber stimulation in this phase of the calcium response (Figure 2g), this later window
was chosen to ensure DREADD manipulation of VIP neurons suppressed or enhanced VIP
responses in the physiologically relevant response period.

<u>Determination of active and rest trials</u>

Activity of mice was monitored at 30.98 Hz using a DALSA M640 CCD camera (Teledyne
Technologies, Thousand Oaks, CA, USA) focused on the animal's face and a portion of the
running wheel. To determine the frames in which mice were active, custom MATLAB script was
used to generate a binary trace that indicated the frames in which movement between frames

occurred (based on a comparison of pixel values between consecutive frames). For each
recording session, movement traces and corresponding videos from ten randomly selected trials
were manually inspected to ensure all frames in which mice were whisking or running were
indicated. Note whisking occurred independently of running, but running seldom occurred
without whisking. Rest trials were defined as any trial with no movement detected at least 400
ms before and after stimulus onset – longer than the evoked event window, and the same as the
window chosen in Ayaz *et. al.*[95] to distinguish rest from running onset. All other trials were
considered active trials.

Statistics and quantifications

All statistics and quantifications were performed using MATLAB. To assess normality, we
applied the Lilliefors test to each dataset before performing statistical calculations. For paired
samples, a paired t-test was used when data were normally distributed, and a Wilcoxon signed-
rank test was used for non-normal data. For unpaired comparisons, an unpaired t-test was used
for normally distributed data, a Welch's t-test for normally distributed data with unequal
variances, and a Wilcoxon rank-sum test for non-normal distributions. Categorical variables
were compared using the Chi-squared test. Two-way ANOVA was used for multi-group
comparisons (i.e., to determine whether plasticity effects across time were significantly different
between groups). To perform statistics across all animals, average responses were first
normalized by dividing the response in the relevant experimental condition by the baseline
response. A one-sample t-test was then performed against a standard value of 1. Statistical
significance was set to $p < 0.05$, with p values presented in figures as follows: $p < 0.05$: *;
$p < 0.01$: **; $p < 0.001$: ***; $p > 0.05$: not significant (n.s.). Bar graphs represent mean ± SEM.
Box plots contain the interquartile range (25th and 75th percentiles) and median.

Resource availability

Lead contact
924   Further information and requests for resources and reagents should be directed and will be
925   fulfilled by the lead contact, Christian Hansel (chansel@bsd.uchicaho.edu).

Data availability
928   All data shown here, as well as additional two-photon and confocal imaging data described in the
929   Methods, will be placed on GitHub prior to publication.

Code availability
931   All code used for the analyses will be provided prior to publication, with accession codes made
932   available for public access.

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Declaration of Interests

The authors declare no competing interests.

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Figure 1. Optogenetic CF co-activation prevents adaptive L2/3 pyramidal cell potentiation in S1 cortex. (**A**) Experimental schematic for (**D** – **L**). (**B** and **C**) ChR2 expression in the inferior olive (IO) and optogenetic CF activation evokes calcium transients in PCs. (**B**) Calbindin staining of PCs marks the cerebellum (shown here in the same plane as the IO; white arrowhead). Scale bar: 1mm. (**C**) Top view of PC dendrites with calcium responses evoked by 470nm, 50ms LED pulses in PCs. Scale bar: 100μm. Right: Distribution of response amplitudes (box: 25th and 75th percentiles; line: median). (**D**) GCaMP6f expression in S1 cortex. Scale bar: 1mm. (**E**) Sagittal view of ChR2-expressing CF terminals in cerebellum that are activated in optogenetic experiments. Scale bar: 100μm. (**F**) Left: Example of the morphological identification of L2/3 pyramidal cells by apical dendrites (green arrowheads) and INs (white arrowheads). Scale bar: 100μm. Right: Pyramidal cell at higher magnification. Scale bar: 15μm. (**G**) Pyramidal cell plasticity after RWS (N=12 mice) and after pairing with CF activation (RWS+CF; N=11 mice). Right: Quantification of evoked responses across neurons. (**H**) Data in (**G**) binned across time. Blue shading: RWS/RWS+CF. (**I**) Quantification across mice, normalized to the average baseline whisker response. (**J-L**) Analyses for INs in the same mice. Data: mean ± SEM. Scale bars: 0.2 ΔF/F; 0.5s.

Figure 2. Optogenetic CF co-activation differentially modulates basic responses to whisker stimulation in different types of neocortical neurons. (**A**) Schematic of the recording configuration (left) and S1 cortex microcircuit (right). (**B**) CF co-activation with whisker stimulation does not significantly alter response amplitudes in morphologically identified L2/3 pyramidal neurons (n=826 cells; N=23 mice) in the default analysis window (0-700ms; light shade), but responses decrease in the late response window (650-850ms; dark shade). Left: averaged traces. Middle: analysis by neuron. Shades distinguish analyzed time windows. Right: analysis by mouse (0-700ms). Blue line indicates mean. (**C**) CF co-activation significantly enhances responses in putative interneurons (INs; n=1104 cells; N=24 mice). (**D** – **F**) Sample two-photon field of view (left) and post-hoc verification of expression in S1 using confocal microscopy (right). Arrowheads indicate neurons co-expressing GCaMP6f and tdTomato in VIP interneurons (**D**), SST interneurons (**E**), or PV interneurons (**F**). Scale bars: 100μm (left); 500μm (right). (**G**) CF co-activation with whisker stimulation significantly reduces VIP population responses in the late analysis window (n=129 cells, N=7 mice; shades as in **B**), but enhances responses in SST interneurons (n=102 cells; N=12 mice; **H**) and PV interneurons (n=177 cells; N=18 mice; **I**). Middle plots show analysis by neuron, right plots analysis by mouse (blue line indicates mean). Scale bars: 0.1 ΔF/F; 0.5s. Data: mean ± SEM.

Figure 3. SST interneurons are recruited to prevent L2/3 pyramidal cell plasticity. (**A**) Circuit diagram highlighting the chemogenetic activation of SST interneurons and recording configuration. (**B**) Expression of activating hM3D(Gq) receptors is localized to neocortical SST interneurons. Scale bar: 100μm (top) and 500μm (bottom). (**C**) DCZ application enhances responses in SST interneurons (n=12 cells; N=7 mice). (**D**) In the presence of DCZ, RWS stimulation suppresses L2/3 pyramidal cell potentiation, and even causes depression, even when the CF is not co-activated (n=255 cells; N=4 mice). Left: average traces; middle: analysis by cell (mean ± SEM); right: response range by cell (whiskers: range; box: 25th and 75th percentiles; line: median). (**E**) Circuit diagram highlighting the chemogenetic inhibition of SST interneurons and recording configuration. (**F**) Expression of inhibiting hM4D(Gi) receptors is localized to neocortical SST interneurons. Scale bars as in (**B**). (**G**) DCZ application reduces responses in SST interneurons (n=53 cells; N=6 mice). (**H**) In the presence of DCZ, pyramidal cell potentiation is observed, despite CF-co-activation (n=183 cells; N=3 mice). Data analysis presented as in (**D**). Scale bars: 0.2 ΔF/F; 0.5s.

Figure 4. VIP interneurons mediate the effects of optogenetic CF activation. (**A**) Circuit diagram
highlighting the chemogenetic inhibition of VIP interneurons and recording configuration. (**B**) Expression
of inhibiting hM4D(Gi) receptors is localized to neocortical VIP interneurons. Scale bars: 100μm (top)
and 500μm (bottom). (**C**) DCZ application reduces responses in VIP interneurons (n=10 cells; N=3 mice).
(**D**) In the presence of DCZ, RWS stimulation suppresses L2/3 pyramidal cell potentiation, and even
causes depression, even when the CF is not co-activated (n=189 cells; N=3 mice). Left: averaged traces;
middle: analysis by cell (mean ± SEM); right: response range by cell (range; box: 25th and 75th
percentiles; line: median). (**E**) Circuit diagram highlighting the chemogenetic activation of VIP
interneurons and recording configuration. (**F**) Expression of activating hM3D(Gq) receptors is localized
to neocortical VIP interneurons. Scale bars as in (**B**). (**G**) DCZ application enhances responses in VIP
interneurons (n=14 cells; N=3 mice). (**H**) In the presence of DCZ, pyramidal cell potentiation is observed,
despite CF co-activation (n=202 cells; N=4 mice). Data analysis presented as in (**D**). Scale bars: 0.1 ΔF/F;
0.5s.

Figure 5. VIP interneurons orchestrate the inhibitory network in S1. (**A**) Circuit diagram highlighting
the chemogenetic inhibition of VIP interneurons and recording configuration. (**B**) Inhibiting hM4D(Gi)
receptors are expressed in neocortical VIP interneurons. Note A-B are identical to Fig. 4.A-B. (**C**) In the
presence of DCZ, RWS stimulation activates L2/3 interneurons, even when the CF is not co-activated
(n=27 cells; N=3 mice). Left: averaged traces; middle: analysis by cell (mean ± SEM); right: response
range by cell (whiskers: range; box: 25th and 75th percentiles; line: median). (**D**) Circuit diagram
highlighting the chemogenetic activation of VIP interneurons and recording configuration. (**E**) Activating
hM3D(Gq) receptors are expressed in neocortical VIP interneurons. Note D-E are identical to Fig. 4E-F.
(**F**) In the presence of DCZ, IN depression is observed, despite CF co-activation (n=105 cells; N=4 mice).
Data analysis presented as in (**C**). Scale bars: 0.2 ΔF/F; 0.5s.

Figure 6. A pathway from the cerebellar nuclei via the zona incerta and thalamic nucleus POm to S1 cortex. (**A**) Schematic of dual-injection strategy to label outputs of zona incerta (ZI) neurons receiving input from contralateral cerebellar nuclei. Scale bar: 1mm. (**B**) EYFP expression in ZI (left) with a higher-magnification image (same slice). Right: labeled axons in the posterior medial nucleus (POm; bottom; scale bar: 20μm) and S1 cortex (top; scale bar: 10μm). (**C**) Distribution of projections from cerebellar nuclei to the ZI along the rostro-caudal axis; dotted line separates ventral (vZI) and dorsal ZI (dZI). Scale bars: 500μm. (**D**) Labeling strategy for recordings shown in (**E**). Scale bar: 1mm. (**E**) Top: Whole-cell patch-clamp recordings from POm-projecting ZI neurons receiving innervation from cerebellar nuclei. Responses to depolarizing current pulses in the absence (left) and presence (right) of TTX (1μM). Bottom: Responses to photostimulation of ChR2-expressing terminals from cerebellar nuclei neurons. (**F**) Amplitude (left) of photostimulation-evoked EPSPs and latency (right) in control recordings (n=5) and after wash-in of TTX (n=3). (**G**) Experimental configuration for recordings from mice expressing inhibiting hM4D(Gi) receptors in PV-expressing ZI interneurons. (**H**) In the presence of DCZ, the suppressive effect of CF co-activation on basic whisker responses in L2/3 pyramidal neurons is blocked (n=80 cells; N=3 mice). (**I**) Bar graphs: analysis by neuron. Data: mean ± SEM. Scale bars: 0.1 ΔF/F; 0.5s.

Supplementary Figure 1. Optogenetic CF activation causes GCaMP6f-encoded calcium transients in Purkinje cells. (**A**) Schematic of IO injection. Top left: Cerebellum (calbindin-staining of PCs) and IO (as in Fig 1B). Bottom: ChR2 expression in the IO principal (IOPr) and dorsal IO (IOD). Scale bars: 1mm (top) and 100μm (bottom). (**B**) Overlap of VGluT2-stained CF terminals with ChR2 expression in the contralateral cerebellar cortex. Scale bar: 40μm. (**C**) Post-hoc verification of ChR2 expression in CF terminals in crus I and II. Scale bars: 50μm. (**D**) Measure of distance of the pial surface to the cerebellar nuclei. While the IO innervates both the cerebellar cortex and nuclei directly, it is unlikely that CF terminals other than the superficial ones innervating PCs are optogenetically activated as the intensity of blue light decreases exponentially in brain tissue. Scale bar: 300μm. Dashed line: 2mm. (**E**) Experimental schematic for F-I. (**F**) Test of various stimulus protocols: 3x15ms, 2x20ms, 1x50ms (interval 50ms) and (3x15ms)*2 (interval 50ms). Traces show response averages (n=102 cells from one mouse). (**G**) Bar graphs showing response peaks (left) and integrals (right) from the corresponding data in f. Data: mean ± SEM. (**H**) Calcium transients evoked by 470nm, 50ms LED light pulses in PCs in crus I/II (n=131 cells; N=2 mice) in comparison to spontaneous events (left). Scale bars: 0.1 ΔF/F; 0.5s. (**I**) Amplitude and latency to peak measures from the corresponding data in h. Data: mean ± SEM. Scale bars: 0.1 ΔF/F; 0.5s.

Supplementary Figure 2. Plasticity in different neuron types. (A) Experimental schematic. (B) Event
probability around stimulus onset with the window for detection of stimulus-evoked events (red; three
standard deviations above the mean event probability, calculated across all frames collected during the
20-second trial periods). (C) Pie chart indicating the percentage of L2/3 pyramidal cells recruited or
suppressed by RWS/RWS+CF activity (only responsive to whisker stimulation in post- or pre- periods,
respectively) and how many remained entirely unresponsive. (D) Left: RWS-evoked response changes in
L2/3 pyramidal neurons (n=329 cells; N=12 mice). Right: RWS+CF-evoked response changes in L2/3
pyramidal neurons (n=597 cells; N=11 mice). Bar graphs quantify calcium responses across all cells and
trials (i.e. including unresponsive neurons and trials without whisker-evoked responses). (E) The same as
D in INs in the same mice (n=616 cells for RWS; n=590 cells for RWS+CF). (F, G) Same as (D, E) in
tdTomato-tagged SST interneurons (F; n=41 cells and N=3 mice for RWS; n=31 cells and N=3 mice for
RWS+CF) and PV interneurons (G; n=83 cells; N=5 mice for RWS; n=113 cells; N=5 mice for
RWS+CF). Scale bars: 0.05 ΔF/F; 0.5s. Data: mean ± SEM.

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Supplementary Figure 3. Impact of RWS- and RWS+CF-stimulation on animal motion. (**A**)
Experimental schematic. Mice are awake and head-fixed but can walk/run on the treadmill. (**B**) Average
cross-correlation coefficient between neural calcium signals and movement activity data, segregated by
neuron type. The numbers below the box charts indicate the number of recorded cells. (**C**) Left:
probability distribution of animal movement in Rest trials, demonstrating the absence of movement both
before and after stimulus onset. Right: probability distribution of animal movement in Active trials. (**D**)
Pie charts indicating the percentage of rest, active and stop trials pre-vs post-RWS (left) or RWS+CF
(right). (**E**) Average probability of animal movement in Rest trials pre- and post-RWS (left) or RWS+CF
(right) over the same time scale as the neural traces shown below. (**F**) and (**H**) RWS-evoked potentiation
in L2/3 pyramidal neurons (n=124; N=12) and RWS+CF-mediated suppression of potentiation (n=388;
N=11) in the absence of animal movement (i.e. in Rest trials). (**G**) and (**I**) Analysis for INs in the same
mice (n=212 cells for RWS; n=296 cells for RWS+CF). Scale bars: 0.1 ΔF/F; 0.5s. Data: mean ± SEM.
Note motion activity traces capture both whisking and running activity. Scale bars: 0.1 ΔF/F; 0.5s.

Supplementary Figure 4. Intrinsic optical imaging confirms CF-mediated suppression of S1
plasticity. (**A**) Experimental schematic. Intrinsic optical imaging is performed using a high-speed camera.
The field of view is illuminated with red light (625 nm). A decrease in reflectance indicates increased
oxygen consumption by neurons activated by single whisker stimulation (blue arrow). (**B**) Top: Example
of S1 plasticity evoked by stimulation of the C2 whisker at 10Hz. Bottom: Example of suppression of
response potentiation when whisker stimulation is paired with optogenetic CF co-activation at 1Hz. (**C**)
Quantification of RWS-evoked potentiation of the single whisker response area (blue; N=4 mice) and
RWS+CF-mediated suppression of response potentiation (red; N=4 mice). No response potentiation is
observed in the absence of repeated whisker stimulation or CF activation (grey; N=4 mice). Data are
normalized to the baseline response area. Light blue line indicates mean.

Supplementary Figure 5. Chemogenetic manipulation of interneuron activity: plasticity effects in L2/3 pyramidal cells vs control. (**A**) Chemogenetic activation of SST interneurons. Left: Experimental schematic. Middle: circuit diagram. Right: DCZ injection activates SST interneurons (yellow). Top: Control protocol. 'Post' marks the period equivalent in time to the period for the measurement of plasticity effects after RWS (bottom). (**B**) Response amplitudes in L2/3 pyramidal neurons during Pre, DCZ wash-in, and Post periods in control experiments. (**C**) The difference in response (ΔAUC) of L2/3 pyramidal cells to whisker stimulation between the DCZ wash-in and Post periods; (control: n=218 cells; N=3 mice; plasticity: n=253 cells; N=4 mice); negative ΔAUC values indicate AUC values in the Post period are lower than in the DCZ wash-in period. The ΔAUC in control and plasticity experiments are normalized to the absolute ΔAUC in control experiments to demonstrate changes in the plasticity experiments relative to controls. Note the absence of PN potentiation in the plasticity experiment. (**D**) – (**F**) Equivalent measures during chemogenetic VIP interneuron inhibition (red; control: n=173 cells; N=3 mice; plasticity: n=164 cells; N=3 mice). Note the absence of PN potentiation in the plasticity experiment. (**G**) – (**I**) Equivalent measures during chemogenetic SST interneuron inhibition (control: n=175 cells; N= 3 mice; plasticity: n=181 cells; N=3 mice). (**J**) – (**L**) Equivalent measures during chemogenetic PV interneuron inhibition (control: n=368 cells; N=4 mice; plasticity: n=258 cells; N=3 mice). PN neurons are significantly potentiated after plasticity experiments beyond changes in controls. (**m**) – (**o**) Equivalent measures during chemogenetic VIP interneuron activation (control: 145 cells; N=3 mice; plasticity: n=197 cells; N=4 mice). PN neurons are significantly potentiated after plasticity experiments beyond changes in controls. Optogenetic CF co-activation is applied during RWS in the experiments shown in (**G**) – (**O**). Data: mean ± SEM.

Supplementary Figure 6. Cerebellar nuclei do not strongly project to thalamic reticular nucleus.
(**A**) Schematic of labeling approach (top) and injection site of retrograde tracer (bottom). Scale bar: 1mm.
(**B**) Image of retrograde label from the somatosensory thalamus (POm and VPM; cyan) and orthograde
label from the contralateral cerebellar nuclei (red) taken approximately -1.0mm posterior from bregma,
demonstrating lack of overlap between neurons in the thalamic reticular nucleus (TRN) and cerebellar
nuclei terminals in the adjacent thalamic nuclei. Representative images are shown from two separate
mice. Scale bars: 1mm (left images) and 250μm (right images). (**C**) Projection diagram. DCN: deep
cerebellar nuclei; ZI: zona incerta; POm: posterior medial thalamic nucleus; TRN: thalamic reticular
nucleus; VPM: ventral posterior medial nucleus; VPL: ventral posterior lateral nucleus; VA: ventral
anterior nucleus; VL: ventral lateral nucleus.

Supplementary Figure 7. Transsynaptic labeling identifies projections from the cerebellar nuclei to
POm. (**A**) Schematic of dual-injection approach to label outputs of ZI neurons receiving input from the
contralateral CBN. (**B**) Top left: EYFP-expressing terminals and cell bodies within ZI (dashed lines) and
axonal projections out of the ZI (arrowheads). Scale bar: 100μm. Bottom left: projections from the ZI
terminate in POm (arrowheads). Scale bar: 50 μm. Right: zoomed-out view showing projections from ZI
to POm in the same plane. Scale bar: 200 μm. (**C**) ZI neurons receiving cerebellar input project to S1.
Synaptic terminals are observed in L1 and L5a. Scale bar: 100 μm. (**D**) Series of coronal sections showing
details of the projections from the contralateral CBN to ZI and POm. Scale bars: 1mm. (**E**) The same as D
in a second mouse, showing robust EYFP expression across the rostro-caudal extent of ZI. Scale bars:
1mm. POt = posterior thalamic nucleus (triangular part). Colored dots mark images taken from the same
sections.

Supplementary Figure 8. Whole-cell patch-clamp recordings are performed from neurons in the ZI
that project to POm and receive input from cerebellar nuclei. (**A**) Left: Differential interference
contrast (DIC) image showing the ZI (white dashed circle) at low magnification (top; scale bar: 500μm),
and a neuron selected for patching at higher magnification (bottom; scale bar: 100μm). Middle left: same
images showing ChR2-expression. Middle right: same images showing retrograde fluoro-Ruby label
originating in the POm. Right: merged images (boxed area with yellow dashed line marks the recordings
area shown in the images below). (**B**) Post-hoc histology of an acute slice used for recordings
demonstrating the details of cerebellar innervation of the ZI, separated by dorsal ZI (dZI) and ventral ZI
(vZI; image corresponds to Fig. 5D, separated by channel). Scale bar: 500μm. (**C**) Quantification of the
terminal fluorescence segregated by dZI and vZI. Data: mean ± SEM.

	panel	mean	median	error	n cells	N mice	test	P value	result	measurement (units)	cell type
	1.C	0.554	0.5004	0.023	102	1	N/A	N/A	N/A	Amplitude (ΔF/F)	PC
1.G	Pre (RWS)	3.9581	2.3941	0.3824	287†	12	Wilcoxon	0.0010	**	AUC ((ΔF*t)/F)	PN
	Post (RWS)	5.246	2.4812	0.6725							
	Pre (RWS+CF)	3.2676	1.9988	0.1811	534†	11	Wilcoxon	0.1126	n.s.		
	Post (RWS+CF)	2.6892	2.0035	0.1049							
1.H	-40 min (RWS)	4.65488	N/A	0.5261	144†	10	Two-way ANOVA (Group comparison)	<0.001	***	AUC ((ΔF*t)/F)	PN
	-30 min (RWS)	3.91113		0.4083	174†	11					
	-20 min (RWS)	3.65761		0.3639	172†	11					
	-10 min (RWS)	3.91723		0.4911	213†	12					
	0 min (RWS)‡	4.19536		0.7388	100†	7					
	10 min (RWS)‡	5.77819		0.8127	212†	12					
	20 min (RWS)‡	6.48013		1.7059	217†	12					
	30 min (RWS)‡	5.91206		0.7591	155†	10					
	40 min (RWS)‡	5.64781		0.8828	157†	10					
	50 min (RWS)‡	5.16045		0.6085	117†	9					
	60 min (RWS)‡	5.02273		0.7503	54†	4					
	- 40 min (RWS+CF)	3.32904		0.2981	351†	10					
	-30 min (RWS+CF)	3.60127		0.3131	317†	10					
	- 20 min (RWS+CF)	3.69046		0.2739	364†	10					
	-10 min (RWS+CF)	3.05876		0.2269	394†	11					
	0 min (RWS+CF)‡	3.87257		0.423	204†	9					
	10 min (RWS+CF)‡	2.37707		0.1063	479†	11					
	20 min (RWS+CF)‡	2.61932		0.1256	511†	11					
	30 min (RWS+CF)‡	2.74055		0.1435	483†	10					
	40 min (RWS+CF)‡	2.32432		0.1067	477†	10					
	50 min (RWS+CF)‡	2.49962		0.1297	362†	6					
	60 min (RWS+CF)‡	2.80165		0.233	352†	4					
1.I	Pre (RWS)	1	N/A	0	287	12†	One-sample t-test	0.0104	*	AUC ((ΔF*t)/F)	PN
	Post (RWS)	1.2629		0.0853							
	Pre (RWS+CF)	1		0	534	11†	One-sample t-test	0.003	**		
	Post (RWS+CF)	0.7723		0.0584							
1.J	Pre (RWS)	3.8936	2.7519	0.1701	570†	12	Wilcoxon	<0.001	***	AUC ((ΔF*t)/F)	IN
	Post (RWS)	3.0852	2.2592	0.1194							
	Pre (RWS+CF)	3.5683	2.3499	0.2021	516†	11	Wilcoxon	<0.001	***		
	Post (RWS+CF)	3.8983	2.5878	0.1797							
1.K	-40 min (RWS)	4.70089	N/A	0.286	342†	10	Two-way ANOVA (Group comparison)	0.0127	*	AUC ((ΔF*t)/F)	IN
	-30 min (RWS)	3.69953		0.2221	380†	11					
	-20 min (RWS)	4.15268		0.266	324†	11					
	-10 min (RWS)	3.83042		0.21	446†	12					
	0 min (RWS)‡	4.50767		0.3892	207†	7					
	10 min (RWS)‡	3.47265		0.1809	438†	12					
	20 min (RWS)‡	3.31423		0.1744	434†	12					
	30 min (RWS)‡	3.47938		0.1832	311†	10					
	40 min (RWS)‡	3.66121		0.1901	335†	9					
	50 min (RWS)‡	3.24975		0.2246	239†	9					
	60 min (RWS)‡	3.16086		0.2327	147†	4					
	- 40 min (RWS+CF)	3.93255		0.3137	320†	10					
	-30 min (RWS+CF)	4.4692		0.396	283†	11					
	- 20 min (RWS+CF)	5.02758		0.3417	344†	10					
	-10 min (RWS+CF)	3.95778		0.278	376†	11					
	0 min (RWS+CF)‡	4.11703		0.4043	219†	10					
	10 min (RWS+CF)‡	3.7648		0.2121	444†	11					
	20 min (RWS+CF)‡	3.87636		0.2234	479†	11					
	30 min (RWS+CF)‡	3.90208		0.2237	423†	10					
	40 min (RWS+CF)‡	3.37561		0.1953	405†	10					
	50 min (RWS+CF)‡	3.8697		0.2619	314†	6					
	60 min (RWS+CF)‡	4.25666		0.2688	273†	4					
1.L	Pre (RWS)	1	N/A	0	570	12†	One-sample t-test	<0.001	***	AUC ((ΔF*t)/F)	IN
	Post (RWS)	0.7983		0.0252							
	Pre (RWS+CF)	1		0	516	11†	One-sample t-test	0.2233	n.s.		
	Post (RWS+CF)	1.1321		0.1018							

† indicates whether statistics performed across cells or mice
‡ time points included in ANOVA

Data S1.
Data and statistical tests corresponding to Figure 1.

	panel	mean	median	error	n cells	N mice	test	P value	result	measurement (units)	cell type
2.B (c)	W (0-700ms)	3.5038	2.3648	0.1358	826†	23	Wilcoxon	0.1856	n.s.	AUC ((ΔF*t)/F)	PN
	W+CF (0-700ms)	3.2344	2.2292	0.1314							
	W (650-850ms)	1.0935	0.4922	0.0672			Wilcoxon	0.0487	*		
	W+CF (650-850ms)	0.937	0.4763	0.0611							
2.C (r)	W	1	N/A	0	826	23†	One-sample t-test	0.3709	n.s.	AUC ((ΔF*t)/F)	PN
	W+CF	0.954		0.0504							
2.C (c)	W	4.0841	2.8209	0.1357	1104†	24	Wilcoxon	<0.001	***	AUC ((ΔF*t)/F)	IN
	W+CF	4.7268	3.1189	0.1637							
2.C (r)	W	1	N/A	0	1104	24†	One-sample t-test	<0.001	***	AUC ((ΔF*t)/F)	IN
	W+CF	1.1961		0.0406							
2.G (c)	W (0-700ms)	2.883	2.529	0.1128	129†	7	Wilcoxon	0.8665	n.s.	AUC ((ΔF*t)/F)	VIP
	W+CF (0-700ms)	2.9589	2.5207	0.1438							
	W (650-850ms)	1.0888	0.9519	0.0545			Wilcoxon	0.0255	*		
	W+CF (650-850ms)	1.0294	0.8138	0.0713							
2.G (r)	W	1	N/A	0	129	7†	One-sample t-test	0.4113	n.s.	AUC ((ΔF*t)/F)	VIP
	W+CF	1.047		0.0533							
2.H (c)	W	2.9453	2.3916	0.18	102†	12	Wilcoxon	0.001	**	AUC ((ΔF*t)/F)	SST
	W+CF	3.2532	2.7647	0.2152							
2.H (r)	W	1	N/A	0	102	12†	One-sample t-test	0.0035	**	AUC ((ΔF*t)/F)	SST
	W+CF	1.1582		0.0427							
2.I (c)	W	2.4758	2.269	0.107	177†	18	Wilcoxon	0.0032	**	AUC ((ΔF*t)/F)	PV
	W+CF	2.6768	2.369	0.1092							
2.I (r)	W	1	N/A	0	177	18†	One-sample t-test	0.0127	*	AUC ((ΔF*t)/F)	PV
	W+CF	1.1122		0.0403							

† indicates whether statistics performed across cells or mice

Data S2.
Data and statistical tests corresponding to Figure 2.

	panel	mean	median	error	n cells	N mice	test	P value	result	measurement (units)	cell type
3.C	Pre	5.2386	4.0088	0.8243	12	7	Wilcoxon	0.0425	*	AUC ((ΔF*t)/F)	SST
	DCZ	9.2671	5.9757	2.3962							
3.D (c)	Pre	5.7312	2.927	0.5212	255	4	Wilcoxon	<0.001	***	AUC ((ΔF*t)/F)	PN
	Post	3.106	1.9677	0.2207							
3.D (r)	Pre	0.3938	0.1934	0.0358	255	4	Wilcoxon	<0.001	***	Amplitude (ΔF/F)	PN
	Post	0.2139	0.1403	0.015							
3.G	Pre	8.9616	7.5298	1.1235	53	6	Wilcoxon	0.0057	**	AUC ((ΔF*t)/F)	SST
	DCZ	7.5387	6.4485	0.7883							
3.H (c)	Pre	7.8992	4.0365	1.0087	183	3	Wilcoxon	<0.001	***	AUC ((ΔF*t)/F)	PN
	Post	9.9871	4.5967	1.2894							
3.H (r)	Pre	0.5334	0.267	0.0682	183	3	Wilcoxon	<0.001	***	Amplitude (ΔF/F)	PN
	Post	0.6759	0.3062	0.0872							
4.C	Pre	1.1339	0.8779	0.3452	10	3	Paired t-test	0.0272	*	AUC ((ΔF*t)/F)	VIP
	DCZ	0.2521	0.2239	0.04							
4.D (c)	Pre	2.6299	1.8696	0.2739	189	3	Wilcoxon	<0.001	***	AUC ((ΔF*t)/F)	PN
	Post	1.4749	1.2773	0.0768							
4.D (r)	Pre	0.179	0.1242	0.02	189	3	Wilcoxon	<0.001	***	Amplitude (ΔF/F)	PN
	Post	0.1021	0.0849	0.0055							
4.G	Pre	1.3677	0.2754	0.8043	14	3	Wilcoxon	0.0494	*	AUC ((ΔF*t)/F)	VIP
	DCZ	1.9361	0.4533	1.0391							
4.H (c)	Pre	2.53	1.9726	0.1644	202	4	Wilcoxon	<0.001	***	AUC ((ΔF*t)/F)	PN
	Post	3.2011	2.2506	0.2818							
4.H (r)	Pre	0.1813	0.1324	0.0142	202	4	Wilcoxon	<0.001	***	Amplitude (ΔF/F)	PN
	Post	0.2309	0.1529	0.0226							
5.C (c)	Pre	2.0677	1.4505	0.2728	27	3	Wilcoxon	0.0174	*	AUC ((ΔF*t)/F)	IN
	Post	2.5314	2.1002	0.2536							
5.C (r)	Pre	0.1400	0.1096	0.0181	27	3	Wilcoxon	0.0198	*	Amplitude (ΔF/F)	IN
	Post	0.1740	0.1366	0.0181							
5.F (c)	Pre	3.7801	2.5438	0.3629	105	4	Wilcoxon	<0.001	***	AUC ((ΔF*t)/F)	IN
	Post	2.9725	2.2871	0.2231							
5.F (r)	Pre	0.2656	0.1778	0.0263	105	4	Wilcoxon	<0.001	***	Amplitude (ΔF/F)	IN
	Post	0.2052	0.1572	0.0162							

Data S3.

Data and statistical tests corresponding to Figures 3, 4, and 5.

panel		mean	median	error	n cells	N mice	test	P value	result	measurement (units)	cell type
6.F	baseline	1.2386	N/A	0.11	5	5	Welch's t-test	<0.001	***	EPSP amplitude (mV)	ZI neuron
	+TTX	0.4094		0.081	3	3					
	baseline	3.4		0.4159	5	5	Welch's t-test	0.62	n.s.	Latency (ms)	
	+TTX	3.633		0.1333	3	3					
6.I	W (Baseline)	5.1648	3.2461	0.778	80	3	Wilcoxon	0.0183	*	AUC ((ΔF*t)/F)	PN
	W+CF (Baseline)	4.7113	3.1395	0.7239							
	W (DCZ)	3.6838	2.5494	0.6804			Wilcoxon	<0.001	***		
	W+CF (DCZ)	4.1972	2.7795	0.6946							

Data S4.

Data and statistical tests corresponding to Figure 5.

panel		mean	median	error	n cells	N mice	test	P value	Result	measurement (units)	cell type
S1.G (l)	3x15	0.3455	N/A	0.0198	102	1	N/A	N/A	N/A	Amplitude (ΔF/F)	PC
	2x20	0.3213		0.0177							
	1x50	0.554		0.023							
	(3x15)*2	0.3616		0.0171							
S1.G (r)	3x15	3.5376	N/A	0.2585	102	1	N/A	N/A	N/A	AUC ((ΔF*t)/F)	PC
	2x20	3.8521		0.2483							
	1x50	5.1302		0.3019							
	(3x15)*2	4.3732		0.261							
S1.I (t)	Spontaneous	0.3768	0.3777	0.0078	131	2	Paired t-test	0.785	n.s.	Amplitude (ΔF/F)	PC
	Evoked	0.3831	0.3184	0.0245							
S1.I (b)	Spontaneous	308.355	290.51	10.308	131	2	Wilcoxon	0.0875	n.s.	Latency to peak (ms)	PC
	Evoked	332.761	258.231	20.567							

Data S5.

Data and statistical tests corresponding to Supplementary Figure 1.

panel	mean	median	error	%	n cells	N mice	test	P value	result	measurement (units)	cell type
S2.C Persistent (RWS) Recruited (RWS) Suppressed (RWS) Unresponsive (RWS)	N/A	N/A	N/A	87.23 6.991 4.255 1.52	329†	12	Chi-squared	0.0103	*	percent total cells (%)	PN
Persistent (RWS+CF) Recruited (RWS+CF) Suppressed (RWS+CF) Unresponsive (RWS+CF)				89.45 8.71 1.508 0.335	597†	11					
S2.D Pre (RWS) Post (RWS)	1.0374 1.3728	0.4502 0.5007	0.1950 0.1998	N/A	329†	12	Wilcoxon	<0.001	***	AUC ((ΔF*t)/F)	PN
Pre (RWS+CF) Post (RWS+CF)	1.1012 0.9152	0.4755 0.5405	0.1203 0.0697		597†	11	Wilcoxon	0.9575	n.s.		
S2.E Pre (RWS) Post (RWS)	1.4409 0.9525	0.7726 0.4952	0.116 0.0832	N/A	616†	12	Wilcoxon	<0.001	***	AUC ((ΔF*t)/F)	IN
Pre (RWS+CF) Post (RWS+CF)	1.4741 1.5336	0.4905 0.643	0.1679 0.1391		590†	11	Wilcoxon	<0.001	***		
S2.F Pre (RWS) Post (RWS)	1.4276 1.1308	1.0298 0.9495	0.1611 0.1378	N/A	41†	3	Wilcoxon	0.0048	**	AUC ((ΔF*t)/F)	SST
Pre (RWS+CF) Post (RWS+CF)	1.1075 1.0747	0.6072 0.5804	0.294 0.1862		31†	3	Wilcoxon	0.7539	n.s.		
S2.G Pre (RWS) Post (RWS)	0.6099 0.5004	0.3682 0.3811	0.0767 0.0651	N/A	83†	5	Wilcoxon	0.0359	*	AUC ((ΔF*t)/F)	PV
Pre (RWS+CF) Post (RWS+CF)	0.5372 0.5028	0.3252 0.3527	0.0779 0.0580		113†	5	Wilcoxon	0.5456	n.s.		

† indicates whether statistics performed across cells or mice

Data S6.
Data and statistical tests corresponding to Supplementary Figure 2.

panel		mean	median	error	n cells	N mice	test	P value	result	measurement (units)	cell type
S3.B	PN	0.0078	-0.0019	0.0023	1147	24				xCorr (r)	PN
	IN	0.0332	0.0123	0.0023	1620	35				xCorr (r)	IN
	PV	-1.47E-04	-2.72E-04	0.0045	221	13	N/A	N/A	N/A	xCorr (r)	PV
	SST	0.041	0.0354	0.0099	91	8				xCorr (r)	SST
	VIP	0.0909	0.0836	0.0078	173	11				xCorr (r)	VIP
S3.D (top l)	Rest	29.61		9.7967							
	Active	69.84		9.9334							
	Stop	0.55		0.2902			12†	Chi-squared	0.01071	*	
S3.D (bottom l)	Rest	24.35		9.5786							
	Active	75.29		9.6393						percent total trials (%)	N/A
	Stop	0.36	N/A	0.2324	N/A						
S3.D (top r)	Rest	33.46		12.1959							
	Active	66.54		12.1959							
	Stop	0		0			11†	Chi-squared	<0.001	***	
S3.D (bottom r)	Rest	42.45		13.1339							
	Active	57.32		13.1921							
	Stop	0.24		0.1441							
S3.H	Pre (RWS)	4.0059	2.1655	0.7509	124†	12	Wilcoxon	0.0147	*		
	Post (RWS)	5.7207	2.2396	1.5931						AUC ((ΔF*t)/F)	PN
	Pre (RWS+CF)	2.677	1.7382	0.1927	388†	11	Wilcoxon	0.5104	n.s.		
	Post (RWS+CF)	2.4344	1.7958	0.1195							
S3.I	Pre (RWS)	3.3914	2.3096	0.2246	212†	12	Wilcoxon	0.0033	**		
	Post (RWS)	2.965	2.0138	0.2076						AUC ((ΔF*t)/F)	IN
	Pre (RWS+CF)	3.3041	2.2275	0.2025	296†	11	Wilcoxon	<0.001	***		
	Post (RWS+CF)	3.5204	2.2994	0.1901							

† indicates whether statistics performed across cells or mice

Data S7.
Data and statistical tests corresponding to Supplementary Figure 3.

panel		mean	error	N mice	test	P value	result	measurement (units)
S4.C	RWS (Pre)	1	0	4	one-sample t test	0.0471	*	Whisker responsive area
	RWS (Post)	3.6978	0.8273					
	RWS+CF (Pre)	1	0	4		0.1138	n.s.	
	RWS+CF (Post)	0.5195	0.2171					
	control (Pre)	1	0	4		0.0649	n.s.	
	control (Post)	1.5067	0.1176					

Data S8.

Data and statistical tests corresponding to Supplementary Figure 4.

panel		mean	median	error	n cells	N mice	test	P value	result	measurement (units)
S5.B	Pre	2.8467	2.507	0.1745	220†	3	Pre vs DCZ: Wilcoxon	Pre vs DCZ: <0.001	Pre vs DCZ: ***	AUC $((\Delta F * t)/F)$
	DCZ	1.587	1.2817	0.0809	218†					
	Post	1.7015	1.4958	0.0726	220†		DCZ vs Post: Wilcoxon	DCZ vs Post: <0.001	DCZ vs Post: ***	
S5.C	Control	0	0.0212	0.0631	218†	3	Wilcoxon	<0.001	***	Relative ΔAUC $(\Delta((\Delta F * t)/F))$
	Plasticity	-0.14	-0.2199	0.0509	253†	4				
S5.E	Pre	4.2185	3.0214	0.3601	181†	3	Pre vs DCZ: Wilcoxon	Pre vs DCZ: <0.001	Pre vs DCZ: ***	AUC $((\Delta F * t)/F)$
	DCZ	2.9159	2.2613	0.2104	177†					
	Post	2.5923	2.1445	0.1419	177†		DCZ vs Post: Wilcoxon	DCZ vs Post: 0.0422	DCZ vs Post: *	
S5.F	Control	0	-0.0686	0.0655	173†	3	Wilcoxon	0.6742	n.s.	Relative ΔAUC $(\Delta((\Delta F * t)/F))$
	Plasticity	-2.469	-0.0227	2.9139	164†	3				
S5.H	Pre	4.3825	2.7846	0.425	181†	3	Pre vs DCZ: Wilcoxon	Pre vs DCZ: 0.0018	Pre vs DCZ: **	AUC $((\Delta F * t)/F)$
	DCZ	4.3789	3.374	0.3058	185†					
	Post	3.8364	2.8455	0.2325	180†		DCZ vs Post: Wilcoxon	DCZ vs Post: <0.001	DCZ vs Post: ***	
S5.I	Control	0	-0.2341	0.0623	175†	3	Wilcoxon	0.0928	n.s.	Relative ΔAUC $(\Delta((\Delta F * t)/F))$
	Plasticity	0.1359	0.0253	0.0409	181†	3				
S5.K	Pre	3.093	2.0421	0.2373	363†	4	Pre vs DCZ: Wilcoxon	Pre vs DCZ: < 0.001	***	AUC $((\Delta F * t)/F)$
	DCZ	9.0378	6.7442	0.3663	368†					
	Post	6.381	4.7294	0.2822	369†		DCZ vs Post: Wilcoxon	Pre vs DCZ: <0.001	***	
S5.L	Control	0	-0.0262	0.0189	368	4	Wilcoxon	<0.001	***	Relative ΔAUC $(\Delta((\Delta F * t)/F))$
	Plasticity	0.6522	0.0344	0.0633	258	3				
S5.N	Pre	2.4242	1.9103	0.2445	136†	3	Pre vs DCZ: Wilcoxon	Pre vs DCZ: <0.001	Pre vs DCZ: ***	AUC $((\Delta F * t)/F)$
	DCZ	3.1922	2.6613	0.2031	137†					
	Post	3.0724	2.5155	0.2135	137†		DCZ vs Post: Wilcoxon	DCZ vs Post:0.0902	DCZ vs Post: n.s.	
S5.O	Control	0	0.0021	0.022	145†	3	Wilcoxon	<0.001	***	Relative ΔAUC $(\Delta((\Delta F * t)/F))$
	Plasticity	0.1723	0.0982	0.0351	197†	4				

† indicates whether statistics performed across cells or mice

Data S9.
Data and statistical tests corresponding to Supplementary Figure 5.