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Traumatic brain injury (TBI) causes cortical dysfunction and can lead to post-traumatic epilepsy. Multiple studies demonstrate that GABAergic inhibitory network function is compromised following TBI, which may contribute to hyperexcitability and motor, behavioral, and cognitive deficits. Preserving the function of GABAergic interneurons, therefore, is a rational therapeutic strategy to preserve cortical function after TBI and prevent long-term clinical complications. Here, we explored an approach based on the ketogenic diet, a neuroprotective and anticonvulsant dietary therapy which results in reduced glycolysis and increased ketosis. Utilizing a pharmacologic inhibitor of glycolysis (2-deoxyglucose, or 2-DG), we found that acute in vitro application of 2-DG decreased the excitability of excitatory neurons, but not inhibitory interneurons, in cortical slices from naïve mice. Employing the controlled cortical impact (CCI) model of TBI in mice, we found that in vitro 2-DG treatment rapidly attenuated epileptiform activity seen in acute cortical slices 3 to 5 weeks after TBI. One week of in vivo 2-DG treatment immediately after TBI prevented the development of epileptiform activity, restored excitatory and inhibitory synaptic activity, and attenuated the loss of parvalbumin-expressing inhibitory interneurons. In summary, 2-DG may have therapeutic potential to restore network function following TBI.

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**Title: Glycolytic inhibitor 2-deoxyglucose prevents cortical hyperexcitability after
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Short Title: 2-DG treatment after traumatic brain injury

One Sentence Summary: Following traumatic brain injury in mice, in vivo treatment with the glycolytic inhibitor 2-deoxyglucose prevented cortical network pathology including cortical hyperexcitability, changes in synaptic activity, and loss of parvalbumin-expressing GABAergic interneurons.

Abstract

Traumatic brain injury (TBI) causes cortical dysfunction and can lead to post-traumatic epilepsy. Multiple studies demonstrate that GABAergic inhibitory network function is compromised following TBI, which may contribute to hyperexcitability and motor, behavioral, and cognitive deficits. Preserving the function of GABAergic interneurons, therefore, is a rational therapeutic strategy to preserve cortical function after TBI and prevent long-term clinical complications. Here, we explored an approach based on the ketogenic diet, a neuroprotective and anticonvulsant dietary therapy which results in reduced glycolysis and increased ketosis. Utilizing a pharmacologic inhibitor of glycolysis (2-deoxyglucose, or 2-DG), we found that acute in vitro application of 2-DG decreased the excitability of excitatory neurons, but not inhibitory interneurons, in cortical slices from naïve mice. Employing the controlled cortical impact (CCI) model of TBI in mice, we found that in vitro 2-DG treatment rapidly attenuated epileptiform activity seen in acute cortical slices 3-5 weeks after TBI. One week of in vivo 2-DG treatment immediately after TBI prevented the development of epileptiform activity, restored excitatory and inhibitory synaptic activity, and attenuated the loss of parvalbumin-expressing inhibitory interneurons. In summary, 2-DG may have therapeutic potential to restore network function following TBI.

Introduction

Traumatic brain injury (TBI) is a leading cause of death and disability worldwide and can lead to motor, behavioral, and cognitive losses (1) as well as post-traumatic epilepsy (PTE) (2-4). Patients who develop PTE show higher mortality than TBI patients without epilepsy, and many PTE patients are refractory to the currently available anticonvulsive therapies (2, 4). Multiple molecular and cellular changes that occur after TBI may play an important role in the pathophysiology of network dysfunction, PTE, and other TBI-induced pathologies. Aberrant electrical activity (5, 6), indiscriminate release of neurotransmitters (7), and altered cerebral glucose utilization (8, 9) all occur after TBI. Additionally, compromised cerebral blood flow, disruption of the blood-brain-barrier (10, 11), increased oxidative stress, and mitochondrial dysfunction (12) create a complex set of neurometabolic disruptions after injury (1, 2, 13). Together, these changes can disturb synaptic function, cause hard-wired remodeling of neuronal connectivity, and lead to circuit dysfunction and PTE.

Many studies suggest that a key step in developing network dysfunction after injury is the loss of inhibitory GABAergic interneurons, a diverse group of neurons that powerfully constrain cortical excitation. Disruption of inhibitory network activity can lead to motor dysfunction, cognitive losses, and epilepsy (14-18). The most abundant subtype of cortical interneurons express the calcium-binding protein parvalbumin (PV), fire action potentials at a high frequency (fast-spiking), and provide perisomatic inhibition of excitatory cortical pyramidal neurons (19). PV+ interneuron loss, atrophy, and dysfunction have been shown in the controlled cortical impact (CCI) (20), lateral fluid percussion (14), partially isolated neocortex (“undercut”) (21, 22), and weight drop (23) models of TBI. Importantly, human post-mortem studies also show that PV-expressing interneurons are specifically lost following TBI (24). The mechanisms by which TBI leads to

interneuron loss are largely unknown. Therefore, it has been difficult to develop therapeutic interventions to preserve inhibitory network function following TBI.

We believe that the aberrant neuronal activity that occurs acutely after TBI is a key contributor to long-term cortical network dysfunction. In support of this idea, early seizure activity and other EEG abnormalities are associated with poor clinical outcomes and a higher risk of developing post-traumatic epilepsy (5, 6, 25). Preclinical data from animal models of epilepsy and brain injury demonstrate that early uncontrolled excitation contributes to later loss of GABAergic interneurons and development of inhibitory circuit dysfunction (26-29). Following TBI in both humans and animal models, this aberrant electrical activity occurs during periods of abnormally elevated glycolytic activity (30-35). We hypothesize that reducing hyper-glycolysis will attenuate injury-induced increases in excitatory synaptic activity, ameliorate losses of PV+ interneurons, and ultimately prevent inhibitory network dysfunction.

2-deoxyglucose (2-DG) is a glucose analog that competitively inhibits hexokinase (the rate-limiting enzyme in glycolysis) (36-38). Multiple studies demonstrate that 2-DG can be anticonvulsive in both in vitro and in vivo models of epilepsy (39, 40). 2-DG pre-treatment also prevents kainic acid-induced hippocampal cell loss in rats (41) and can reduce seizure severity (42, 43). While 2-DG has been previously explored as an anticonvulsant therapy, it has not yet been utilized as a disease-modifying agent to prevent epileptogenesis after TBI.

In this study, we tested whether 2-DG reduces TBI-induced hyperexcitation and attenuates the development of inhibitory network dysfunction after TBI. We utilized the controlled cortical impact (CCI) model of focal brain contusion, which results in significant cell loss, network hyperexcitability, and spontaneous behavioral and electrographic seizures (44). Our study shows that acute 2-DG treatment of cortical brain sections attenuated hyperexcitability in the injured

brain. In addition, 2-DG treatment reduced the excitability of excitatory neurons, but not inhibitory interneurons. Finally, in vivo treatment with 2-DG for one week following CCI prevented the development of epileptiform activity and reduced the loss of parvalbumin-expressing interneurons. Together, these studies support the potential application of glycolytic inhibitors to reduce cortical hyperexcitability after TBI.

Results

In vitro 2-DG treatment decreases the excitability of excitatory neurons, but not inhibitory interneurons

2-DG has been previously shown to attenuate epileptiform activity in multiple in vitro and animal models of epilepsy (39, 40, 45). Pharmacologic inhibition of glycolysis can directly suppress synaptic transmission, reduce membrane resistance (46), and broaden the action potential (AP) waveform (47). It is unknown, however, whether glycolytic inhibition has differential effects on the excitability of glutamatergic excitatory neurons and GABAergic inhibitory interneurons. Therefore, we assessed the effects of 2-DG on both cell types in naïve (uninjured) cortex. Whole-cell recordings were established in acute cortical brain slices from adult (>P28) animals of both sexes. Recordings were performed on layer V pyramidal neurons or fast-spiking PV+ GABAergic interneurons. PV interneurons were identified using the *G42* mouse line, a BAC transgenic line in which >70% of genetically GFP-labeled interneurons are PV+ on immunohistochemistry (Figure S1, (48)). In current clamp mode, hyperpolarizing and depolarizing steps were injected in the presence of synaptic receptor blockers (10 μ M CPP, 20 μ M DNQX, and 10 μ M gabazine) and the number of evoked action potentials (APs) was quantified. The results were compared between baseline aCSF (containing 10 mM glucose) and following 10 minutes of slice perfusion with 2-DG-aCSF (8 mM 2-DG, 2 mM glucose). 2-DG-aCSF maintained the same solution osmolarity as baseline aCSF (10 mM total saccharide), while constricting glucose availability to concentrations similar to those present in mouse or human cerebrospinal fluid (2 mM, (49)). In the presence of 2-DG, excitatory layer V pyramidal neurons fired significantly fewer APs at a given level of current injection (Figure 1A, 1C; see Table S1 for all Linear Mixed Model statistical results) while GABAergic interneurons were not affected (Figure 1B, 1D). Consistent with reduced excitability

of glutamatergic excitatory neurons following 2-DG perfusion, rheobase (current injection required to elicit the first AP) was increased in excitatory neurons but was not affected in GABAergic interneurons (Figure 1E, 1H). The decrease in the intrinsic excitability of excitatory neurons was associated with a decrease in membrane resistance which was not observed in GABAergic interneurons (Figure 1F, 1I). No change in resting membrane potential was observed in either cell type upon application of 2-DG (Figure 1G, 1J).

To control for reduced 2 mM glucose in 2-DG-aCSF, these experiments were repeated in low glucose aCSF (LG-aCSF: 2 mM glucose, 8 mM sucrose (a metabolically inert sugar)). LG-aCSF had no effects on AP firing, input-output curves, or membrane resistance in either excitatory neurons or inhibitory interneurons (Figure S2). This demonstrates that simply lowering glucose to 2 mM is not sufficient to alter neuronal excitability. Together, these experiments show that inhibition of glycolysis has cell type-specific effects on neuronal excitability.

2-DG ameliorates CCI-induced increases in synaptic excitation onto GABAergic interneurons

Acutely after TBI, glycolytic activity and neuronal activity are aberrantly increased (35). We hypothesized that TBI results in increased synaptic excitation of GABAergic interneurons in the peri-injury cortex acutely after injury. If so, this hyperexcitation may contribute to later cell death or dysfunction of PV+ (and other GABAergic) interneurons (see Discussion). To test this hypothesis, we utilized controlled cortical impact (CCI) to model focal, contusional TBI in adult male mice. Briefly, a craniectomy was made over left sensorimotor cortex and an impact was delivered to generate a moderate-to-severe injury (3 mm impactor probe diameter, 3.5 m/s velocity, 400 ms dwell time, 1 mm impact depth) (44). Acute cortical slices were prepared from

sham- and CCI-injured brains 3 days following injury and whole-cell recordings were made from fast-spiking GABAergic interneurons within 200 μm of the lateral edge of the CCI lesion. Interneurons were identified with GFP labeling in layer V-VI cortex in *G42* mice, which as above, were primarily PV+ (Figure S1). Neurons were voltage-clamped at -70 mV to isolate spontaneous excitatory postsynaptic currents (sEPSCs). Consistent with our hypothesis, there was an increase in the mean frequency of sEPSCs after CCI compared to sham (Figure 2B). CCI also resulted in a significant leftward shift in the sEPSC inter-event interval cumulative distribution (Figure 2C). There was a trend towards an increase in sEPSC amplitude, without statistical significance based on LMM or 2-sample Kolmogorov-Smirnov test (Figure 2F-G).

Because glucose utilization is elevated acutely after TBI and we had previously observed specific effects of 2-DG on excitatory (but not inhibitory) neuronal activity, we assessed whether 2-DG application to the acute slices would reduce injury-induced excitation of GABAergic interneurons. When 2-DG-aCSF was applied to acute cortical slices prepared 3 days following CCI, the mean frequency of sEPSCs onto interneurons was reduced to approximately sham levels (Figure 2B), corresponding with a rightward shift the sEPSC inter-event interval cumulative distribution (Figure 2D). Application of 2-DG to sham tissue did not significantly affect sEPSC frequency. There was, however, a large variability in the response to 2-DG treatment in sham animals and a trend toward more frequent excitatory events (data not shown), demonstrating unique effects of 2-DG in sham versus CCI-injured brain. 2-DG treatment also resulted in a leftward shift of the cumulative event amplitude following CCI (Figure 2H), although the comparisons of the mean amplitude by LMM and cumulative distributions by 2-sample Kolmogorov-Smirnov test did not reach statistical significance. Overall, these results suggest that

GABAergic interneurons receive increased excitation in the days following TBI and that these effects are attenuated by acute application of 2-DG.

Acute 2-DG treatment attenuates epileptiform activity in cortical brain slices following CCI

Next, we tested whether 2-DG application attenuates network-level cortical hyperexcitability, which is observable in field excitatory postsynaptic potentials (fEPSPs) as early as 2 weeks following TBI (17), and is robustly present by 3-5 weeks following TBI. Brain slices were prepared from mice 3-5 weeks after CCI or sham surgery and electrical stimulation was delivered to ascending cortical inputs to evoke fEPSPs. In sham animals, fEPSPs were brief and low-amplitude. Slices taken 3-5 weeks after CCI, however, exhibited stimulus-evoked fEPSPs with epileptiform activity (increased amplitude, duration, area, and coastline), consistent with network hyperexcitability ((20), Figure 3A). As shown previously in other slice models of epileptiform activity (40), in vitro application of 2-DG markedly reduced the percentage of stimulus-evoked fEPSP traces that exhibited epileptiform activity after CCI from 90% to 18% (Figure 3B). Additionally, 2-DG treatment resulted in field potentials with significantly decreased area (Figure 3C) and high-frequency activity as measured by fEPSP coastline (Figure 3D). In brain slices from sham-injured animals, 2-DG-aCSF did not significantly change the area or coastline of the fEPSPs, or the percentage of epileptiform traces. In low glucose conditions (LG-aCSF), post-CCI epileptiform activity was not attenuated (Figure S3). These results show that in vitro application of the glycolytic inhibitor 2-DG rapidly attenuates cortical network hyperexcitability 3-5 weeks after TBI.

In vivo 2-DG treatment attenuates epileptiform cortical activity following CCI

We next hypothesized that *in vivo* 2-DG treatment during a critical period following TBI reduces subsequent cortical hyperexcitability. To test this hypothesis, we treated mice with 2-DG throughout the first week after injury (250 mg/kg or vehicle intraperitoneally, once daily for seven days following CCI or sham surgery) and assessed network hyperexcitability utilizing cortical field recordings. Acute cortical brain slices were prepared 3-5 weeks following the initial surgery (2-4 weeks following the end of 2-DG treatment), when epileptiform activity is expected in the CCI animals (20). As predicted, sham-injured slices from both treatment groups had normal fEPSPs (Figure 4A-B). Consistent with previous studies, acute cortical brain slices from vehicle-treated CCI-injured animals exhibited epileptiform stimulus-evoked fEPSPs (Figure 4C). However, *in vivo* 2-DG treatment following CCI dramatically attenuated epileptiform fEPSPs (from 95% to 13%) in CCI-injured animals (Figure 4D-E). Area and coastline measurements of the evoked fEPSPs following CCI also exhibited significant attenuation with *in vivo* 2-DG treatment (Figure 4F-G).

CCI also disrupts the input/output (I/O) relationship of the cortical network. Cortical I/O responses were assessed by comparing fEPSP area evoked by 1X and 2X threshold stimulation. In the uninjured brain, cortical networks generated graded output in response to increased stimulation. After CCI, however, cortical networks generated all-or-none responses and lacked graded output in response to stimulation (Figure 4H). *In vivo* 2-DG treatment partially restored the disrupted I/O relationship, resulting in graded fEPSP amplitude and area in response to increased stimulation. 2-DG treatment had a large effect size on I/O relationship (0.54 compared to the effect of CCI, -0.81) but did not reach statistical significance by LMM, likely due to the high degree of variability (Figure 4H). Together, these experiments show that *in vivo* 2-DG treatment

during the first week after injury attenuates network hyperexcitability and reduces long-term TBI-associated cortical network dysfunction.

In vivo 2-DG treatment transiently affects systemic glucose metabolism, but does not affect body temperature

Since systemic metabolic changes, such as hypothermia, may have independent neuroprotective effects after TBI, we assessed the effects of *in vivo* 2-DG on whole body metabolism. After a single 2-DG injection (250 mg/kg, intraperitoneal) in fasted animals, blood glucose levels were transiently increased (Figure 4I) and β -hydroxybutyrate (a ketone body) levels were transiently decreased (Figure 4J), consistent with decreased glucose utilization and increased ketosis. These changes in blood glucose and β -hydroxybutyrate returned to baseline by 4 hours post-injection. There was also a transient decrease in home cage locomotor activity that resolved to vehicle levels after the first 6 hours post-injection (Figure 4K). Importantly, there was no change in flank temperature in 2-DG-injected animals versus vehicle-injected controls (Figure 4L), suggesting that hypothermia does not contribute to the neuroprotective effect of 2-DG.

Then, daily 2-DG injections were performed in naïve animals for one week to parallel the *in vivo* dosing regimen post-CCI. Animals were assessed at rest, immediately before their daily 2-DG injection (~24 hours following their most recent injection). There were no changes observed in blood glucose, blood β -hydroxybutyrate, flank temperature, body weight, or daily food intake between vehicle- and 2-DG-treated animals across the entire week of treatment (Figure S4). For all experiments reported in this study, *in vivo* treatment with 2-DG did not affect animal weight or

recovery from surgery. These data suggest that there were no significant adaptive metabolic changes across the week-long in vivo 2-DG treatment.

In vivo 2-DG treatment prevents CCI-induced dysregulation of excitatory and inhibitory synaptic activity

TBI causes significant disruption of cortical synaptic activity including increased glutamatergic excitation and reduced GABAergic inhibition of excitatory cortical neurons (20), accordant with network hyperexcitability. In order to test whether in vivo 2-DG treatment ameliorates post-CCI changes in synaptic activity, we recorded spontaneous excitatory and inhibitory postsynaptic currents onto perilesional layer V pyramidal neurons 3-5 weeks following CCI. Consistent with previous studies, CCI induced a significant increase in mean spontaneous excitatory postsynaptic current (sEPSC) frequency (Figure 5A), resulted in a significant leftward shift in the sEPSC inter-event interval cumulative distribution, and had no effect on sEPSC amplitude (Figure 5C). Inhibitory synaptic transmission was decreased by CCI, again consistent with previous studies. CCI decreased mean spontaneous inhibitory postsynaptic current (sIPSC) frequency (Figure 5G), caused a significant rightward shift in the sIPSC inter-event interval cumulative distribution, and significantly increased mean sIPSC amplitude (Figure 5I).

Consistent with its effects on network hyperexcitability, in vivo 2-DG treatment significantly attenuated CCI-induced increases in excitation. 2-DG treatment reduced mean sEPSC frequency (Figure 5A, 5E) and caused a significant rightward shift in the sEPSC inter-event interval cumulative distribution (Figure 5B), with no effect on sEPSC amplitude (Figure 5C-D). In vivo treatment with 2-DG also restored inhibitory synaptic transmission after CCI. 2-DG

had a large, but not significant, effect on mean sIPSC frequency (Figure 5G), caused a significant leftward shift in the sIPSC inter-event interval cumulative distribution (Figure 5H), significantly reduced mean sIPSC amplitude (Figure 5I), and caused a leftward shift in the cumulative distribution of sIPSC amplitude (Figure 5J). Interestingly, we noted a difference in the distribution of mean sIPSC frequency following CCI; in cortical pyramidal cells recorded after sham injury, there was consistently a population of neurons (5 out of 25 cells in the Sham+Vehicle group) that received sIPSCs at a high rate (>12 Hz). This population was absent after CCI (0 out of 21 cells in CCI+Vehicle) but was present after in vivo 2-DG treatment (7 out of 21 cells in CCI+2-DG), suggesting that CCI causes a loss of robust network inhibition that is prevented by 2-DG treatment. Together, these data demonstrate that in vivo 2-DG treatment ameliorates network deficits in spontaneous excitatory and inhibitory neurotransmission caused by CCI.

In vivo 2-DG treatment attenuates the loss of parvalbumin-expressing interneurons following CCI

Because 2-DG treatment attenuated the development of epileptiform activity and prevented the decrease in sIPSCs after CCI, we hypothesized that it also prevents the loss of parvalbumin-expressing interneurons that had been previously reported (20). An additional question that we wished to address was whether CCI caused a loss of PV cells, or merely a loss of PV protein. To test our hypothesis and address the loss of PV cells versus protein, we used a combination of genetic and immunohistochemical approaches. *PV^{Cre}* mice (50), in which Cre-recombinase is expressed under the control of the *Pvalb* promoter, were crossed with *Ai9* reporter mice (51),

producing *PV^{Cre}/Ai9* mice which express tdTomato (tdT) in cells containing Cre. This provides a genetic label of PV cells, even if the PV protein itself is lost.

PV^{Cre}/Ai9 mice underwent CCI or sham surgery and were treated in vivo with 2-DG or vehicle as described above. Brains were prepared for immunohistochemical analysis 3-5 weeks post-CCI. First, lesion volume was approximated across serial sections in CCI animals treated with vehicle or 2-DG. In vivo 2-DG treatment had no significant effect on cavitory lesion volume (2-sample t-test; $n = 5$ animals/group, $p = 0.557$). Then, the densities of PV-immunolabeled and tdT-labeled neurons were quantified in five 100- μ m-wide rectangular regions of interest (ROIs) in the cortex lateral to the site of CCI injury (or in isotopic cortex in sham animals). Consistent with previous studies (20), the density of PV-immunolabeled cells was significantly reduced in the cortex of vehicle-treated CCI animals relative to sham (Figure 6A-E, 6G). PV+ cell density was significantly reduced in the first 400 μ m adjacent to the lesion (Figure 6G). A similar result was seen after counting tdT-labeled neurons (Figure 6H) although the decrease was significant only in the first 200 μ m lateral to the lesion. These results show that CCI causes both a loss of PV cells and a decrease in PV immunoreactivity adjacent to the lesion.

In vivo 2-DG treatment attenuated the loss of PV+ neurons relative to vehicle-treated CCI animals (Figure 6F-G). 2-DG prevented PV+ cell loss in the first 300 μ m adjacent to the lesion. 2-DG did not globally reduce the loss of tdT+ neurons, although a significant increase in tdT+ neurons was observed in the 100 μ m closest to the site of injury (Figure 6H), suggesting a focal effect of 2-DG. These results indicate that 2-DG broadly attenuates the loss of PV expression after CCI and focally reduces the loss PV (tdT+) neurons.

In order to further examine the effects of CCI and 2-DG on PV cell loss, versus changes in PV protein expression, we assessed the proportion of tdT-labeled neurons that had also been identified as immuno-positive for PV. In sham-injured animals, approximately 50% of tdT-labeled neurons were PV immuno-positive. The abundance of tdT-labeled neurons which lacked PV immunolabeling resembles that found in similar studies (52), and is partially attributable to the high threshold that the blinded experimenter used to identify a “PV+” cell by immunohistochemistry. The ~50% colocalization of tdT with the PV immunolabel may also result from PV expression below the detection limit of an immunolabeling approach, transient expression of PV during development, or non-specific Cre expression. After CCI, a significantly higher proportion of tdT-labeled neurons lacked PV immunoreactivity (Figure 6I-J) with the effect being most prominent in the first 200 μ m adjacent to the lesion. This suggests that a subset of PV (tdT+) interneurons remain after CCI but lose their expression of PV protein. Because PV is a calcium binding protein that is necessary for physiologic interneuron AP firing, loss of PV expression is consistent with decreased inhibitory function and cortical hyperexcitability (53, 54). In vivo treatment with 2-DG restored the ratio of PV (tdT+) cells expressing PV to levels seen in sham animals (Figure 6J). Together, these results show that CCI causes a loss of PV (tdT+) interneurons and reduces PV expression in remaining PV interneurons. In vivo 2-DG treatment broadly increases PV expression in surviving PV interneurons, consistent with enhanced inhibitory function.

Discussion

2-DG, a hexokinase inhibitor that reduces glycolytic activity, has been previously explored for its potential neuroprotective, anticonvulsant, and anti-epileptogenic properties, but the mechanisms of its rapid actions on neuronal function are unknown. Here, we demonstrate that 2-DG treatment attenuates excitatory, but not inhibitory, cortical neuron excitability. This provides a potential mechanism by which 2-DG mediates its acute anticonvulsant effects. Furthermore, we show that 2-DG has therapeutic potential to treat traumatic brain injury (TBI). While glycolysis is known to be abnormally elevated acutely following brain injury (55), it is unknown whether early metabolic disruptions contribute to chronic cortical hyperexcitability. Using the CCI model of TBI combined with electrophysiological approaches, we find that excitation onto inhibitory interneurons is elevated 3 days after injury and can be attenuated by acute 2-DG treatment. Further, we find that in vivo 2-DG treatment during the first week after TBI prevents epileptiform network activity, restores aspects of synaptic function, and reduces parvalbumin-positive inhibitory interneuron loss associated with CCI. Together, our work provides novel insight into acute synaptic dysfunction following TBI, implicates glycolysis in the hyperexcitation of inhibitory interneurons after TBI, and shows that in vivo 2-DG treatment reduces post-traumatic synaptic and network dysfunction.

Our findings, summarized above, demonstrate a complex set of synaptic changes following CCI, in both the acute (Figure 2) and chronic (Figure 5) settings. Acutely after TBI (3 days post-CCI), we observe increased excitation of PV interneurons that we suspect contributes to later inhibitory dysfunction, although our study does not confirm that interneuron loss and dysfunction occur as a direct result of hyperexcitation. Alternatively, increased excitation of interneurons acutely after CCI may represent a compensatory mechanism to increase inhibitory control of

cortical activity in the injured brain. If true, however, one would predict that 2-DG's reduction of IN hyperexcitation would be detrimental to TBI-related outcomes. We did not observe any increases in adverse events, seizures, or mortality during 2-DG treatment in our studies, consistent with 2-DG's known neuroprotective effects and suggesting that acute hyperexcitation of interneurons is, in fact, pathologic. Further work is required to clarify how early aberrant excitation of interneurons relates to and may result in long-term network pathology, including changes in parvalbumin expression, altered interneuron function and synaptic output, and interneuron cell loss.

Next, in the chronic period (3-5 weeks post-CCI), we observed changes in both excitatory and inhibitory synaptic activity that were ameliorated with in vivo 2-DG treatment. Consistent with our previous work (20), CCI increased spontaneous synaptic excitation and decreased spontaneous synaptic inhibition onto perilesional layer V pyramidal neurons. Both of these changes were attenuated by in vivo 2-DG treatment, suggesting that 2-DG ameliorates synaptic dysfunction after TBI. In line with the sIPSC frequency data we report, we also found that in vivo 2-DG treatment reduced the loss of parvalbumin-expressing interneurons. While changes in inhibitory cell density may contribute to our synaptic findings, changes in inhibitory cell function may also play a role in the overall cortical hyperexcitability evident 3-5 weeks after CCI. For example, loss of parvalbumin expression can cause inhibitory dysfunction, hyperexcitability, and worsened seizure phenotype (53, 56, 57), independent of loss of inhibitory synapses. Additionally, altered chloride transport (reduced KCC2/increased NKCC1 expression (58-62)) and changes in GABA receptor subunit composition and density (63-66) may further contribute to altered inhibitory circuit function. Thus, complex changes in both excitation and inhibition at the synaptic, cellular, and circuit levels result in cortical dysfunction and hyperexcitability after TBI.

Our study demonstrates that in vivo 2-DG treatment significantly ameliorates many of the effects of CCI on synaptic and network activity, and thus, strongly supports that modulation of metabolism in the injured brain may have therapeutic potential to improve patient outcomes by maintaining proper network function. Of note, we do not directly demonstrate that in vivo 2-DG treatment reduces brain glycolysis after TBI. 2-DG is well-established as an inhibitor of glycolysis (67) across multiple organ systems, animal models, and disease states (68-75). We observe acute changes in blood glucose and ketone body levels (Figure 4I-J) upon administration of 2-DG, consistent with 2-DG's ability to inhibit glycolysis and decrease glucose utilization. While a recent publication suggests that systemic delivery of 2-DG does not result in appreciable changes in brain hexokinase activity (76), multiple studies demonstrate that 2-DG treatment broadly inhibits glycolysis in whole brain and brain slices (36, 39, 40, 77). Systemic delivery of 2-DG has been shown to result in transcriptional changes consistent with reduced glycolysis in the brain, including reductions in hexokinase, PGK-1, and LDH-A expression in the rat hippocampus (39). Furthermore, TBI-associated breach of the blood-brain-barrier (78) may allow greater penetration of 2-DG into the brain, resulting in improved inhibition of brain glycolysis. While we observe robust CNS effects after delivering 2-DG post-CCI, it is possible that they are mediated through a peripheral effect of 2-DG or through a mechanism not directly related to hexokinase inhibition.

From a translational standpoint, the therapeutic rationale to reduce glycolysis after TBI is well-founded on both preclinical and clinical literature (35). Using PET scanning, studies report that a majority of patients exhibit increased cerebral glucose utilization following severe TBI (55, 79) that later develops into a chronic, hypometabolic state (80-82). Additionally, patients with aberrant glucose utilization and network activity during the first week post-TBI have worsened outcomes 6 months later (83). Preclinical work from multiple animal models of TBI shows early

increases in glucose utilization (33, 84-87) which are activity-dependent (34). Consistent with focal hyper-glycolysis and network dysfunction, our previous studies show heightened glutamate signaling and PV+ interneuron loss in the peri-injury cortex (20). Finally, there is abundant evidence supporting the use of the ketogenic diet (which results in decreased glycolysis and increased ketosis) as a strategy to manipulate neural network activity in animals and humans (reviewed in (88-94)). While dietary approaches and glycolytic inhibition have shown efficacy in controlling seizure activity, manipulations of glucose metabolism have not yet been harnessed effectively in the context of TBI. In our studies, we did not find evidence of chronic changes in animal health, systemic metabolism, or body temperature (Figure S4) over the course of 2-DG treatment. This is important as long-term exposure to 2-DG (>2 weeks) has been previously associated with cardiotoxicity (95, 96) and hypothermia alone may neuroprotective after TBI (97-99). 2-DG is already in use clinically to limit tumor growth in cancer patients (based on the Warburg effect, which describes increased glycolytic activity in neoplastic cells), and short-term treatments are well-tolerated in humans (75). This underscores that short-term use of 2-DG as a disease-modifying agent is a feasible clinical strategy.

A particularly important finding of our study is the cell type-specific coupling of metabolism to neuronal excitability. Our results are concordant with published studies showing that GABAergic interneurons have unique energetic requirements and are known to be enriched with mitochondria for energy production via oxidative phosphorylation (reviewed in (100)). Our findings suggest that glutamatergic pyramidal neurons rely on glycolysis to maintain their intrinsic excitability. While glycolysis is traditionally thought to occur mainly in astrocytes (astrocyte-neuron lactate shuttle model (101)), there is significant evidence that glycolysis also occurs in neurons. Multiple glycolytic enzymes are enriched in neurons relative to astrocytes (102, 103) and

recent work implicates increased neuronal glycolysis in conditions of heightened energy need (104, 105). Thus, increased energetic requirements, glycolytic flux, and cell type-specific neuronal activity may all be interdependent during the acute period following TBI, and deserve further investigation. In future studies, we hope to explore the molecular mechanisms of decreased excitability in excitatory neurons in the presence of 2-DG. We found that decreased excitability in excitatory neurons was associated with significantly decreased membrane resistance, suggesting that the opening of specific ion channels may underly the change in excitability. A possible candidate for this mechanism is an ATP-dependent potassium channel, as K_{ATP} channels have been previously shown to link neuronal energy status with excitability (106-108) and have been implicated in network hyperexcitability and epilepsy (43, 109). A deeper understanding of metabolic demands following TBI and how different CNS cells utilize energy will aid in the development of metabolic approaches to manipulate neural network activity with greater precision.

In summary, we report that in vitro and in vivo 2-DG treatments attenuate multiple pathologic cellular and network changes following CCI in mice. Future work is needed to investigate the use of 2-DG to prevent injury pathology in other settings, including in diffuse or concussive mechanisms of TBI, in larger animal models, and in pilot clinical trials. Additionally, studies are needed to explore whether reducing cortical hyperexcitability is sufficient to ameliorate patient-oriented outcomes including mortality, seizure frequency/severity in post-traumatic epilepsy, and cognitive/behavioral outcomes. Finally, improving our understanding of the specific energy demands of the neuronal activity, hemodynamic response, and inflammation that occur after TBI will enable us to more effectively harness metabolic manipulations to reduce brain injury-associated pathologies.

Materials and Methods

Study Design

Basic Design: All experiments were hypothesis-driven, rigorously designed, and performed in a controlled laboratory environment. Sample size was set based on power analysis calculations using GPower to provide power >0.80 for $\alpha = 0.05$ for most experiments. We also utilized standards in the field, i.e. ≥ 3 animals per group, to ensure proper statistical power.

Randomization and blinding: Mice were randomized to groups within each experiment (sham vs. CCI; vehicle vs. 2-DG). All studies were blinded at the level of data analysis. Data was numerically encoded and analyzed by a trained experimenter blinded to condition. To further prevent bias during analysis, automated analysis approaches were utilized for all synaptic data sets (Figures 2, 5). After initial analysis was finalized, the blinding was lifted to compare data across groups. Because CCI lesions are easily discernable from sham and in vitro drug applications cannot be reasonably blinded, blinding during data collection of electrophysiological experiments was not feasible (Figures 1-3). Due to constraints of animal tracking, blinding was not performed during in vivo drug injections (Figures 4-6). Immunohistochemical and anatomical experiments were blinded throughout the data collection process (Figure 6).

Exclusion criteria and statistical approach: Each experiment had one primary endpoint based on the experimental design. We did not define rules for stopping data collection outside of humane endpoints based on animal protocols. Data was excluded based on quality control of whole-cell recordings (recordings were excluded when access resistance changed $>20\%$ during an experiment). Outliers were removed in the context of fEPSP recordings due to inherent variability in the responses evoked with this technique (Figures 3-4). Outliers were defined as slices with

response area >2 standard deviations from the group mean and were excluded from the subsequent analysis. The number of experimental replicates is included in the figure captions and in Table S1. The number of individual animals, cells, or slices were factored into our analysis using a linear mixed model approach. This approach includes terms to account for inter- and intra-animal variability.

Animals

All animal procedures were performed in accordance with the Tufts University School of Medicine's Institutional Animal Care and Use Committee. All experiments were performed on adult mice. C57/BL6 were obtained from Jackson Labs (Strain #000664), Charles River Laboratories (Strain #027), or bred in-house. *PV^{Cre}* (Stock #017320), *Ai9* (Stock #07909), and *G42* (Stock #007677) mice were obtained from Jackson Laboratories. Animals were kept on a standard 12-hour light cycle and fed *ad libitum* with regular chow diet and water. Male mice were used for all experiments, unless noted.

Controlled cortical impact

Traumatic brain injury was modeled with controlled cortical impact (CCI), as previously described (20, 110). Briefly, 10-14-week-old male mice were anesthetized using inhaled isoflurane in oxygen (4% for induction, 2% for maintenance). Following placement in a Kopf stereotaxic frame, the surgical area was sterilized and a vertical, midline skin incision was made. A 5 mm craniectomy was performed lateral to midline, between bregma and lambda (over the left somatosensory cortex) and the skull flap was removed. The surgical field was flushed with sterile saline throughout the procedure to cool the surgical area during the craniectomy drilling. Impact was performed with a Leica Benchmark Stereotaxic Impactor, using a 3 mm diameter piston, 3.5

m/s velocity, 400 ms dwell time, and 1 mm depth. After the CCI procedure, sutures were used to close the incision. The bone flap was not replaced in order to accommodate swelling after the procedure and to prevent pressure-induced damage to the injury site. Sham animals received the anesthesia and craniotomy drilling, but did not receive the CCI. Animals were singly housed following surgery until time of sacrifice.

In vivo 2-DG treatment

Animals were injected intraperitoneally (I.P.) with 250 mg/kg 2-DG dissolved in sterile saline to a final injection volume of 200 μ L per dose. Vehicle injections consisted of 200 μ L of sterile saline per dose. For the in vivo 2-DG treatment paradigm, the first dose was administered ~20 minutes after CCI or sham surgery, and the doses were continued daily for 7 days at the same time of day. Animal weight was recorded daily to monitor animal health.

Acute brain slice preparation

Brain slices were prepared as previously described (20, 111). Briefly, mice were anesthetized in an isoflurane chamber and then decapitated by guillotine. Brains were rapidly removed and placed in chilled slicing solution (in mM: 234 sucrose, 11 glucose, 24 NaHCO₃, 2.5 KCl, 1.25 NaH₂PO₄, 10 MgSO₄, 0.5 CaCl₂) equilibrated with 95% O₂/5% CO₂. The brain was then glued to the slicing stage of a Leica VT1200S vibratome and the slicing chamber was filled with chilled slicing solution, again equilibrated with 95% O₂/5% CO₂. Coronal slices (300 or 400 μ m-thick) were taken at 0.05 mm/s and hemisected. The slices were placed in a chamber filled with artificial cerebrospinal fluid (aCSF, in mM: 126 NaCl, 26 NaHCO₃, 3 KCl, 1.25 NaH₂PO₄, 2 MgCl₂, 2 CaCl₂, 10 glucose) equilibrated with 95% O₂/5% CO₂. The chamber with the acute

cortical slices was incubated in a water bath at 34°C for one hour, and then stored at room temperature until recording.

In vitro treatment of acute cortical slices with 2-DG

For 2-DG wash-on experiments, 2-DG-aCSF solution was prepared by adding 8 mM 2-DG and removing 8 mM glucose from the standard aCSF recipe (to maintain solution osmolarity, for a final concentration of 2 mM glucose, 8 mM 2-DG). Baseline field or whole-cell recordings were obtained, and then the regular aCSF perfusate was replaced with 2-DG-aCSF for the remaining recordings. For low glucose wash-on experiments, aCSF containing 2 mM glucose and 8 mM sucrose (an inert sugar) was prepared (LG-aCSF), and LG-aCSF was washed on following baseline field or whole-cell recording collection.

Whole-cell recordings

Acute cortical slices were placed in a submersion recording chamber of an Olympus BX51 microscope with continuous perfusion of oxygenated aCSF at 32°C. Layer V pyramidal neurons were visually identified with infrared differential interference contrast microscopy, and fast-spiking parvalbumin-positive cortical interneurons were identified based on GFP expression in *G42* mice (Jax Stock #007677). Whole-cell recording mode was achieved with a borosilicate glass electrode (3-5 M Ω) filled with internal solution optimized for each experiment (described below). Access resistance, membrane resistance, and capacitance were monitored throughout each experiment. Cells with >20% change in access resistance during the experiment were excluded from analysis. Data were collected using an Axon Multiclamp 700B amplifier, Digidata 1440A digitizer, and pClamp software.

For recordings of synaptic activity, the internal utilized was, in mM: 140 CsMs, 10 HEPES, 5 NaCl, 0.2 EGTA, 5 QX314, 1.8 MgATP, 0.3 NaGTP, pH = 7.25, mOsm = 290. The recording electrode was placed within 200 μ m of the lateral edge of the cavitory cortical lesion. Spontaneous EPSCs and IPSCs were recorded by voltage-clamping the cell at -70 and 0 mV, respectively, for 2 minutes. Synaptic activity analysis was performed using MiniAnalysis (SynaptoSoft) with the experimenter blinded to experimental group.

For current injection experiments, the internal utilized was, in mM: 120 KGluconate, 11 KCl, 10 HEPES, 10 EGTA, 2 MgCl₂, 2 MgATP, 0.3 NaGTP, pH = 7.3, mOsm = 290. The perfused aCSF also included DNQX (20 μ M), CPP (10 μ M), and gabazine (10 μ M) to block synaptic activity. Current injection steps were applied to each cell (ranging from -200 \rightarrow 400 pA in 25 pA steps) and cell type was confirmed using AP shape and firing rate.

Field recordings

Brain slices were placed in an interface chamber perfused with 34°C, oxygenated aCSF at a rate of 2 mL/min. A tungsten stimulating electrode was placed at the layer VI-white matter boundary to stimulate ascending white matter tracts. A borosilicate glass micropipette (pulled to a resistance of \sim 1 M Ω) was filled with aCSF and placed in the corresponding area of layer V cortex directly above the stimulating electrode. Recordings were performed in the perilesional cortex (within 100 μ m of the injury site) or in isotopic cortex in sham animals. Electrical stimulus (8-25 μ A, 100 μ s pulse length) was delivered using a stimulus isolator (World Precision Instruments) at 30 second intervals. The signal was amplified with a Multiclamp 700A amplifier, digitized with a Digidata 1322A digitizer (sampling rate = 20 kHz), and recorded with pClamp software (Molecular Devices). Threshold stimulus intensity was defined as the minimal required stimulus

to obtain a detectable field response (≥ 0.05 mV). “2X threshold” was defined by doubling the threshold stimulus intensity for a given slice.

Analysis of field recordings

Evoked field potentials were analyzed using pClamp software and MATLAB. First, traces were adjusted by subtracting the baseline (the average amplitude directly before the stimulus), imposing a low-pass filter (Bessel 8-pole, 1000 Hz cutoff), and filtering for electrical 60 Hz noise. The area of the field potential was measured by integrating the charge transfer in the first 250 ms following the stimulus. Coastline measurements describe the relative amount of high-frequency activity in the field response and were measured by summing the distance between each point taken every 50 μ s over a 250 ms time window following the stimulus. The percentage of epileptiform sweeps for a given slice was determined by counting the number of field potentials that exhibited epileptiform activity (prolonged, high frequency, and high amplitude) out of 10 total sweeps (over a 5 minute period).

Immunohistochemistry

Animals were transcardially perfused with 100 mL of phosphate-buffered saline (PBS) followed by 100 mL of 4% paraformaldehyde (PFA) in 0.4 M phosphate buffer (PB). Brains were dissected out, placed in 4% PFA in PB overnight, and then moved to a 30% sucrose solution for 3 days. Forty μ m thick coronal slices were taken on a Thermo Fisher Microm HM 525 cryostat. For CCI or sham animals, the brains were serial sectioned, so that each collection well contained a representative set of coronal sections throughout the entire CCI lesion, with each section 400 μ m apart from the next. Slices were washed with PBS and then incubated in blocking buffer (10% normal goat serum (NGS), 5% bovine serum albumin (BSA) in PBS with 0.2% Triton X-100

(PBS-T)) at room temperature for 1 hour. The slices were incubated overnight at 4°C with primary antibody in PBS-T with 5% NGS/1% BSA. The primary antibodies used were as follows: parvalbumin (PV; mouse; Swant PV235; 1:1000 for Figure 6, 1:2000 for Figure S1), Green Fluorescent Protein (GFP; chicken; Abcam Ab13970; 1:1000), Glutamate decarboxylase 67 (GAD67; mouse; Millipore MAB5406; 1:10000), Somatostatin (SST; rat; Millipore MAB354; 1:100), and Calretinin (CR; mouse; Swant 6B3; 1:5000). Then slices were washed with PBS and incubated with secondary antibody (anti- mouse, rat, or chicken antibodies tagged with Alexa 488, Cy3, or 647, applied at 1:500) in PBS-T with 5% NGS/1% BSA for 2 hours at room temperature. Finally, slices were rinsed with PBS and mounted, in anatomical order, onto slides using Vectashield with DAPI.

Imaging for cell counting was performed using a Keyence epifluorescence microscope with a 10x or 20x optical zoom. Additional images for Figure 6 were taken on a Leica SPE confocal microscope with a 63x oil objective; additional images for Figure S1 were taken on a Nikon A1R confocal microscope with a 40x oil objective (Olympus). For each experiment, identical laser and capture parameters were utilized across all experimental groups.

Analysis of imaging data

Image analysis was performed using Fiji/ImageJ software. For Figure 6, 5 regions of interest (ROIs) were drawn lateral to the edge of the cavitory CCI lesion, with each ROI 100 μ m wide. PV+ and tdT+ cells were counted independently, by an experimenter blinded to treatment group. The cell counts were then divided by the area of the selected region to find the cell density. The number of PV+tdT+ colocalized cells was determined by comparing the locations of cell bodies selected in the individual PV+ and tdT+ counting phases. Three sections from each brain

(representing the same anatomical regions across all subjects) were quantified and averaged to obtain a single cell density value per cell type in each ROI in each animal. For Figure S1, GFP+ cells were identified manually in layer V-VI cortex of sections from *G42* mice, and then examined for their co-labeling with PV, GAD67, SST, or CR immunolabels.

Lesion volume was evaluated in >10 serial sections from each animal, mounted and imaged in anatomical order. A blinded experimenter traced the size of the cavitory lesion in each section and the area of missing tissue was recorded. Each area was multiplied by 400 μm (the distance between the serial sections) to estimate the lesion volume.

Metabolic studies

Blood glucose was measured using a OneTouch glucose meter and blood β -hydroxybutyrate was measured using a PrecisionXtra Ketone Monitor with blood taken from the tail. Flank temperature was measured using a subcutaneous temperature probe (BioMedic Data Systems Implantable Electronic ID Transponders IPTT-300). Food intake was measured by subtracting the final weight from the initial weight of the chow during the indicated study period. Home cage locomotor activity was measured utilizing a photobeam-break system (Tufts Behavior Core). Animals were fasted for 16-18 hours before acute measurements following 2-DG injection (Figure 4I-J, 4L).

Statistical analysis

Most experiments used a 2X2 factorial design. To assay the statistical significance of each factor (fixed effect) and the interaction between factors, we performed linear mixed model (LMM) analysis (see Table S1 and Supplementary Methods, (112)). For comparison of cumulative distributions of inter-event interval or amplitude of synaptic events, we utilized a 2-sample

Kolmogorov-Smirnov test with α level corrected for multiple comparisons. For analyzing home cage locomotor activity and lesion volume, we used a 2-sample Student's t-test (2-tailed) with $\alpha = 0.05$.

Study approval

All animal procedures were performed in accordance with (and approved by) the Tufts University School of Medicine's Institutional Animal Care and Use Committee. No human subjects or tissues were used in this study.

Drugs and reagents

Salts were obtained from Sigma-Aldrich or Fisher Scientific. Glucose was obtained from Fisher Scientific. 2-deoxy-D-glucose and DNQX were obtained from Sigma-Aldrich. CPP was obtained from Abcam. Gabazine was obtained from Tocris.

Author Contributions

CD and JK conceived of and designed experiments examining 2-DG's effects on neuronal excitability and synaptic activity. CD, DC, and JK conceived of and designed experiments examining 2-DG's effects on epileptiform activity and interneuron density. CD, JK, CL, and DK conceived of and designed experiments examining effects of 2-DG on metabolism. JK performed experiments and analyzed the data for Figures 1, 2, 5, 6, S2, and S3. DC performed experiments for Figures 3 and 4A-H; DC and JK analyzed the data for Figures 3 and 4A-H. JK and CL performed metabolic experiments and analyzed the data for Figures 4I-L and S4. MS performed experiments and analyzed the data for Figure S1. FN performed all LMM statistical analysis (Tables S1-S2). JK and CD wrote first draft of manuscript; comments from all authors were incorporated into the final draft.

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Competing interests

There are no competing interests to declare.

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Figures

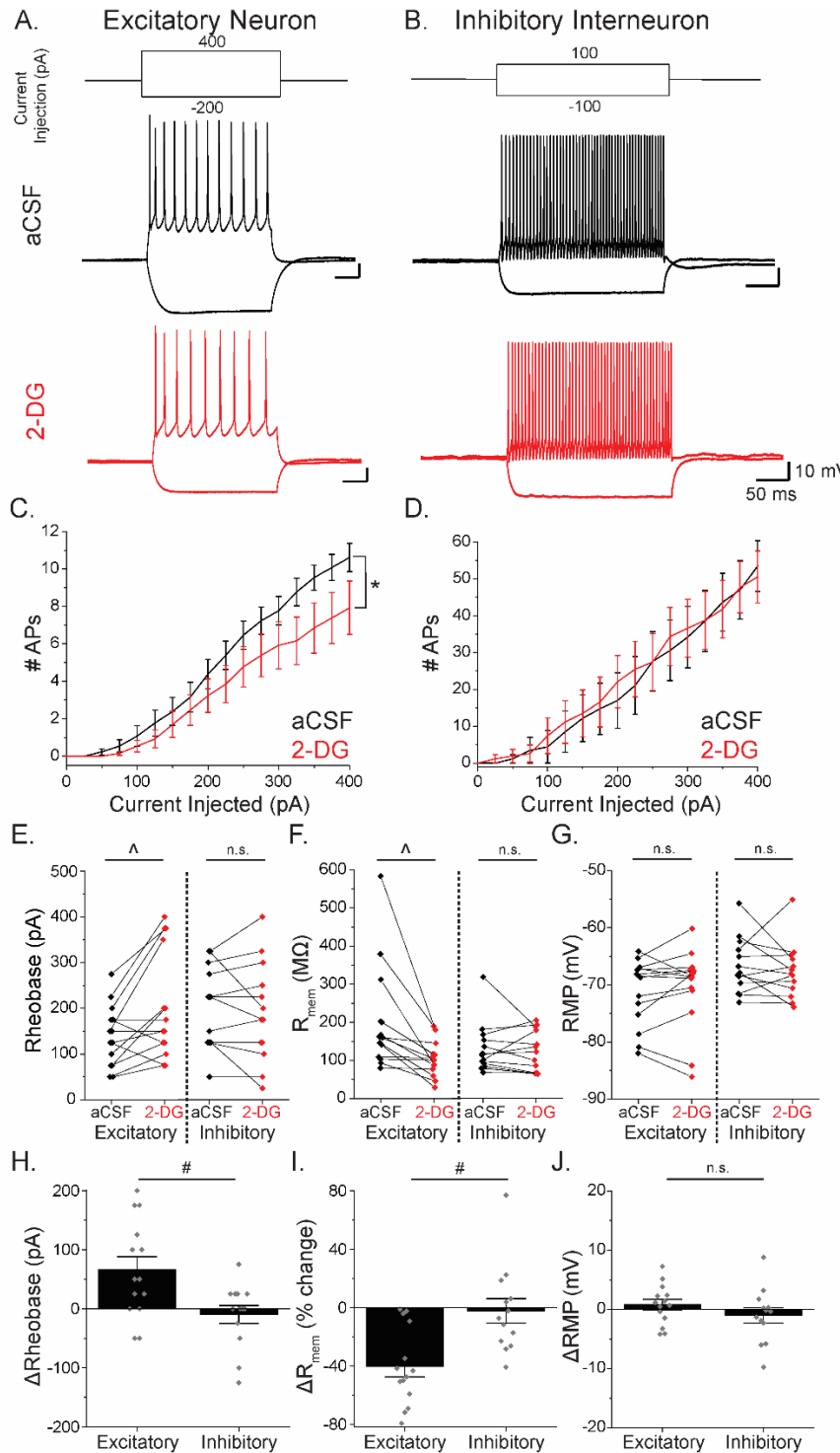


Figure 1: In vitro 2-DG treatment decreases the intrinsic excitability of excitatory pyramidal neurons. A-B. Representative traces following current injection into layer V cortical excitatory

pyramidal neurons (**A**) or interneurons (**B**) before (black) or after (red) treating the cortical slice with 2-DG for 10 minutes. **C-D**. Input-output curves from excitatory pyramidal neurons (**C**) and inhibitory interneurons (**D**). **E-G**. Rheobase (**E**, current injection required to fire the first action potential), membrane resistance (**F**), and resting membrane potential (**G**, RMP) before (black) and after (red) 2-DG treatment in each cell type. **H**. Δ Rheobase (rheobase in 2-DG versus baseline) in excitatory neurons and interneurons. **I**. % change in membrane resistance of excitatory neurons and interneurons following 2-DG treatment. **J**. Change in resting membrane potential (RMP) in excitatory neurons and interneurons in 2-DG. (Error bar = SEM. n=14 excitatory neurons from 9 animals, 13 inhibitory interneurons from 10 animals. LMM: * $t > \pm 1.96$, effect: interaction between current injection and 2-DG; # $t > \pm 1.96$, effect: cell type. 2-sample t-test: $p < 0.05$. n.s. not significant.)

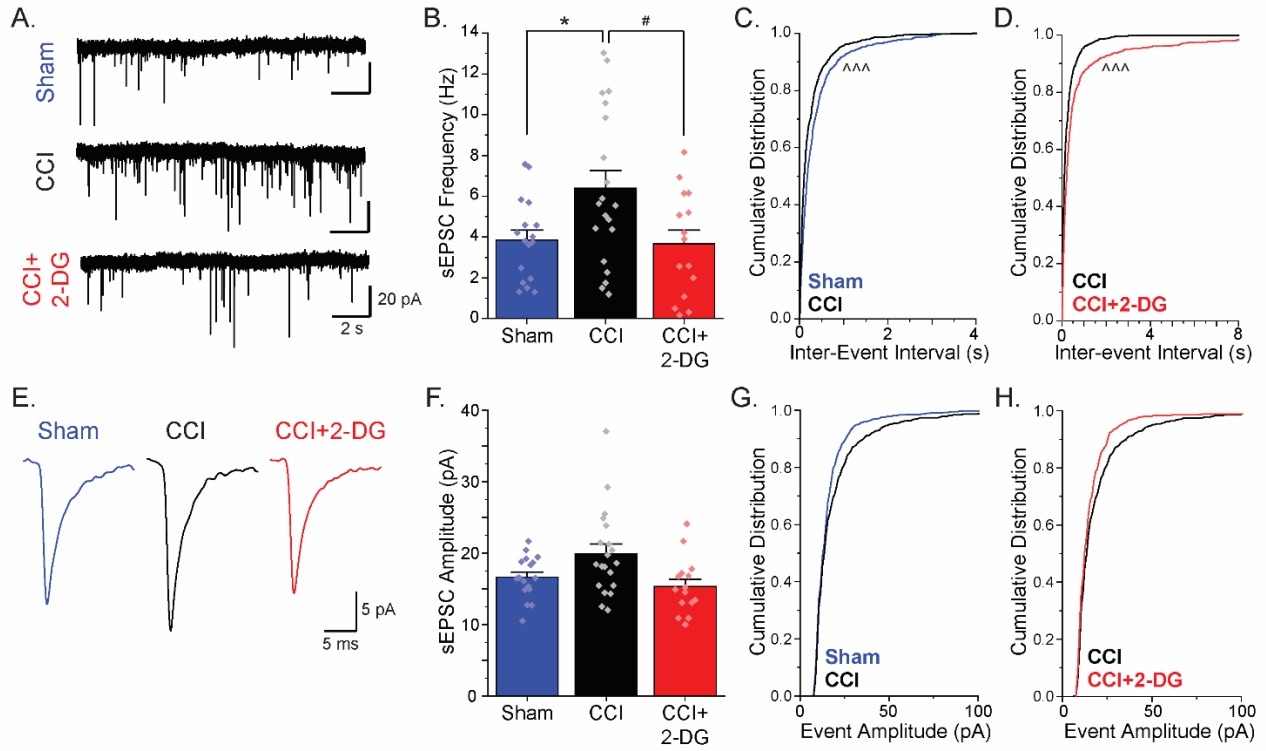


Figure 2: In vitro 2-DG attenuates CCI-induced increases of excitatory synaptic activity onto interneurons 3 days after injury. **A.** Representative sEPSC traces from acute cortical slices from sham or CCI-injured animals, and a representative trace from a CCI slice following 10 minutes of 2-DG wash-on. **B.** Mean sEPSC frequencies. **C-D.** Cumulative distributions of inter-event interval generated from 100 random events per recorded cell. **E.** Representative events from each condition, generated by averaging 100 events from a representative trace. **F.** Average event amplitude. **G-H.** Cumulative distributions of event amplitude generated from 100 randomly selected events per cell. (Error bar = SEM. n=15-20 cells from 4 animals/group. LMM: * $t > \pm 1.96$, effect: CCI; # $t > \pm 1.96$, effect: interaction between CCI and 2-DG. 2-sample K-S test with correction for multiple comparisons: *** $p < 1E-5$)

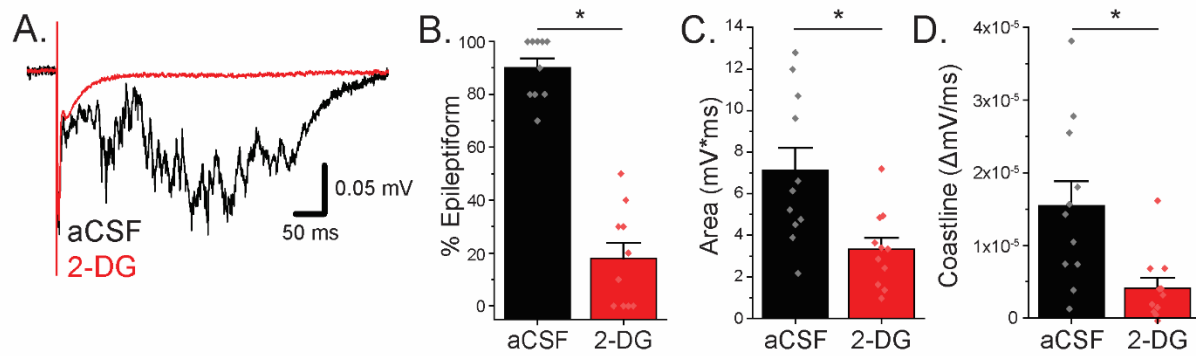


Figure 3: Acute 2-DG treatment decreases epileptiform activity in vitro following CCI. A. Representative stimulus-evoked field potentials in an acute cortical slice from a CCI-injured animal before (black) and after (red) local perfusion of 2-DG for 30 minutes. **B.** Percentage of traces with epileptiform activity after CCI. **C-D.** Area (C) and coastline (D) of the field potentials in baseline aCSF and 2-DG-aCSF in CCI-injured slices. (Error bar = SEM. n=10 slices from 6 animals. LMM: $*t > \pm 1.96$, effect: interaction between CCI and 2-DG.)

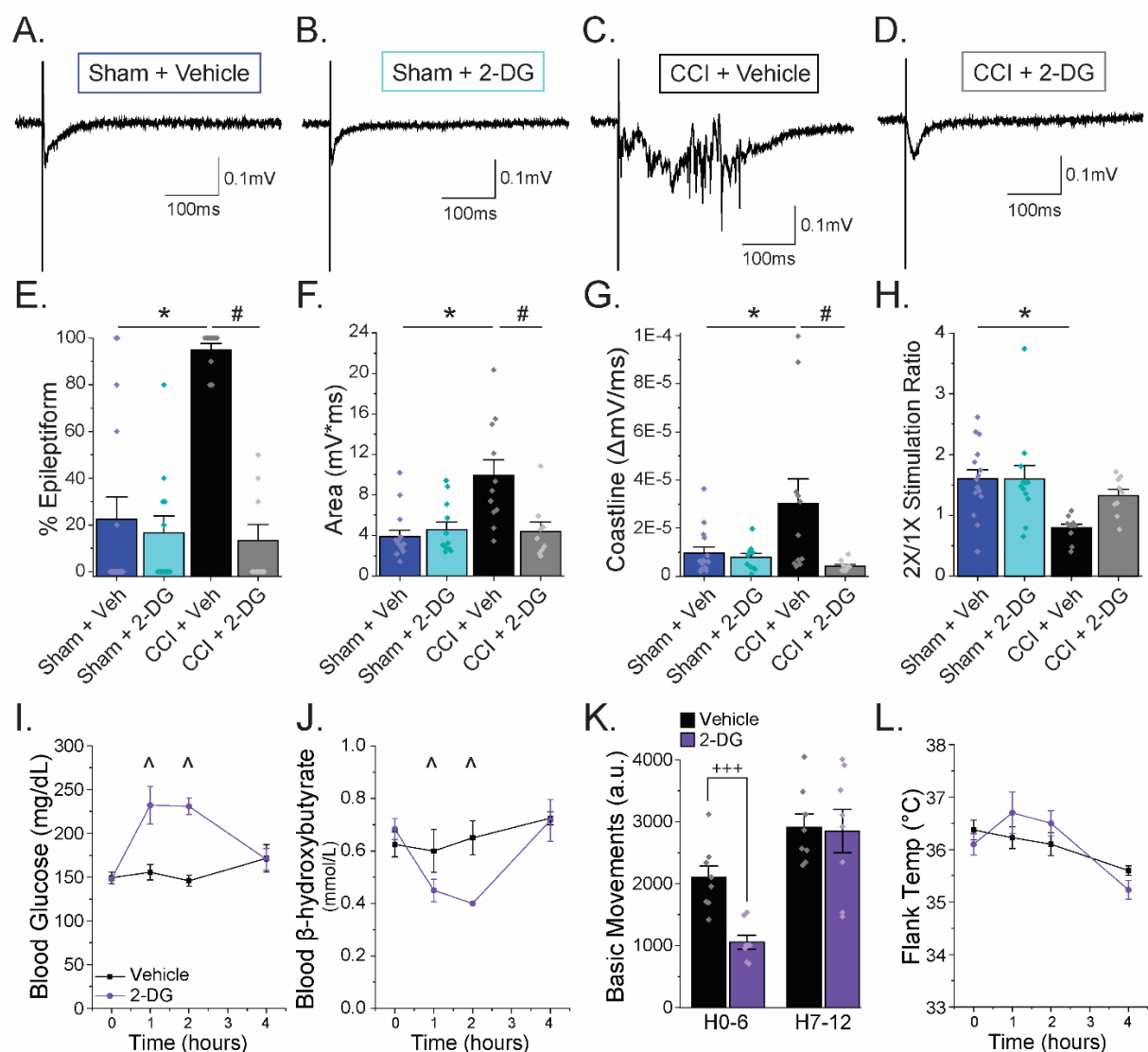


Figure 4: In vivo 2-DG treatment following CCI decreases epileptiform activity in acute cortical slices. Animals were given sham or CCI surgery, followed by 1 week of daily intraperitoneal injections of vehicle or 2-DG (250 mg/kg). **A-D.** Representative stimulus-evoked field potentials taken from acute cortical slices 3-5 weeks following sham or CCI surgery, with vehicle or 2-DG treatment. **E.** Percentage of sweeps per slice exhibiting epileptiform activity. **F-G.** Area (F) and coastline (G) of fEPSP traces. **H.** Input-output ratios. **I.** Blood glucose in naïve fasted mice after a single vehicle (black) or 2-DG (purple) injection. **J.** Blood ketone body levels (as measured by β -hydroxybutyrate) after vehicle or 2-DG injection. **K.** Basic locomotor activity in vehicle- or 2-DG-treated animals. **L.** Flank temperature after vehicle or 2-DG injection. (Error bar = SEM. $n=9-15$ slices from 3 animals/group for fEPSPs. $n=4$ vehicle, 6 2-DG animals for metabolism studies. $n=8$ animals/group for locomotor activity. LMM: * $t>\pm 1.96$, effect: CCI; # $t>\pm 1.96$, effect: interaction between CCI and 2-DG; ^ $t>\pm 1.96$, effect: 2-DG at each time point. 2-sample t-test: +++ $p<0.001$)

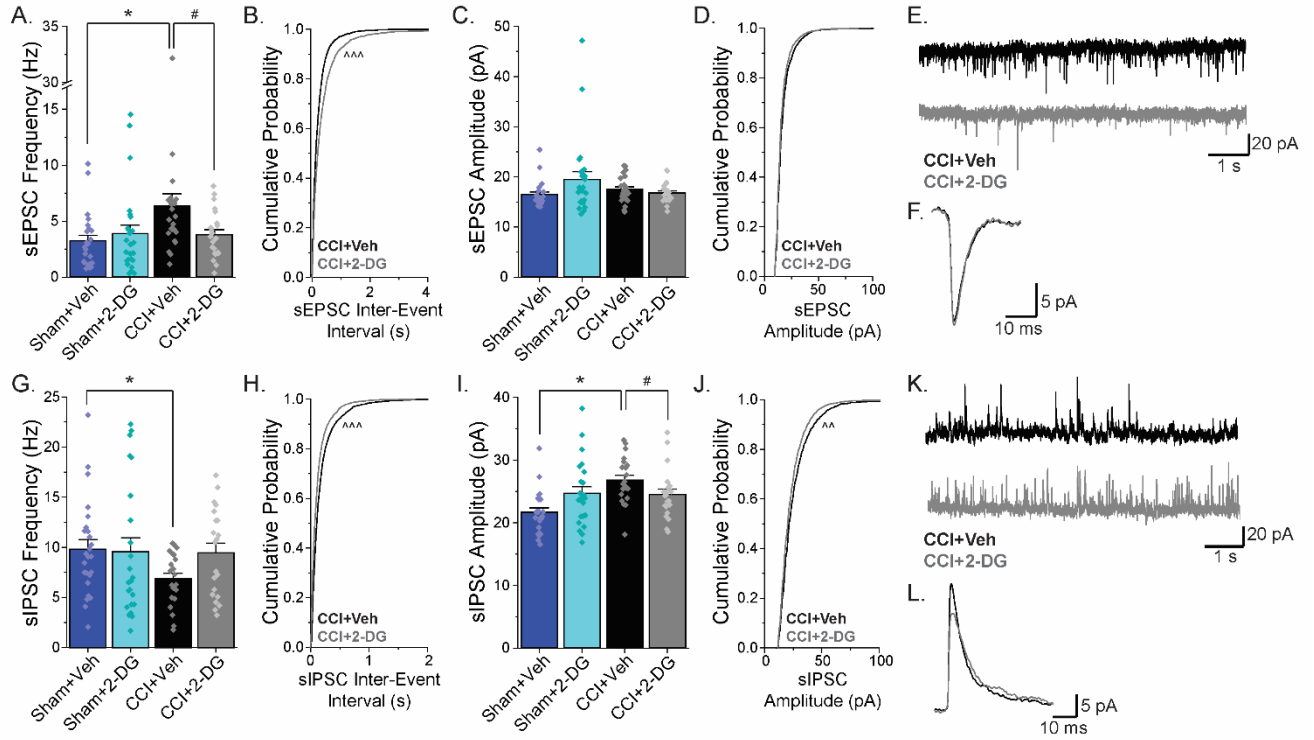


Figure 5: In vivo 2-DG treatment attenuates changes in synaptic activity onto perilesional layer V pyramidal neurons 3-5 weeks after injury. A. Mean sEPSC frequency. B. Cumulative distribution of sEPSC inter-event interval. C. Mean sEPSC amplitude. D. sEPSC amplitude cumulative distribution. E. Representative sEPSC traces from vehicle-treated or 2-DG-treated CCI brains. F. Representative example of sEPSC events (generated by averaging 100 events from a representative trace). G. Mean sIPSC frequency. H. Cumulative distribution of sIPSC inter-event interval. I. Mean sIPSC amplitude. J. Cumulative distribution of sIPSC amplitude. K. Representative sIPSC traces from vehicle-treated or 2-DG-treated CCI brains. L. Representative example of sIPSC events. (Error bar = SEM. $n=21-26$ cells from 5 animals/group. LMM: * $t > \pm 1.96$, effect: CCI; # $t > \pm 1.96$, effect: interaction between CCI and 2-DG. 2-sample K-S test corrected for multiple comparisons: $^{AA}p < 1E-4$; $^{AAA}p < 1E-5$)

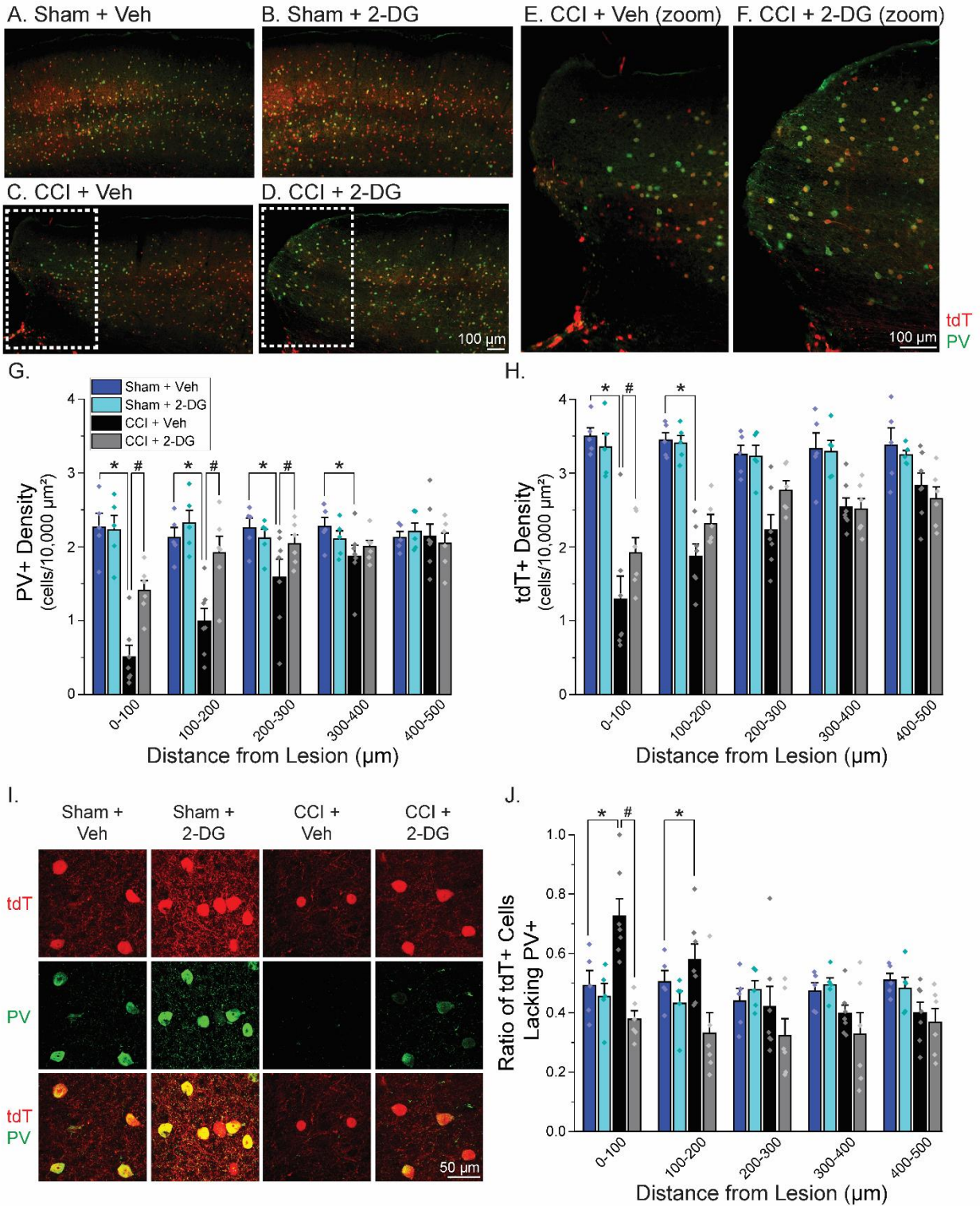


Figure 6: In vivo 2-DG treatment attenuates the decrease in PV+ cells in the perilesional cortex. A-D. Representative images with PV (green) and tdT (red – in PV^{Cre} cells) in animals from the different treatment groups, collected 3-5 weeks after sham or CCI surgery. E-F. Zoom-in of perilesional area showing loss of colocalized perilesional PV+tdT+ cells after CCI. G.

Quantification of PV⁺ cell density in five regions lateral to the CCI cavitory lesion. **H.** Quantification of cells identified with tdT⁺ in *PV^{Cre}/Ai9* reporter mice. **I.** Example confocal images of cells in the perilesional area with genetic tdT expression (red) and PV immunolabel (green). **J.** The density of tdT⁺ cells not colocalized with PV immunolabel was divided by the total density of tdT⁺ cells in each ROI to calculate the ratio reported. (Error bar = SEM. n=5-7 animals/group (3 sections from each animal were analyzed and averaged to generate a single value for each animal). LMM: * $t > \pm 1.96$, effect: CCI in each ROI; # $t > \pm 1.96$, effect: interaction of CCI and 2-DG in each ROI.)

Supplementary Methods

Linear mixed model analysis

For most of the statistical analysis, we utilized linear mixed model (LMM) analysis. This approach estimates the effect size of each factor while accounting for intra- and inter-animal variability. LMMs were fitted with random intercepts to assess for the correlation between repeated measurements on the same mouse, and experiment-specific effects were analyzed for statistical significance. t values > 1.96 and < -1.96 were considered to be statistically significant and corresponded to 95% confidence intervals that did not cross zero. Each LMM examined both main fixed effects and interactions between the effects.

Experiments with a traditional 2 X 2 factorial design (including those for Figures 3-5) used LMM to examine fixed effects of CCI, 2-DG treatment, and the interaction between these two effects. To compare input-output curves in Figure 1, the fixed effects were current injection, 2-DG treatment, and the interaction between the two effects. LMM was also utilized to compare excitatory and inhibitory neurons in Figure 1, by examining cell type as the fixed effect.

In Figure 2, the statistical approach also paralleled the experimental design. The primary experimental question was whether synaptic activity onto interneurons was altered acutely after CCI (as this had not been shown previously); thus “Stage 1” of the statistical approach was to examine CCI as the solitary fixed effect. Then, in “Stage 2” of the experiment and statistical analysis, we examined the effects of in vitro 2-DG wash-on to slices taken from either CCI or sham animals (and thus used both CCI and 2-DG as fixed effects).

In Figure 6, there was another fixed effect (ROI), introduced as a categorical variable. We performed LMM with fixed effects of CCI, 2-DG, ROI, and interactions between each of these.

Comparisons were made with ROI5 (the furthest region from the CCI lesion) as the reference point. To further interpret these complex results, we utilized LMM with the Type III analysis of variance (ANOVA) test with Satterthwaite's method to assess the global significance of "ROI" as a single factor instead of each ROI as an independent factor relative to ROI5. Thus, the Type III ANOVA reports global effects across all ROIs (Table S2).

Cumulative distribution generation

Cumulative distributions were generated by randomly selecting 100 events from each recording to ensure that data from more active cells were not more heavily weighted than data from cells with fewer events. Random event selection was accomplished using a custom-written MATLAB script. Within each treatment group, randomly selected events from each cell were pooled to generate a single cumulative distribution, and distributions were compared using a 2-sample Kolmogorov-Smirnov (K-S) test. To account for the large degrees of freedom associated with comparing distributions and to prevent false positives, we decreased α to 0.001. We further corrected for repeated comparisons when doing multiple 2-sample K-S tests on the same data set by dividing α by 6 possible comparisons across the four treatment groups (final $\alpha = 1.67\text{E-}4$).

Supplementary Figures

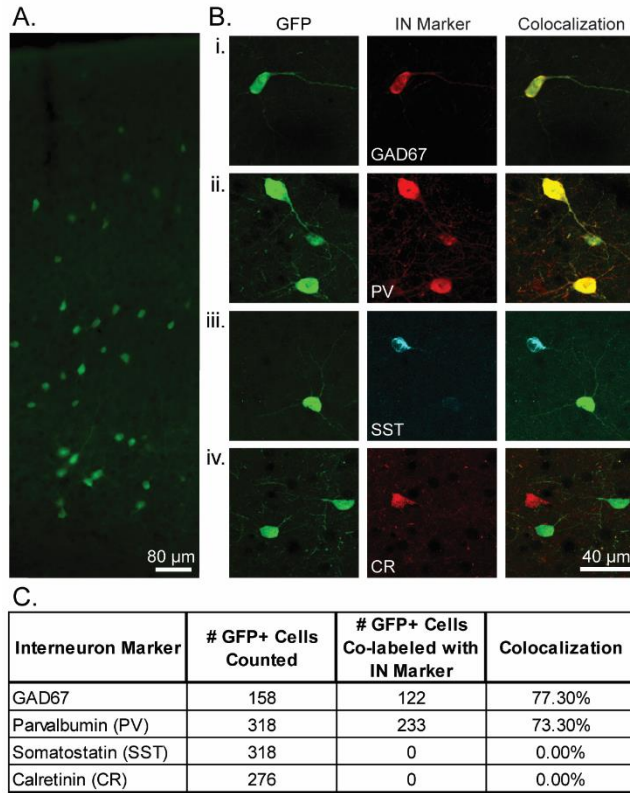


Figure S1: Characterization of *G42* mouse line. **A.** Immunohistochemical labeling of genetically-encoded GFP (green) in the somatosensory cortex (SSC) of adult *G42* mice. **B.** Co-labeling of GFP in *G42* mice with markers of inhibitory interneurons in layers V-VI of SSC (**i.** GAD67, **ii.** Parvalbumin (PV), **iii.** Somatostatin (SST), **iv.** Calretinin (CR)). **C.** Table showing abundant co-labeling of GFP+ cells with PV and GAD67. (n=4 mice)

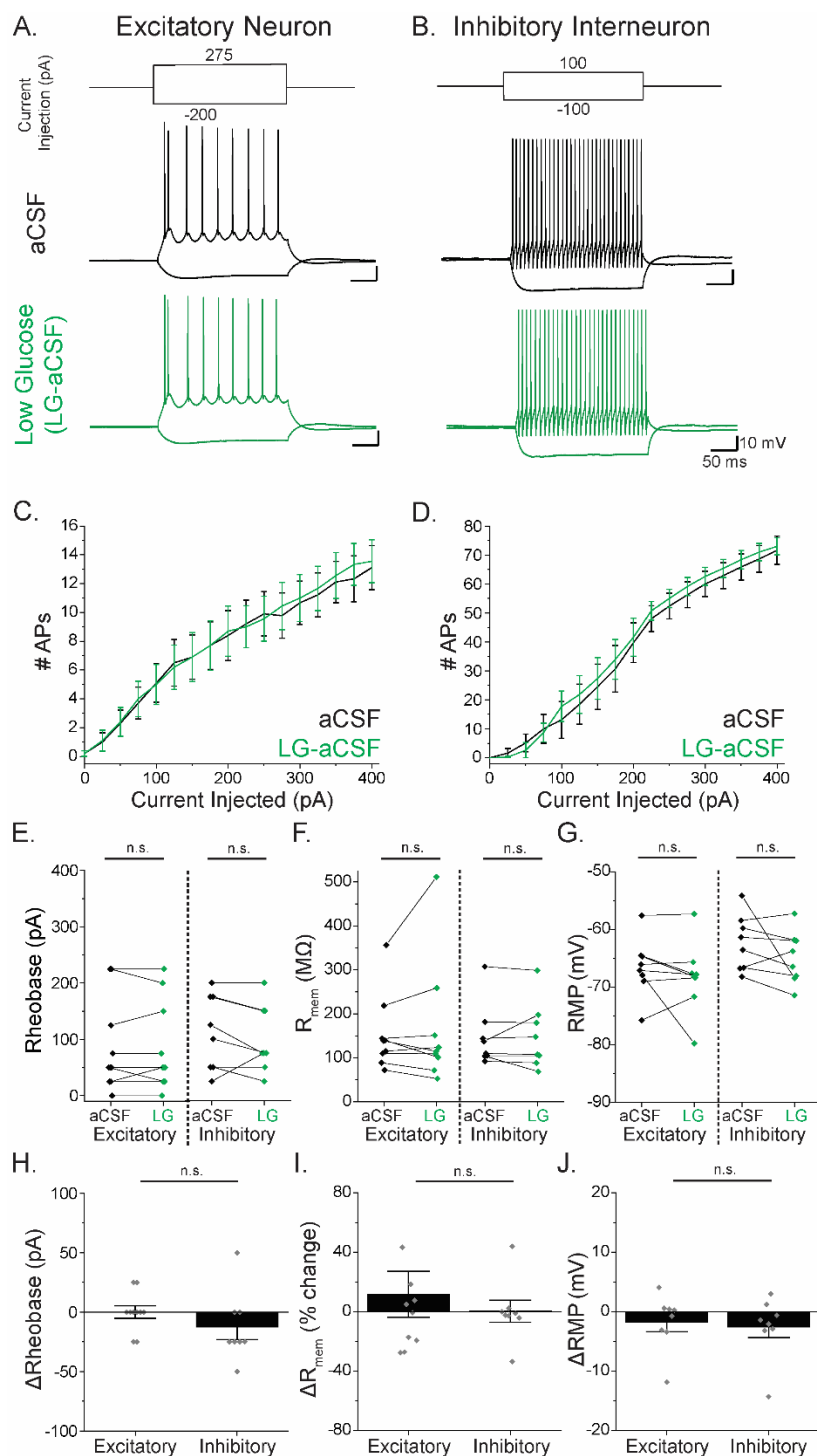


Figure S2: Treatment with Low Glucose (LG)-aCSF did not affect the excitability of excitatory or inhibitory neurons. A-B. Representative traces following current injection into layer V cortical excitatory pyramidal neurons (A) or interneurons (B) before (black) or after (green) treating the cortical slice with LG-aCSF for 10 minutes. C-D. Input-output curves in excitatory (C) or inhibitory (D) neurons. E. Rheobase (current injection required to fire the first action potential) before (black) and after (green) LG-aCSF treatment in each cell type. F.

Membrane resistance before and after LG-aCSF in each cell type. **G.** Resting membrane potential (RMP) before and after LG-aCSF treatment in each cell type. **H.** Δ Rheobase (rheobase in LG-aCSF versus baseline) in excitatory neurons and interneurons. **I.** % change in membrane resistance of excitatory neurons and interneurons following LG-aCSF treatment. **J.** Change in resting membrane potential (RMP) in excitatory neurons and interneurons in LG-aCSF. (Error bar = SEM. n=10 excitatory neurons from 5 animals, 8 inhibitory neurons from 5 animals. n.s. not significant.)

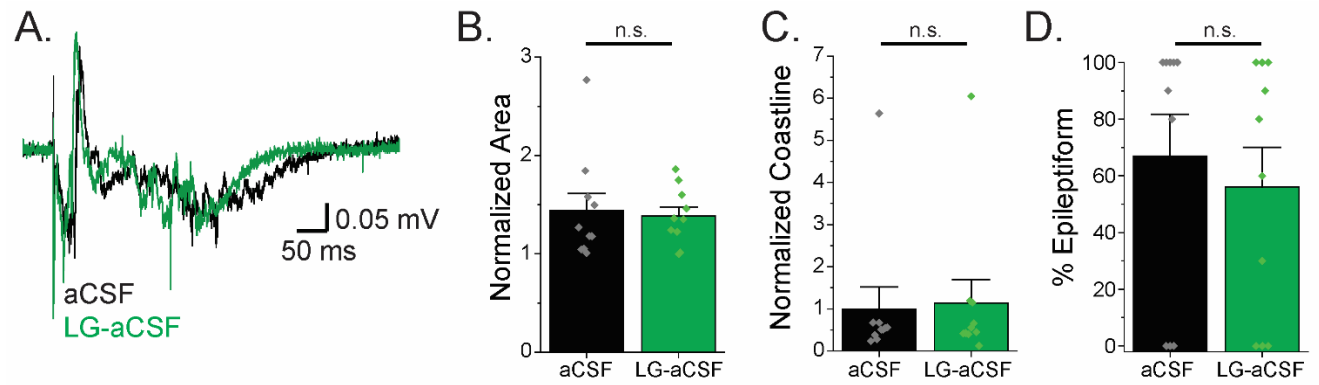


Figure S3: Low glucose (LG) conditions do not attenuate epileptiform activity following CCI.
A. Representative stimulus-evoked field potentials in an acute cortical slice 3-5 weeks following CCI surgery. **B-C.** Area (B) and coastline (C) measurements from fEPSPs from CCI-injured animals with or without 30 minutes of LG-aCSF. **D.** The percentage of sweeps exhibiting epileptiform activity in slices from CCI animals with or without LG-aCSF treatment. (Error bar = SEM. n=10 slices from 4 animals. n.s. not significant.)

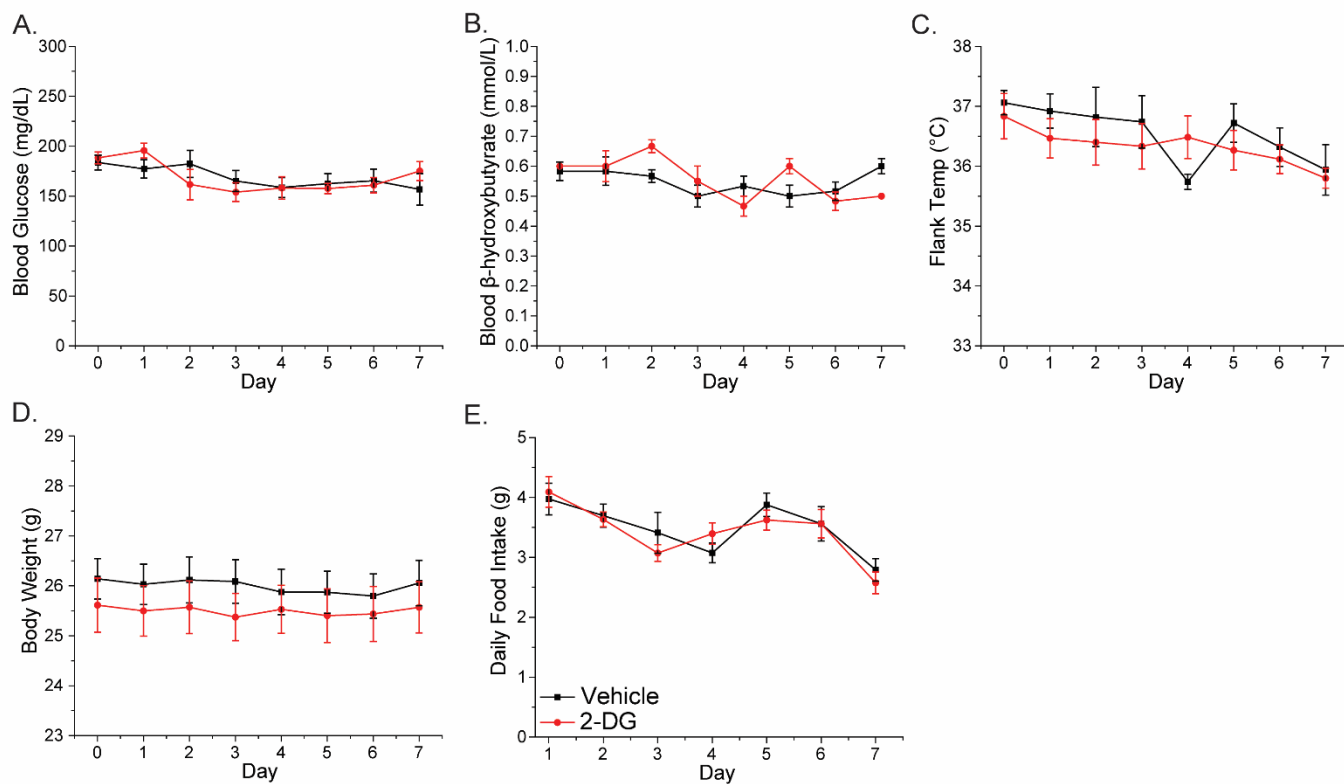


Figure S4: 2-DG has no effect on daily blood glucose, blood β -hydroxybutyrate, temperature, body weight, or food intake during a week-long dosing regimen. Animals were injected daily with 2-DG or vehicle. Shown are blood glucose (A), β -hydroxybutyrate (B), flank temperature (C), body weight (D), and daily food intake (E) measured immediately prior to each daily injection. (Error bar = SEM. n=6 animals/group.)

| Linear Mixed Model Effects | | | | | | | | | | | |
|----------------------------|--|---|-------------|----------|----------|-------------|---|-------------|----------|----------|-------------|
| Figure | N | Fixed effect | Coefficient | SE | t value | Significant | Fixed effect | Coefficient | SE | t value | Significant |
| 1 | Excitatory: n=14 cells from 9 animals Inhibitory: n=13 cells from 10 animals | Excitatory Neuron Input/Output Units: # APs | | | | | Inhibitory Interneuron Input/Output Units: # APs | | | | |
| | | Current Injection | 0.03133 | 0.001336 | 23.451 * | | Current Injection | 0.14475 | 0.008215 | 17.62 * | |
| | | 2-DG (<i>in vitro</i>) | 0.273077 | 0.456739 | 0.598 | | 2-DG (<i>in vitro</i>) | 2.522917 | 2.808498 | 0.898 | |
| | | Interaction Current/2-DG | -0.007801 | 0.001889 | -4.129 * | | Interaction Current/2-DG | -0.005598 | 0.011618 | -0.482 | |
| | | $\Delta R_{heobase}$ (excitatory versus inhibitory) Units: pA | | | | | % ΔR_{mem} (excitatory versus inhibitory) Units: % change | | | | |
| | | Cell Type | 75.687 | 26.982 | 2.805 * | | Cell Type | -38.62% | 11.14% | -3.468 * | |
| | | ΔR_{MP} (excitatory versus inhibitory) Units: mV | | | | | | | | | |
| 2 | Sham+aCSF: n=17 cells Sham+2-DG: n=11 cells CCI+aCSF: n=20 cells CCI+2-DG: n=15 cells 4 animals/group | Frequency - sEPSCs onto INs (PID3) Units: Hz | | | | | Amplitude - sEPSCs onto INs (PID3) Units: pA | | | | |
| | | Stage 1: CCI only | 2.559 | 1.0263 | 2.493 * | | Stage 1: CCI only | 3.144 | 1.773 | 1.773 | |
| | | Stage 2: CCI + 2-DG Effects | | | | | Stage 2: CCI + 2-DG Effects | | | | |
| | | CCI | 2.559 | 1.619 | 1.581 | | CCI | 2.653 | 2.251 | 1.178 | |
| | | 2-DG (<i>in vitro</i>) | 2.387 | 1.272 | 1.877 | | 2-DG (<i>in vitro</i>) | -1.709 | 1.619 | -1.056 | |
| | | Interaction CCI/2-DG | -5.148 | 1.696 | -3.036 * | | Interaction CCI/2-DG | -2.815 | 2.158 | -1.304 | |
| | | | | | | | | | | | |
| 3 | Sham: n=6 slices from 3 animals CCI: n=10 slices from 6 animals | fEPSP % Epileptiform - <i>In vitro</i> 2-DG treatment Units: % sweeps | | | | | fEPSP Area - <i>In vitro</i> 2-DG treatment Units: mV*ms | | | | |
| | | CCI | 90.20% | 7.27% | 12.412 * | | CCI | 3.1432 | 1.4637 | 2.147 * | |
| | | 2-DG (<i>in vitro</i>) | -2.36E-16 | 6.45% | 0 | | 2-DG (<i>in vitro</i>) | 0.8189 | 1.2192 | 0.672 | |
| | | Interaction CCI/2-DG | -72.00% | 8.16% | -8.825 * | | Interaction CCI/2-DG | -4.6187 | 1.5157 | -3.047 * | |
| | | fEPSP Coastline - <i>In vitro</i> 2-DG treatment Units: $\Delta mV/ms$ | | | | | | | | | |
| | | CCI | 1.23E-05 | 4.05E-06 | 3.034 * | | | | | | |
| | | 2-DG (<i>in vitro</i>) | 5.36E-06 | 3.48E-06 | 1.542 | | | | | | |
| 4 | Sham+Veh: n=15 slices Sham+2-DG: n=12 slices CCI+Veh: n=10 slices CCI+2-DG: n=9 slices 3 animals/group | fEPSP % Epileptiform - <i>In vivo</i> 2-DG treatment Units: % sweeps | | | | | fEPSP Area - <i>In vivo</i> 2-DG treatment Units: mV*ms | | | | |
| | | CCI | 67.50% | 12.13% | 5.566 * | | CCI | 6.0403 | 1.3315 | 4.537 * | |
| | | 2-DG (<i>in vivo</i>) | -5.83% | 11.83% | -0.493 | | 2-DG (<i>in vivo</i>) | 0.6882 | 1.2991 | 0.53 | |
| | | Interaction CCI/2-DG | -59.72% | 18.26% | -3.27 * | | Interaction CCI/2-DG | -6.2079 | 1.9901 | -3.119 * | |
| | | fEPSP Coastline - <i>In vivo</i> 2-DG treatment Units: $\Delta mV/ms$ | | | | | fEPSP Input/Output - <i>In vivo</i> 2-DG treatment Units: ratio | | | | |
| | | CCI | 2.05E-05 | 7.58E-06 | 2.701 * | | CCI | -0.81373 | 0.23292 | -3.494 * | |
| | | 2-DG (<i>in vivo</i>) | -1.97E-06 | 7.65E-06 | -0.258 | | 2-DG (<i>in vivo</i>) | -0.01304 | 0.23067 | -0.057 | |
| | | Interaction CCI/2-DG | -2.39E-05 | 1.14E-05 | -2.102 * | | Interaction CCI/2-DG | 0.5443 | 0.34703 | 1.568 | |
| | Vehicle: n=4 animals 2-DG: n=6 animals | Blood Glucose - single 2-DG injection Units: mg/dL | | | | | Blood β -hydroxybutyrate - single 2-DG injection Units: mmol/L | | | | |
| | | Interaction (Hour 1)/2-DG | 78.083 | 26.131 | 2.988 * | | Interaction (Hour 1)/2-DG | -0.20833 | 0.08598 | -2.423 * | |
| | | Interaction (Hour 2)/2-DG | 86.5 | 26.131 | 3.31 * | | Interaction (Hour 2)/2-DG | -0.30833 | 0.08598 | -3.586 * | |
| | | Interaction (Hour 4)/2-DG | 0.25 | 26.131 | 0.01 | | Interaction (Hour 4)/2-DG | -0.06667 | 0.08598 | -0.775 | |
| | | Flank Temperature - single 2-DG injection Units: °C | | | | | | | | | |
| | | Interaction (Hour 1)/2-DG | 0.75 | 0.50827 | 1.476 | | | | | | |
| 5 | sEPSCs: n=22-26 cells/group from 5 animals/group | Frequency - sEPSCs onto pyramidal neurons (PID21-35) Units: Hz | | | | | Amplitude - sEPSCs onto pyramidal neurons (PID21-35) Units: pA | | | | |
| | | CCI | 3.0884 | 1.0952 | 2.82 * | | CCI | 1.151 | 1.551 | 0.742 | |
| | | 2-DG (<i>in vivo</i>) | 0.654 | 1.1057 | 0.591 | | 2-DG (<i>in vivo</i>) | 3.066 | 1.555 | 1.972 * | |
| | | Interaction CCI/2-DG | -3.1991 | 1.5744 | -2.032 * | | Interaction CCI/2-DG | -3.902 | 2.215 | -1.762 | |
| | sIPSCs: n=21-25 cells/group from 5 animals/group | Frequency - sIPSCs onto pyramidal neurons (PID21-35) Units: Hz | | | | | Amplitude - sIPSCs onto pyramidal neurons (PID21-35) Units: pA | | | | |
| | | CCI | -2.9437 | 1.4551 | -2.023 * | | CCI | 5.504 | 1.56 | 3.528 * | |
| | | 2-DG (<i>in vivo</i>) | -0.258 | 1.4203 | -0.182 | | 2-DG (<i>in vivo</i>) | 2.97 | 1.533 | 1.938 | |
| | | Interaction CCI/2-DG | 2.8493 | 2.0781 | 1.371 | | Interaction CCI/2-DG | -5.487 | 2.21 | -2.483 * | |
| 6 | 3 slices analyzed to generate 1 average value per animal Sham+Veh: 5 animals Sham+2-DG: 5 animals CCI+Veh: 7 animals CCI+2-DG: 6 animals | PV+ Density Units: cells/10,000 μm^2 | | | | | tdT+ Density Units: cells/10,000 μm^2 | | | | |
| | | Interaction CCI/ROI1 | -1.771 | 0.2083 | -8.501 * | | Interaction CCI/ROI1 | -1.665 | 0.2567 | -6.487 * | |
| | | Interaction CCI/ROI2 | -1.148 | 0.2083 | -5.513 * | | Interaction CCI/ROI2 | -1.027 | 0.2567 | -4 * | |
| | | Interaction CCI/ROI3 | -0.6815 | 0.2083 | -3.271 * | | Interaction CCI/ROI3 | -0.4762 | 0.2567 | -1.855 | |
| | | Interaction CCI/ROI4 | -0.4185 | 0.2083 | -2.009 * | | Interaction CCI/ROI4 | -0.2377 | 0.2567 | -0.926 | |
| | | Interaction CCI/2-DG/ROI1 | 1.111 | 0.2997 | 3.707 * | | Interaction CCI/2-DG/ROI1 | 0.8247 | 0.3693 | 2.233 * | |
| | | Interaction CCI/2-DG/ROI2 | 0.9103 | 0.2997 | 3.037 * | | Interaction CCI/2-DG/ROI2 | 0.532 | 0.3693 | 1.44 | |
| | | Interaction CCI/2-DG/ROI3 | 0.7652 | 0.2997 | 2.553 * | | Interaction CCI/2-DG/ROI3 | 0.6053 | 0.3693 | 1.639 | |
| | | Interaction CCI/2-DG/ROI4 | 0.4751 | 0.2997 | 1.585 | | Interaction CCI/2-DG/ROI4 | 0.5779 | 0.3693 | 0.156 | |
| | | PV+ tdT+ Colocalized Density Units: cells/10,000 μm^2 | | | | | Ratio of tdT+ Cells Lacking PV+ Units: ratio | | | | |
| | | Interaction CCI/ROI1 | -1.393 | 0.1829 | -7.614 * | | Interaction CCI/ROI1 | 0.364877 | 0.05684 | 6.419 * | |
| | | Interaction CCI/ROI2 | -0.9167 | 0.1829 | -5.011 * | | Interaction CCI/ROI2 | 0.191243 | 0.05684 | 3.365 * | |
| | | Interaction CCI/ROI3 | -0.5225 | 0.1829 | -2.856 * | | Interaction CCI/ROI3 | 0.091818 | 0.05684 | 1.615 | |
| | | Interaction CCI/ROI4 | -0.248 | 0.1829 | -1.356 | | Interaction CCI/ROI4 | 0.027345 | 0.05684 | 0.481 | |
| | | Interaction CCI/2-DG/ROI1 | 0.761 | 0.2632 | 2.892 * | | Interaction CCI/2-DG/ROI1 | -0.336425 | 0.081767 | -4.114 * | |
| | | Interaction CCI/2-DG/ROI2 | 0.5429 | 0.2632 | 2.063 * | | Interaction CCI/2-DG/ROI2 | -0.157418 | 0.081767 | -1.925 | |
| | | Interaction CCI/2-DG/ROI3 | 0.6884 | 0.2632 | 2.616 * | | Interaction CCI/2-DG/ROI3 | -0.139133 | 0.081767 | -1.702 | |
| | | Interaction CCI/2-DG/ROI4 | 0.2361 | 0.2632 | 0.897 | | Interaction CCI/2-DG/ROI4 | -0.087048 | 0.081767 | -1.065 | |

Table S1. Linear mixed model results, organized by figure.

| Type III Analysis of Variance with Satterthwaite's Method | | | |
|---|--------------------|-----------|-------------|
| Data | Effect/Interaction | p value | Significant |
| PV+ Density | CCI | 1.83E-04 | * |
| | 2-DG | 0.074809 | |
| | ROI | 1.27E-11 | * |
| | CCI/ROI | 5.83E-15 | * |
| | 2-DG/ROI | 1.21E-04 | * |
| | CCI/2-DG | 0.058903 | |
| | CCI/2-DG/ROI | 0.002792 | * |
| tdT+ Density | CCI | 6.07E-09 | * |
| | 2-DG | 0.35164 | |
| | ROI | 1.02E-06 | * |
| | CCI/ROI | 1.48E-11 | * |
| | 2-DG/ROI | 0.09617 | |
| | CCI/2-DG | 0.10985 | |
| | CCI/2-DG/ROI | 0.11161 | |
| PV+ tdT+ Colocalized Density | CCI | 0.0037177 | * |
| | 2-DG | 0.0729436 | |
| | ROI | 9.72E-09 | * |
| | CCI/ROI | 7.91E-15 | * |
| | 2-DG/ROI | 2.34E-04 | * |
| | CCI/2-DG | 0.101366 | |
| | CCI/2-DG/ROI | 0.0202878 | * |
| Ratio of tdT+ Lacking PV+ | CCI | 0.307487 | |
| | 2-DG | 0.067198 | |
| | ROI | 1.58E-06 | * |
| | CCI/ROI | 2.78E-06 | * |
| | 2-DG/ROI | 1.34E-06 | * |
| | CCI/2-DG | 0.127384 | |
| | CCI/2-DG/ROI | 0.001373 | * |

Table S2. Type III ANOVA results.