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1 **Astrocytes mediate cell non-autonomous correction of aberrant firing in**
2 **human FXS neurons**

3
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35 **Summary**

36
37 Pre-clinical studies of fragile X syndrome (FXS) have focused on neurons with the role of glia
38 remaining largely underexplored. We examined the astrocytic regulation of aberrant firing of FXS
39 neurons derived from human pluripotent stem cells. Human FXS cortical neurons, co-cultured with
40 human FXS astrocytes, fired frequent short duration spontaneous bursts of action potentials
41 compared to less frequent, longer duration, bursts of control neurons co-cultured with control
42 astrocytes. Intriguingly, bursts fired by FXS neurons co-cultured with control astrocytes are
43 indistinguishable from control neurons. Conversely, control neurons exhibit aberrant firing in the
44 presence of FXS astrocytes. Thus, the astrocyte genotype determines the neuronal firing
45 phenotype. Strikingly, astrocytic conditioned medium, and not the physical presence of astrocytes,
46 is capable of determining the firing phenotype. The mechanistic basis of this effect indicates that
47 the astroglial-derived protein, S100 β , restores normal firing by reversing the suppression of a
48 persistent sodium current in FXS neurons.

49 **Introduction**

50 Fragile X syndrome (FXS), a leading genetic cause of intellectual disability and autism spectrum
51 disorder, is caused by the absence of the Fragile X messenger ribonucleoprotein 1 (FMRP)
52 produced by the *FMRI* gene¹⁻⁴. Accumulating evidence from rodent models of FXS have given
53 shape to a useful framework for investigating how the absence of FMRP contributes to various
54 molecular, neuronal, and behavioral abnormalities⁵⁻⁸. As valuable as these findings have been in
55 advancing our understanding of the mechanisms by which absence of FMRP leads to
56 pathophysiology and for identifying potential therapeutic strategies for treating FXS, none has
57 translated into effective clinical outcomes for patients⁹⁻¹². This underscores the need for alternative
58 strategies to model FXS using ‘all-human’ based platforms.

59 It is in this context that human pluripotent stem cell (hPSC)-derived neurons offer a useful model
60 system¹³⁻¹⁵. The comparatively few studies that have used human *FMRI* null neurons to model
61 FXS have reached different conclusions regarding the nature of the changes in cellular excitability
62 and action potential firing properties. Nevertheless, they each share the common methodological
63 approach of focusing only on neuronal function¹⁶⁻²¹. Such a neuro-centric approach ignores the
64 pivotal role of glia on the development, maintenance and regulation of neuronal function and
65 homeostasis^{22,23}, including neuronal rhythmogenesis²⁴. In the context of FXS, co-culture of

66 hippocampal neurons from *Fmr1*^{-/-} mice with wild-type astrocytes prevents the emergence of an
67 altered dendritic morphology that is observed in co-cultures where neurons and astrocytes each do
68 not express FMRP²⁵. Furthermore, astrocyte-specific loss of FMRP expression has been reported
69 to result in increased spine density in cortical neurons and impaired motor learning in mice²⁶.
70 Moreover, it has been proposed that molecules secreted by astrocytes are required for synapse
71 formation and when absent in the astrocyte secretome from *Fmr1*^{-/-} mice this leads to alterations
72 in both spine morphology and synapse maturation²⁷. Together, these rodent studies have
73 contributed to steadily increasing evidence that astrocytes play essential roles in modulating the
74 function of neurons and neural circuits. Nevertheless, studies examining the glia contribution to
75 modulating neuronal electrophysiological phenotypes in a human model of FXS are lacking.
76 Thus, in the present study we addressed this gap in knowledge by carrying out detailed
77 electrophysiological analyses in co-cultures of cortical neurons and astrocytes each derived from
78 human pluripotent stem cells (hPSCs) lacking FMRP. Our results identify an important cell non-
79 autonomous contribution of human astrocytes in correcting aberrant electrical activity in human
80 FXS neurons, thereby suggesting a framework for exploring new therapeutic strategies aimed at
81 human neuron-glia interactions.

82

83 **Results**

84 Fibroblast-derived iPSCs were generated from one control (CON) and two FXS patient lines
85 (FXS1, FXS2). Neural precursor cells (NPCs) and astrocyte precursor cells (APCs), generated
86 from hPSCs, were terminally differentiated into cortical neurons²¹ and astrocytes (Figure S1A),
87 respectively. In addition, to ensure that the differences we observe in FXS patient-derived lines
88 are due to the silencing of the *FMRI* gene and the absence of FMRP, and not due to differences in
89 genetic backgrounds of the iPSC-derived neurons, we also analyzed a pair of isogenic embryonic
90 stem cell (ESC) derived cells²⁸ that includes one in which the *FMRI* gene was genetically deleted
91 (*FMRI*^{-/-}), and an otherwise genetically identical control line (*FMRI*^{+/-}). In each of these cell
92 lines, the absence of FMRP (Figure S1B) and expression of pluripotent stem cell markers were
93 confirmed (Figure S1C, Table S1).

94 **Astrocyte genotype determines the electrophysiological phenotype of neurons**

95 As a first step in our analysis, we used current-clamp recordings to characterize spontaneous bursts
96 of action potentials fired in human cortical neurons, co-cultured with human astrocytes, each
97 derived from induced pluripotent stem cells (iPSCs) generated from a healthy individual (CON,
98 Figure 1A; Table S2). These control cortical neurons, co-cultured with control astrocytes, fired a
99 low number of bursts, but the durations of individual bursts were long (Figure 1B, D; CON).
100 However, human cortical neurons lacking FMRP, co-cultured with FXS astrocytes, derived from
101 an iPSC line generated from a FXS patient (FXS1), exhibited a strikingly different pattern of
102 activity consisting of a significantly higher number of bursts that were of a much shorter duration
103 (FXS1, Figure 1C, D; Table S2)²¹. This same effect of the loss of FMRP on burst firing was also
104 seen in a second patient iPSC-derived line (FXS2, Figure S2; Table S3). Furthermore, the same
105 aberrant firing pattern was seen in neurons generated from a pair of isogenic embryonic stem cell
106 lines (*FMRI*^{+/-}; *FMRI*^{-/-}), in which the *FMRI* gene was deleted using CRISPR/Cas9-mediated
107 genome editing (Figure S2; Table S4). This aberrant pattern of activity in cortical neurons lacking
108 FMRP, manifested as higher number of shorter bursts (Figure 1C) compared to those seen in
109 control neurons (Figure 1B), will henceforth be referred to as the “FXS” firing pattern (Figure 1C).
110 Surprisingly, the equivalent FXS neurons, when co-cultured with control astrocytes, exhibited
111 normal burst firing consisting of a lower number of longer bursts (Figure 1E, G, H; Figure S2).
112 Conversely, in the presence of FXS astrocytes, control neurons exhibited aberrant FXS firing
113 patterns (Figure 1F, G, I). In other words, whether cortical neurons exhibit normal or FXS bursting
114 activity is determined by the genotype of the astrocyte, not the genotype of the neuron (Figure 1H,
115 I).

116 **FMRP re-expression in the FXS astrocytes rescues the aberrant bursting in FXS neurons**

117 To ascertain causality of this aberrant electrophysiological phenotype to FXS repeat expansion
118 mutation and loss of FMRP expression in astrocytes, we removed CGG repeats in FXS2 iPSC line
119 via two gRNAs flanking repeat expansion mutation using CRISPR/Cas9 methodology^{29,30}. The
120 successful gene corrected iPSC clone, FXS2Δ, displayed re-expression of FMRP protein (Figure
121 1J, K) following the removal of CGG repeats (Figure S3) and associated reversal of the
122 hypermethylation of *FMRI* promoter as shown via bi-sulfite sequencing (Figure 1L, Figure S3).
123 Moreover, FXS2 neurons co-cultured with FXS2Δ astrocytes exhibited normal burst firing – less
124 frequent bursts of longer duration (Fig. 1M, N).

125 **Conditioned medium from control astrocytes by itself restores normal firing in FXS neurons,**
126 **while medium from FXS astrocytes elicits aberrant firing in control neurons**

127 Next, we asked if the astrocytes must be physically present in the same culture to elicit this cell
128 non-autonomous effect on neuronal firing. Specifically, we asked if the astrocytic conditioned
129 medium (ACM) alone, i.e., the cell supernatant from pure astrocyte cultures, can influence the
130 burst firing properties of the cortical neurons even in the absence of the astrocytes themselves. To
131 test this, we examined all four combinations of control/FXS neurons grown in the presence of the
132 control/FXS ACM alone (Figure 2A; Table S5-7). Similar to the experiments described above,
133 current-clamp recordings were carried out from cortical neurons during the 8th week of cell culture.
134 For these experiments the neurons were grown in ACM, and not in the physical presence of
135 astrocytes, for the preceding 5 weeks prior to electrophysiological analysis. Control neurons grown
136 in control ACM exhibited normal burst firing (Figure 2B, D). Similarly, FXS neurons, grown in
137 FXS ACM, also fired aberrant bursts as seen earlier in neurons grown in the presence of FXS
138 astrocytes (Figure 1C, D). These results show that just the conditioned medium obtained from
139 astrocytes of the same genotype is adequate for the neurons to exhibit the same firing patterns
140 described in Figure 1B, C.

141 We next asked whether conditioned medium from astrocytes of one genotype can alter the
142 bursting property of neurons of the *other* genotype? Strikingly, FXS cortical neurons, when grown
143 in control ACM, displayed normal burst firing (Figure 2E, G). On the other hand, control neurons
144 exhibited FXS bursting patterns when grown in FXS ACM (Figure 2F, G). Therefore, the physical
145 proximity of astrocytes in co-cultures is not essential to switch the firing patterns of cortical
146 neurons – the ACM alone is adequate to elicit the cell non-autonomous effects (Figure 2H, I). This
147 was also seen in the isogenic pair and the second patient-derived lines (*FMRI*^{+y}, *FMRI*^{-y}; FXS2;
148 Figure S4).

149 **Persistent sodium currents are reduced in neurons exhibiting aberrant FXS firing**

150 The results presented so far show that co-cultured control astrocytes and their conditioned media
151 can correct the aberrant FXS bursting pattern. Conversely, co-cultured FXS astrocytes and their
152 conditioned medium can induce aberrant FXS bursting in cortical neurons. The following question
153 arises: what is the underlying electrophysiological basis of the alterations in firing caused by FXS,
154 and its correction? To address this question, we turned to several earlier findings on rhythmic burst
155 firing in neurons. For instance, analysis of altered neuronal excitability in the entorhinal cortex of

156 *Fmr1* KO mice³¹ implicated changes in the persistent sodium current (I_{NaP}). This current, which
157 is active at subthreshold voltages, was also shown to determine the properties of neuronal burst
158 firing in other studies³²⁻³⁵. More recently a role for I_{NaP} has been identified in bursting activity of
159 human neurons lacking FMRP co-cultured with mouse astrocytes²¹. Hence, based on previously
160 published experimental protocols^{21,31}, a slow depolarizing voltage ramp (20 mV/s; Figure 3A, *top*)
161 was applied to control cortical neurons, co-cultured with control astrocytes, and the I_{NaP} was
162 pharmacologically isolated (Figure 3A, *bottom, black trace*). Compared to control neurons, the
163 magnitude of I_{NaP} was reduced significantly in FXS neurons co-cultured with FXS astrocytes
164 (Figure 3A, *bottom, red trace*). Similarly, we observed a significant decrease in I_{NaP} in control
165 neurons, co-cultured with FXS astrocytes (Figure 3B, *red trace*), compared to FXS neurons co-
166 cultured with control astrocytes (Figure 3B, *black trace*). Therefore, in the presence of FXS
167 astrocytes, cortical neurons exhibited a reduction in I_{NaP} (Figure 3C, D), and this reduced level of
168 current is correlated with aberrant burst firing.

169 **Enhancement of persistent sodium currents reverses aberrant FXS firing**

170
171 Taken together, these results suggest a potential link between the reduction in the I_{NaP} current to
172 the aberrant FXS bursting. This raises the possibility that a pharmacological reagent specifically
173 designed to enhance I_{NaP} should also be effective in restoring normal bursting activity in FXS
174 neurons. We tested this prediction by applying the I_{NaP} opener, veratridine, during our recordings
175 from cortical neurons. First, after confirming the decrease in I_{NaP} in FXS neurons co-cultured with
176 FXS astrocytes (Figure 4A, *-veratridine*), veratridine (0.5 μ M) was bath applied for 20 minutes
177 (Figure 3A, *+veratridine*). This treatment elicited the expected effect, i.e., an increase in I_{NaP} in the
178 same neuron (Figure 4C). The same *in vitro* treatment with veratridine also reversed the
179 impairment in I_{NaP} control neurons co-cultured with FXS astrocytes (Figure 4B, D).

180 Having established the efficacy of veratridine in enhancing I_{NaP} in two different human
181 neuron-astrocyte combinations, we addressed the key question – will the same treatment also
182 correct the aberrant FXS bursting pattern? To test this, we first established a baseline of aberrant
183 bursting for 10 minutes (Figure 4E, “1”) followed by bath application of veratridine for 20 minutes
184 (Figure 4E, “2”, “3”) while continuing to record from the same neuron. Indeed, the same
185 concentration of veratridine that reversed the reduction in I_{NaP} also restored normal burst firing in
186 FXS neurons co-cultured with FXS astrocytes (Figure 4F, H, I). Furthermore, aberrant bursting in
187 control neurons, co-cultured with FXS astrocytes, was also rectified by the same veratridine

188 treatment (Figure 4G, H, K). Thus, in both co-culture combinations, FXS neurons with FXS
189 astrocytes, and control neurons with FXS astrocytes, that exhibit aberrant bursting patterns
190 veratridine restores normal firing. In other words, this correction of aberrant activity can be
191 achieved by just reversing the deficit in the neuronal I_{NaP} , despite the presence of FXS astrocytes
192 in the co-culture.

193 **Antibody inactivation of S100 β alters neuronal firing patterns in control but not FXS co-**
194 **cultures while addition of S100 β restores normal firing by reversing the suppression of**
195 **persistent sodium currents**

196 Taken together, these results strongly suggest that an astrocyte derived soluble factor(s) determine
197 if a neuron, irrespective of its genotype, will fire like a control or FXS neuron. This led us to ask
198 what compound(s) in the secretome from the astrocytes may mediate this switch between the
199 control and FXS burst firings patterns. In our search for a potential mechanism that mediates the
200 cell non-autonomous effect of correcting aberrant activity in FXS neurons, we reasoned that such
201 an astroglial-derived factor should be capable of eliciting the transition from FXS to control burst
202 firing. Specifically, this factor should, similar to veratridine, be capable of restoring normal firing
203 by reversing the reduction in I_{NaP} . Earlier findings suggested a promising candidate in S100 β , a
204 calcium-binding protein that is expressed primarily by astrocytes³⁶⁻³⁸. Importantly, rhythmic
205 bursting patterns, similar to those seen here, were shown to be regulated by S100 β released from
206 astrocytes in mice by modulating the I_{NaP} current³⁹. Thus, its glial specificity and previously
207 established role in neuronal rhythmogenesis, led us to explore if S100 β is capable of correcting
208 aberrant FXS firing in neurons. If S100 β were to fulfil such a role it would be predicted that its
209 inactivation would have a differential effect on ‘control neuron/control astrocyte’ compared with
210 ‘FXS neuron/FXS astrocyte’ co-cultures. As illustrated in Figure 5A, C the typical long-lasting
211 but less frequent firing pattern of control neurons (co-cultured with control astrocytes) changes to
212 one displaying shorter bursts of firing upon addition of an antibody to S100 β . Importantly, the
213 equivalent antibody treatment of FXS neurons (co-cultured with FXS astrocytes) did not alter the
214 firing pattern of FXS neurons (Figure 5B, D). As illustrated in Figure 5E we also confirmed that
215 the concentration of S100 β was lower in ACM from FXS astrocyte cultures compared to that in
216 ACM obtained from control astrocytes.

217 If the reduction in a potential astrocyte-derived candidate, S100 β , is mediating the aberrant
218 FXS firing patterns, then this raises the possibility that addition of S100 β would reverse the
219 decrease in I_{NaP} associated with this pattern. To test this, we first confirmed the reduction in I_{NaP}
220 in FXS neurons co-cultured with FXS astrocytes was confirmed (Figure 6A, -S100 β). Next,
221 S100 β (5 μ M) was bath applied for 20 minutes while continuing to record from the same neuron
222 (Figure 6A, +S100 β). Similar to veratridine, this treatment elicited an increase in I_{NaP} in the same
223 neuron (Figure 6C). Treatment with S100 β also reversed the impairment in I_{NaP} seen in control
224 neurons co-cultured with FXS astrocytes (Figure 6B, D). After confirming that S100 β potentiates
225 the I_{NaP} , we tested if it also reverses the aberrant FXS burst firing. We first confirmed that FXS
226 neurons in the presence of FXS astrocytes exhibit the expected aberrant firing (Figure 6F, “1”).
227 But, when S100 β was bath applied after the baseline recording (Fig, 6F “2”), the same FXS neuron
228 exhibited normal bursting patterns (Figure 6F, “3”). We also checked the efficacy of the same
229 treatment on control neurons co-cultured with FXS astrocytes (Figure 6G). Here too the equivalent
230 bath application of S100 β switched FXS firing to normal firing (Figure 6G, “1” & “3”). Thus,
231 under both conditions that give rise to aberrant FXS bursting in neurons, S100 β succeeded in
232 restoring normal firing despite the presence of the FXS astrocytes (Figure 6H-K).

233 Thus, in our human cell co-culture model to study FXS, the observation that whether burst firing
234 activity in human cortical neurons is normal or aberrant depends on the genotype of the astrocytes
235 present and not that of the neurons. Moreover, the physical presence of astrocytes is not required
236 as astrocyte conditioned medium by itself is able to elicit the same cell non-autonomous effects.
237 Finally, our data revealed that the concentration of the astroglial-derived protein, S100 β , is reduced
238 in the secretome of FXS astrocytes and that addition of S100 β to co-cultures of either control or
239 FXS neurons with FXS astrocytes restores normal firing patterns by reversing the suppression, in
240 neurons, of a persistent sodium current.

241

242 **Discussion**

243 Here, we identify a pivotal role for astrocytes in correcting aberrant electrical activity in human
244 FXS neurons, thereby shifting the focus to astrocytes in what was traditionally thought of as a
245 developmental disorder involving neuronal dysfunction. Earlier analyses of deficits caused by

246 FXS, and strategies to reverse them, were aimed primarily at understanding synaptic and molecular
247 signaling abnormalities^{15,40,41}. More recent work has also examined alterations in intrinsic and
248 network excitability, as well as modulation of ion channels by FMRP⁴²⁻⁴⁵. However, these
249 analyses centered exclusively on neurons, and primarily were conducted in animal models of FXS.
250 Recently, a handful of studies have characterized the impact of the loss of FMRP on excitability
251 in human neurons. In one study, depolarizing current injections into human neurons lacking FMRP
252 elicited at most single action potentials, not the more complex patterns of burst firing observed
253 here¹⁸. Another study revealed higher spontaneous firing in neurons lacking FMRP, but used
254 extracellular multi-electrode arrays, thereby lacking single-cell resolution²⁰. We have also recently
255 reported that human cortical neurons lacking FMRP display aberrant burst firing properties
256 compared to control neurons²¹. However, and importantly for the present study, these earlier
257 reports did not use human astrocytes in the co-culture but rather, if astrocytes were indeed present,
258 these were of rodent origin. Specifically, in our previous study²¹ the genotype of the co-cultured
259 mouse-derived astrocytes had no effect on the electrophysiological phenotype observed in the
260 human cortical neurons. We do not have an explanation as to why in our previous study²¹ co-
261 culture of human FXS neurons with (wild-type) mouse astrocytes did not result in FXS neurons
262 displaying a control firing pattern. Consistent between our previous and current studies is the fact
263 that the magnitude of the I_{NaP} predicts the firing pattern that is observed; it is greater in control
264 neurons than FXS neurons and pharmacological interventions that enhance I_{NaP} in FXS neurons
265 result in control-like firing patterns. Nevertheless, the presence of wild-type mouse astrocytes
266 does not appear to influence the expression of I_{NaP} in human FXS neurons albeit that our earlier
267 study show that, pharmacologically, these levels can be increased, at least functionally. Thus,
268 while future studies will be needed to investigate if different mechanisms of regulation are at play
269 when human FXS neurons are co-cultured with human versus rodent astrocytes, there is a
270 convergence at the level of the functional alterations of I_{NaP} . The present study underscores the
271 importance of the ‘all-human’ co-culture system in revealing new mechanistic insights into how
272 astrocytes mediate the effects reported here. Our analyses in human neuron-astrocyte co-cultures
273 revealed striking differences in the duration and number of spontaneous bursts of action potentials
274 fired by FMRP-lacking cortical neurons. Importantly, we find the transitions between normal and
275 aberrant bursting to be cell non-autonomous. This, in turn, enabled us to probe the mechanistic
276 basis of neuron-glia interactions underlying alterations in excitability of human FXS cortical
277 neurons.

278 Our results also add to accumulating evidence on how astrocytes modulate neuronal function in
279 both health and disease. Disease models have shown that astrocytes can, depending on disease
280 context, be either protective or injurious⁴⁶⁻⁵⁰. In the context of neurodevelopmental disorders,
281 astrocytes have been implicated in the progression of Rett's syndrome in a mouse model wherein
282 selective re-expression of methyl-CpG-binding protein 2 (*Mecp2*) in astrocytes restored normal
283 dendritic morphology *in vivo*⁵¹. Studies using iPSCs from Down's Syndrome (DS) patients also
284 reported a role for astroglia, as well as astrocytic conditioned medium, in abnormal neuronal
285 phenotypes in DS pathology^{52,53}. Consistent with our results, these studies also emphasized the
286 role of S100 β in the interaction between DS astroglia and DS neurons. Specifically, in rodent
287 models of FXS, wild-type astrocytes, when co-cultured of hippocampal neurons lacking FMRP,
288 reverse the altered dendritic morphology that is present when astrocytes also lack FMRP²⁵.
289 Indeed, loss of expression of FMRP from astrocytes alone leads to increased spine density in
290 (FMRP-expressing) cortical neurons and impaired motor learning in mice demonstrating astrocyte
291 modulation of neuronal morphology and behaviour²⁶. Consistent with our current study showing
292 that the physical presence of astrocytes is not required to impart changes in neuronal firing
293 phenotype is the demonstration that the astrocyte secretome from *Fmr1*^{-/-} mice influences spine
294 morphology and synapse maturation²⁷. It is also worth noting that the correction achieved by
295 S100 β in the present study does not completely re-capitulate the firing patterns seen in control
296 neurons (Figure 6H, J), raising the possibility that other factors besides S100 β may also contribute
297 to this process. For instance, based on earlier studies, the efficacy of other promising candidate
298 astroglial factors, such as IL6, remains to be explored. Our results provide an electrophysiological
299 framework to examine these in future studies.

300
301 Finally, analysis of how S100 β mediates its protective effect (i.e., switching FXS firing to control)
302 revealed the persistent sodium conductance (I_{NaP}) as a key element. Specifically, we found a
303 reduction in the I_{NaP} in neurons that exhibited the aberrant FXS firing pattern, irrespective of the
304 genotype of the neuron in a range of combinations (i.e., FXS neurons, control neurons in the
305 presence of FXS astrocyte or its conditioned medium; Figure 3A, B, *red traces*). Moreover,
306 concentrations of S100 β that reversed this aberrant FXS burst firing to control patterns, also
307 restored suppressed I_{NaP} currents to control levels (Figure 6A, B, *grey traces*). Importantly,
308 veratridine, a specific I_{NaP} opener, corrected the FXS burst firing to normal patterns, thereby
309 confirming the mode of action through which S100 β achieves its corrective action on aberrant

310 excitability seen in FXS neurons (Figure 4A-G). Interestingly, a similar mechanism was also seen
311 earlier in the rat sensorimotor circuit wherein astrocytes were involved in generating neuronal
312 rhythmic burst firing³⁹. Furthermore, when inactivation of astrocytes prevented neuronal bursting
313 in this study, it was restored by adding S100 β to the extracellular space, which in turn was mediated
314 by an enhancement in I_{NaP} . Similar to what we report here (Figure 6C, D), locally applied S100 β
315 also increased the peak amplitude of the isolated I_{NaP} . Thus, our findings also open up possibilities
316 to investigate the *in vivo* functional consequences of these cell non-autonomous glial phenotypes
317 in rodent models of FXS. The co-culture system described here offers a new framework for
318 exploring mechanisms underlying neuron-glia interactions and their efficacy in correcting
319 aberrations in neuronal excitability caused by FXS. Together, these findings elucidate previously
320 undescribed mechanisms of FXS pathogenesis and identify potential therapeutic targets in
321 astrocytes.

322 **Limitations of the study**

323 We have identified a key role for astrocytic regulation of neuronal firing phenotypes in an ‘all-
324 human’ culture-based model of FXS. While our data support the notion that astrocytic-derived
325 S100 β restores control-like firing in FXS neurons by enhancing a persistent sodium current we
326 cannot exclude the possibility that other factors in the astrocyte secretome also can act in a similar
327 manner. Equally, while our study has highlighted that larger magnitude persistent sodium currents
328 are present in control neurons and we can phenocopy control neuronal firing patterns in FXS
329 neurons by co-culturing with control astrocytes or by adding control ACM, veratridine or S100 β
330 each of which enhances a persistent sodium current, we remain open to the idea that other ion
331 channels could also play a pivotal role in generating the distinct firing patterns seen in control and
332 FXS neurons. In other words, simply phenocopying the firing pattern does not imply that each are
333 generated in the same manner, mechanistically. Nevertheless, if the temporal nature of neuronal
334 output is central to circuit dysfunction present in models of FXS then ‘correcting’ aberrant firing
335 to restore normal firing might be an approach, while agnostic to its mechanistic basis, that allows
336 further identification of potential therapeutic targets.

337

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350 **Author contributions:** SDS, BKR, RP, TER, JDC, BTS, PCK, SiC, DJAW, and SuC were
351 responsible for the conception and design of the experiments. SDS performed all the
352 electrophysiological recordings. BKR and RP carried out human stem cell (hPSC) culture
353 including propagation and characterization of the cell lines. BKR carried out all the astrocyte
354 culture, including propagation and characterization of the hPSC-derived astrocytes. SDS, BKR,
355 RP, TER, JDC and BTS were responsible for the collection and assembly of data. SDS, BKR, RP,
356 TER, JDC, BTS, PCK, SiC, DJAW and SuC were responsible for the analysis and interpretation
357 of data. SDS, BKR, RP, BTS, PCK, SiC, DJAW, and SuC wrote the manuscript, and all authors
358 had the opportunity to contribute to its editing. All persons designated as authors qualify for
359 authorship, and all those who qualify for authorship are listed. The authors read and approved the
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361 **Declaration of interests:** Authors declare that they have no competing interests.

362 **Inclusion and diversity:** One or more of the authors of this paper self-identifies as an under-
363 represented ethnic minority in their field of research or within their geographical location. We
364 support inclusive, diverse and equitable conduct of research.

365 **Figure Legends**

366 **Figure 1. Distinct patterns of bursting activity of hPSC-derived cortical neurons is**
367 **determined by the genotype of the astrocyte and its secretome, not the genotype of the**
368 **neuron. (A)** Representative confocal images of hiPSC-derived cortical neurons (CON and FXS1)
369 expressing Map2ab co-cultured with hiPSC derived astrocytes expressing GFAP. *Left to right,*
370 neurons were co-cultured in four combinations: CON neurons with CON astrocytes, FXS1 neurons

371 with FXS1 astrocytes, FXS1 neurons with CON astrocytes, and CON neurons with FXS1
372 astrocytes. *Right*, legend for icons. Scale bar = 50 μm . **(B)** (i) Representative current-clamp
373 recording ($V_{\text{HOLD}} = -70 \text{ mV}$) from a CON neuron, co-cultured with CON astrocyte, firing
374 spontaneous bursts of action potentials that occur at low frequencies but with long durations. (ii)
375 Single burst shown in expanded time-base. **(C)** (i) Representative current-clamp recording of
376 aberrant spontaneous activity in a FXS neuron, co-cultured with FXS1 astrocyte, containing a
377 significantly higher number of bursts but of shorter duration. (ii) Single burst shown in expanded
378 time-base. **(D)** Comparison of mean burst duration (CON neurons with CON astrocytes, $45.19 \pm$
379 4.065 s , $n = 24$, $N = 4$; FXS1 neurons with FXS1 astrocytes, $10.97 \pm 1.126 \text{ s}$, $n = 13$, $N = 3$) and
380 mean burst number per 10 min of recording (CON neurons with CON astrocytes, 6.167 ± 0.89 ;
381 FXS1 neurons with FXS1 astrocytes, 25.92 ± 3.482). **(E)** Representative trace from a FXS1
382 neuron, co-cultured with CON astrocyte, exhibits a normal activity with low burst number and
383 longer burst duration. **(F)** Conversely, a CON neuron, co-cultured with FXS1 astrocyte, shows
384 aberrant activity containing higher number of bursts of shorter duration. **(G)** Comparison of mean
385 burst duration (FXS1 neurons with CON astrocytes, $31.6 \pm 4.42 \text{ s}$, $n = 17$, $N = 3$; CON neurons
386 with FXS1 astrocytes, $9.37 \pm 1.32 \text{ s}$, $n = 18$, $N = 3$) and mean burst number per 10 min of recording
387 (FXS1 neurons with CON astrocytes, 5.18 ± 1.033 ; 33.39 ± 3.96). **(H)** Summary: control astrocytes
388 correct aberrant burst firing in FXS neurons. **(I)** Summary: control neurons switch to aberrant
389 firing in the presence of FXS astrocytes. **(J)** Immunoblot and **(K)** quantification depicting re-
390 expression of FMRP ($n=3$) in FXS2 patient iPSC line and its isogenic corrected control (FXS 2Δ),
391 expressed as % of an unrelated control (C). **(L)** Bisulfite sequencing of the FMR1 promoter
392 showing its demethylation in the edited lines. **(M)** Representative current-clamp trace of
393 spontaneous activity recorded from a FXS2 neuron co-cultured with FXS2 Δ astrocytes exhibits
394 normal firing consisting of a low number of longer bursts. **(N)** Comparison of mean burst duration
395 ($37.6 \pm 3.84 \text{ s}$, $n = 15$, $N = 3$) and mean burst number per 10 min of recording (7.4 ± 0.80) in FXS2
396 neurons co-cultured with FXS2 Δ astrocytes. All representative traces of aberrant activity patterns
397 in cortical neurons are depicted in red, whereas normal spiking activity is depicted in black. On
398 the other hand, for the genetic identity of the neuron and astrocyte, controls are depicted in black
399 and FXS in red. *** $p < 0.001$, one-way ANOVA and post-hoc Sidak's test. All values are mean
400 \pm SEM.

401
402

403 **Figure 2. Astrocyte secretome is sufficient to determine the bursting activity patterns of**
404 **hPSC-derived cortical neurons. (A)** Representative confocal images of Map2ab-expressing
405 hiPSC-derived cortical neurons (CON and FXS1) grown in astrocytic conditioned media (ACM).
406 Left to right, neurons were grown in ACM in four combinations: CON neurons with CON ACM,
407 FXS1 neurons with. FXS1 ACM, FXS1 neurons with CON ACM, and CON neurons with FXS1
408 ACM. Right, legend for icons. Scale bar = 50 μm . **(B)** Recording from a CON neuron, grown in
409 CON ACM, exhibits normal firing consisting of a low number of longer bursts. **(C)** Recording
410 from FXS1 neuron grown in FXS1 ACM shows a high number of bursts of shorter duration. **(D)**
411 Comparison of mean burst duration (CON neurons in CON ACM, 42.29 ± 5.66 s, $n = 16$, $N = 3$;
412 FXS1 neurons in FXS1 ACM, 8.501 ± 0.98 s, $n = 12$, $N = 3$) and mean burst number per 10 min
413 of recording (CON neurons in CON ACM, 3.19 ± 0.46 ; FXS1 neurons in FXS1 ACM, $19.67 \pm$
414 2.13). **(E)** Representative recording from a FXS1 neuron, grown in CON ACM, exhibiting control
415 burst firing, i.e., less frequent bursts of longer duration. Thus, control ACM exerts a non-cell
416 autonomous effect in reversing aberrant FXS bursting. **(F)** Representative recording from a CON
417 neuron, grown in FXS1 ACM, shows the aberrant FXS bursting. **(G)** Comparison of mean burst
418 duration (FXS1 neurons in CON ACM, 51.66 ± 7.027 s, $n = 12$, $N = 3$; CON neurons in FXS1
419 ACM, 9.945 ± 1.026 s, $n = 13$, $N = 3$) and mean burst number per 10 min of recording (FXS1
420 neurons in CON ACM, 4.17 ± 0.78 s; CON neurons in FXS1 ACM, 28.38 ± 3.61). **(H)** Summary:
421 control ACM corrects aberrant burst firing in FXS neurons. **(I)** Summary: FXS ACM alone elicits
422 aberrant firing in control neurons. $***p < 0.001$, one-way ANOVA and post-hoc Sidak's test. All
423 values are mean \pm SEM.

424

425 **Figure 3. Neurons co-cultured with FXS astrocytes have reduced I_{NaP} densities. (A)** Persistent
426 sodium currents (I_{NaP}) isolated by subtracting current traces evoked by a slow depolarizing ramp
427 (*top*, -100 to -20 mV, 20 mV/s) in the absence and presence of TTX. Representative traces (*bottom*)
428 are from a CON neuron co-cultured with CON astrocyte (*black*), and a FXS1 neuron co-cultured
429 with FXS1 astrocyte (*red*). **(B)** Same as (A) but representative traces recorded from a CON neuron
430 co-cultured with FXS1 astrocyte (*red*) and a FXS1 neuron co-cultured with a CON astrocyte
431 (*black*). **(C)** Current density–voltage (I-V) curves plotted from the ramp-evoked persistent sodium
432 current (I_{NaP}). Currents are normalized to the corresponding cell capacitance. *Inset*: Comparison
433 of current densities at -20 mV reveals a significant reduction in I_{NaP} in FXS1 neurons co-cultured
434 with FXS1 astrocytes (*red*; -0.32 ± 0.17 , $n = 6$, $N = 1$) compared to CON neurons co-cultured with

435 CON astrocytes (*black*; -1.96 ± 0.6 , $n = 6$, $N = 1$). **(D)** Current density–voltage (I-V) curves plotted
436 from the ramp-evoked persistent sodium current (I_{NaP}). Currents are normalized to the
437 corresponding cell capacitance. *Inset*: Comparison of current densities at -20 mV reveals a
438 significant reduction in I_{NaP} in CON neurons co-cultured with FXS1 astrocytes (*red*; -0.322 ± 0.17 ,
439 $n = 6$, $N = 1$) compared to FXS1 neurons co-cultured with CON astrocytes (*black*; -1.62 ± 0.28 , n
440 $= 7$, $N = 2$). $***p < 0.001$, $**p < 0.01$, $*p < 0.05$, Two-way, repeated measures ANOVA with post
441 hoc Sidak's test (I-V plots), Mann-Whitney test. All values are mean \pm SEM.

442 **Figure 4. Pharmacological enhancement of the I_{NaP} corrects aberrant FXS firing. (A)**
443 Representative traces of persistent sodium currents (I_{NaP}) recorded from a FXS1 neuron co-
444 cultured with FXS1 astrocyte (*red*). The I_{NaP} amplitude increases after addition of veratridine
445 (*grey*). **(B)** Reduced I_{NaP} in a CON neuron, co-cultured with FXS1 astrocyte (*red*), is reversed by
446 application of veratridine (*grey*). **(C)** I-V plots for FXS1 neurons with FXS1 astrocytes in before
447 (*red*) and after veratridine (*grey*). *Inset*: Comparison of current densities at -20 mV reveals a
448 significant increase in I_{NaP} in FXS1 neurons co-cultured with FXS1 astrocytes before (*red*; -0.44
449 ± 0.106 , $n = 8$, $N = 2$) and after veratridine (*grey*; -1.15 ± 0.21). **(D)** I-V plots for CON neurons
450 with FXS1 astrocytes in before (*red*) and after veratridine (*grey*). *Inset*: Comparison of current
451 densities at -20 mV reveals a significant increase in I_{NaP} in CON neurons co-cultured with FXS1
452 astrocytes before (*red*; -0.293 ± 0.099 , $n=5$, $N=1$) and after veratridine (*grey*; -1.635 ± 0.34). **(E)**
453 Experimental protocol: baseline recording for 10 min (1), followed by bath application of
454 veratridine (2), and then continuing to record from the same neuron for another 20 min; bursting
455 properties from last 10 min were compared with first 10 min (3). **(F)** Baseline (*left trace, 1*) current-
456 clamp recording ($V_{HOLD} = -70$ mV) from a FXS1 neuron, co-cultured with FXS1 astrocyte, shows
457 more frequent bursts of shorter duration. Application of veratridine elicits a transition (*middle*
458 *trace, 2*) that eventually culminates in control patterns of bursting (*right trace, 3*), i.e., less frequent
459 bursts of longer duration. **(G)** As in (F) but traces recorded from a CON neuron co-cultured with
460 FXS1 astrocyte. **(H)** Comparison of mean burst duration (7.4 ± 0.51 s; 20.36 ± 1.74 s, $n=7$, $N=2$)
461 and mean burst number per 10 min of recording (44.14 ± 6.9 ; 26.71 ± 6.42) in FXS1 neurons with
462 FXS1 astrocytes in the absence (*red*) and presence (*grey*) of veratridine (Vera). **(I)** Summary:
463 veratridine reverses aberrant FXS burst firing to control firing. **(J)** Comparison of mean burst
464 duration (8.05 ± 1.1 s; 36.74 ± 4.6 s, $n=9$, $N=3$) and mean burst number per 10 min of recording
465 (29.22 ± 6.35 ; 9 ± 2.12) in CON neurons with FXS1 astrocytes in the absence (*red*) and presence

466 (grey) of veratridine. **(K)** Summary: veratridine reverses aberrant firing to control firing in CON
467 neurons co-cultured with FXS astrocytes. *** $p < 0.001$, ** $p < 0.01$, * $p < 0.05$, Two-way, repeated
468 measures ANOVA with post hoc Sidak's test (I-V plots), paired t -test, Wilcoxon test. All values
469 are mean \pm SEM.

470 **Figure 5. Blocking S100 β with an antibody alters the bursting pattern of neurons co-cultured**
471 **with control astrocytes (A)** Baseline (left) current-clamp recording ($V_{\text{HOLD}} = -70$ mV) from a
472 CON neuron, co-cultured with CON astrocyte, shows less frequent bursts of longer duration. In
473 presence of S100 β antibody (right) the control burst firing changes to that of aberrant FXS burst
474 firing. **(B)** Baseline (left) current-clamp recording ($V_{\text{HOLD}} = -70$ mV) from a FXS1 neuron, co-
475 cultured with FXS1 astrocyte, shows more frequent bursts of shorter duration. The aberrant FXS
476 burst firing is unaltered in the presence of S100 β antibody (right). **(C)** Comparison of mean burst
477 duration (38.69 ± 6.14 s; 7.48 ± 0.98 s, $n=8$, $N=2$) and mean burst number per 10 min of recording
478 (10.13 ± 2.3 ; 12.5 ± 3.13) in CON neurons with CON astrocytes in the absence (black) and
479 presence (red) of S100 β Ab. **(D)** Comparison of mean burst duration (5.18 ± 0.76 s; 5.98 ± 0.93 s,
480 $n=7$, $N=2$) and mean burst number per 10 min of recording (21.71 ± 2.17 ; 21.71 ± 1.82) in FXS1
481 neurons with FXS1 astrocytes in the absence and presence of S100 β Ab. **(E)** S100 β concentration
482 is lower (2362 ± 230 pg/ml) in ACM derived from FXS astrocytes compared with that measured
483 in ACM derived from control astrocytes (3301 ± 297 pg/ml; $N=5$). ** $p < 0.01$, * $p < 0.05$ unpaired
484 t -test, Mann-Whitney tests. All values are mean \pm SEM.

485 **Figure 6. S100 β corrects aberrant FXS bursting in neurons by enhancing I_{NaP}** **(A)** Persistent
486 sodium currents (I_{NaP}) isolated by subtracting current traces evoked by a slow depolarizing ramp
487 (*top*, -100 to -20 mV, 20 mV/s) in the absence and presence of TTX. Representative traces (*bottom*)
488 are from a FXS1 neuron co-cultured with FXS1 astrocyte in the absence (*red*) and presence of
489 S100 β (*grey*). **(B)** Same as (A) but representative traces recorded from a CON neuron co-cultured
490 with FXS1 astrocyte in the absence (*red*) and presence of S100 β (*grey*). **(C)** Current density–
491 voltage (I-V) curves plotted from the ramp-evoked persistent sodium current (I_{NaP}). Currents are
492 normalized to the corresponding cell capacitance. *Inset*: Comparison of current densities at -20
493 mV reveals a significant increase in I_{NaP} in FXS1 neurons co-cultured with FXS1 astrocytes before
494 (*red*; -0.44 ± 0.1 , $n = 8$, $N=3$) and after S100 β (*grey*; -1.41 ± 0.36). **(D)** Current density–voltage
495 (I-V) curves plotted from the ramp-evoked persistent sodium current (I_{NaP}). Currents are

496 normalized to the corresponding cell capacitance. *Inset*: Comparison of current densities at -20
497 mV reveals a significant increase in I_{NaP} in CON neurons co-cultured with FXS1 astrocytes before
498 (*red*; -0.41 ± 0.082 , $n = 8$, $N=3$) and after S100 β (*grey*; -1.64 ± 0.43). **(E)** Experimental protocol:
499 baseline recording for 10 min (1), followed by bath application of S100 β (2), and then continuing
500 to record from the same neuron for another 20 min; bursting properties from last 10 min were
501 compared with first 10 min (3). **(F)** Baseline (*left trace, 1*) current-clamp recording ($V_{HOLD} = -70$
502 mV) from a FXS1 neuron, co-cultured with FXS1 astrocyte, shows more frequent bursts of shorter
503 duration. Application of S100 β elicits a transition (*middle trace, 2*) that eventually culminates in
504 control patterns of bursting (*right trace, 3*), i.e., less frequent bursts of longer duration. **(G)** As in
505 (h) but traces recorded from a CON neuron co-cultured with FXS1 astrocyte. **(H)** Comparison of
506 mean burst duration (9.58 ± 1.71 s; 25.51 ± 2.76 s, $n = 10$, $N = 3$) and mean burst number per 10
507 min of recording (22.4 ± 2.76 ; 14.9 ± 1.88) in FXS1 neurons with FXS1 astrocytes in the absence
508 (*red*) and presence (*grey*) of S100 β . **(I)** Summary: S100 β reverses aberrant FXS burst firing to
509 control firing. **(J)** Comparison of mean burst duration (8.406 ± 1.4 s; 30.76 ± 4.5 s, $n = 8$, $N = 2$)
510 and mean burst number per 10 min of recording (31.63 ± 5.49 ; 11.13 ± 1.55) in CON neurons with
511 FXS1 astrocytes in the absence (*red*) and presence (*grey*) of S100 β . **(K)** Summary: S100 β reverses
512 aberrant firing to control firing in CON neurons co-cultured with FXS astrocytes. $***p < 0.001$,
513 $**p < 0.01$, $*p < 0.05$, Two-way, repeated measures ANOVA with post hoc Sidak's test (I-V plots),
514 paired *t*-test, Wilcoxon test. All values are mean \pm SEM.

515 **STAR Methods**

516 **RESOURCE AVAILABILITY**

517 **Lead Contact:**

518 Sumantra (Shona) Chattarji (shona@ncbs.res.in)

519 **Materials Availability:**

520 Further information and requests for resources and reagents should be directed to and will be
521 fulfilled by the Lead Contact, Sumantra (Shona) Chattarji (shona@ncbs.res.in).

522

523 **Data and Code Availability:**

- 524
- All data generated in this study will be made available upon reasonable request.
 - This paper does not report original code.
- 525

- 526 • Any additional information required to reanalyze the data reported in this work paper is
527 available from the Lead Contact upon request.

528 **EXPERIMENTAL MODEL AND SUBJECT DETAILS**

529 **Cell line source**

530 ND30625 (apparently healthy control male), GM05848 (fragile X syndrome male) and GM07072
531 (fragile X syndrome male) fibroblasts and hereafter referred to as CON, FXS1, FXS2 lines,
532 respectively, were obtained from the Coriell Institute for Medical Research under their consent
533 and privacy guidelines as described on their website (<http://catalog.coriell.org/>). Induced human
534 pluripotent stem cells were generated at Cedar-Sinai Medical Centre (Los Angeles, CA) using
535 standard protocols as described previously²¹. Briefly, fibroblasts were re-programmed into non-
536 integrating and virus-free hiPSC by nucleofection using episomal plasmids OCT4, SOX2, KLF4,
537 L-MYC, LIN28, and shRNA to TP53.

538 The human embryonic stem cell (hESC) line Shef 4, referred to as *FMRI*^{+y}, was obtained from
539 the UK Stem Cell Bank⁵⁴. The Shef 4 *FMRI* null line, referred to as *FMRI*^{-y}, was generated using
540 CRISPR-Cas9 technology as described previously²⁸. Further details about each of the cell lines
541 are provided in Table S1.

542 **Generation of FXS isogenic control iPSC line**

543 The FXS-2 iPSC line carrying the CGG repeat expansion mutation (Figure S3) was gene corrected
544 using CRISPR/Cas9 genome editing. In brief, two guide RNAs (gRNAs; gRNA-1: 5'.
545 GACGGAGGCGCCGCTGCCAG - 3' and gRNA-2 5'- GCCCGCAGCCCACCTCTCGG-3')
546 flanking the repeat expansion mutation were cloned in px458 vector using published protocols²⁹.
547 Gene editing in iPSCs was performed as per previously published protocol³⁰. Briefly, iPSCs were
548 dissociated into single-cell suspension with 1X Accutase (Sigma-Aldrich). 8 x 10⁵ cells were
549 transfected with 2 µg Cas9-gRNA-1 plasmid, 2 µg Cas9-gRNA-2 plasmid and 1 µg eGFP-
550 puromycin resistance plasmid using the Amaxa 4D-Nucleofector system (Lonza) according to the
551 manufacturer's instructions with a pulse load of CA137. Cells were plated down onto MatrigelTM
552 (Corning) coated plates in E8 medium with 10 µM ROCK inhibitor (Tocris). After 24 hours,
553 transfected cells were selected for with addition of 1 µg/ml puromycin (Sigma-Aldrich) for a
554 further 24 hours. Clonal analyses were performed on manually picked individual clones and

555 screened for successful gene-editing via immunofluorescence for FMRP and PCR flanking repeat
556 expansion mutation. Successful gene editing clone was determined by re-expression of FMRP and
557 presence of positive PCR amplicon. Successful gene-edited clones were further validated using
558 repeat primed PCR, using AmplideX FMRI PCR kit from Asuragen, which showed negative for
559 gene-edited clone (Figure S3).

560 **Maintenance and expansion of hPSCs**

561 hiPSC and hESC colonies were cultured and propagated in xeno-free and feeder-free conditions
562 using Essential 8 medium (Thermo Fisher Scientific) in reduced growth-factor MatrigelTM-coated
563 6-well plates (Nunc Nunclon delta surface, Thermo Fisher Scientific) at 37 °C, 5% CO₂ in a
564 humidified incubator. Colonies were grown to 90% confluence; enzymatically passaged using 2:1
565 ratio of collagenase (ThermoFisher Scientific, USA) and dispase (ThermoFisher Scientific, USA)
566 for further propagation and cryopreservation⁵⁵. All cells showed normal G banding karyotype
567 (Table S1) and were free from mycoplasma. All cell lines and experimental procedures used in the
568 present study (Study number: 22/SCR/VI-29.4.16) were approved by the Institutional Committee
569 for Stem Cell Research (IC-SCR), InStem, Bangalore, India.

570 **Derivation of Neural Progenitor Cells (NPCs)**

571 NPCs were derived from each of the cell lines mentioned above using previously published
572 protocols^{21,55}. Briefly, hPSCs were enzymatically lifted and cellular aggregates were placed on to
573 non-adherent dishes forming a suspension culture. The medium was supplemented with LDN
574 193189 (Strattech), SB431542 (Tocris) and basic fibroblast growth factor (FGF) (PeproTech) to
575 form neural spheres and rosettes at 37°C, 5% CO₂. Rosettes were dissociated using accutase and
576 plated as a monolayer at 37°C, 3% O₂ and 5% CO₂, which were then maintained, characterized,
577 expanded, and cryopreserved as NPCs.

578 **Generation of Astrocytic Progenitor Cells (APCs) and astrocyte differentiation**

579 hPSCs were neuralized and then converted to spheres as described earlier^{21,56}. Next, to induce
580 astroglialogenesis spheres were subjected to glial enrichment medium (EF20) containing Advanced
581 DMEM/F12 with 1% Anti-anti, 1% N2, 1% Glutamax, 0.1% B27, (each from ThermoFisher
582 Scientific) 20 ng/ml epidermal growth factor (EGF; R&D Systems), 20 ng/ml FGF, 5 mg/ml
583 heparin (Sigma-Aldrich) for 2 weeks before being cultured in glial maturation medium. For

584 maturation of early gliospheres, the medium (EL20) was supplemented with 20 ng/ml leukemia
585 inhibitory factor (LIF) (Sigma-Aldrich) and 20 ng/ml EGF for 4 weeks. After maturation, spheres
586 were propagated in EF20 medium with mechanical dissociation every fortnight to prevent
587 aggregation and loss of viability. The gliospheres were dissociated into monolayers of APCs using
588 papain dissociation kit (Worthington Biochemical Corporation) and plated onto Matrigel™ (1:80
589 dilution) coated plates. Further, APCs were propagated in EF20 medium until confluent and
590 enzymatically passaged using accutase or cryopreserved.

591 A population of APCs were differentiated into astrocytes for 14 days using astrocyte
592 differentiation media containing Neurobasal medium (Thermo Fisher Scientific, USA), 1% Anti-
593 anti, 1% Glutamax, 1% N2, 0.2% B27, 1% non-essential amino acid medium (NEAA)
594 (ThermoFisher Scientific) and 10 ng/ml ciliary neurotrophic factor (CNTF) (R&D Systems) at
595 37°C, 5% CO₂ and maintained for co-culture experiments or harvest of conditioned medium.

596 **Astrocyte Conditioned Media (ACM)**

597 After 14 days of differentiation of astrocytes, the astrocyte differentiation medium was completely
598 replaced with fresh media at 48 hrs intervals. The cell supernatant was carefully harvested at 48
599 hrs intervals and stored at -20°C. Henceforth, this has been referred to as astrocytes conditioned
600 media (ACM).

601 **Astrocyte and neuron co-culture**

602 Astrocytes were plated on to a 13-mm plastic coverslip (Thermo Fisher Scientific) coated with
603 poly-L-ornithine, laminin, fibronectin (each from Sigma-Aldrich) and Matrigel™ at 35,000
604 cells/coverslip; cultured for 48 hrs at 37°C, 5% CO₂ using CNTF. After 48 hrs NPCs (35,000
605 cells/coverslip) were plated on to astrocytes and maintained in hypoxic conditions i.e., 37°C, 5%
606 CO₂ and 3% O₂ in medium containing Advanced DMEM-F12 supplemented with 1% Anti-anti,
607 0.5% Glutamax, 0.5% N2, 0.2% B27, and 2 µg/ml heparin for 1 week, followed by 10 µM/ml
608 forskolin (R&D Systems) for 2 weeks and maintained in neuronal differentiation medium
609 containing 5 ng/ml each of BDNF and GDNF (R&D Systems) for 5 weeks or up to recording.
610 Immunophenotype and electrophysiological recordings were carried out at 6 - 8 weeks²¹.

611 **Neuron-ACM co-culture**

612 Neurons were differentiated from NPCs as described earlier²¹. Briefly NPCs were plated on to 13-
613 mm glass coverslips (VWR, Radnor, PA) coated with poly-L-ornithine, laminin, fibronectin
614 (Sigma, St. Louis, MO) and MatrigelTM at 35,000 cells/coverslip; maintained in hypoxic
615 environment for 8 weeks. Cells were fed with 50% of neuronal maturation medium and 50% of
616 ACM from 3rd week to 8th week, followed by immunostaining and electrophysiological
617 characterization.

618

619 **METHOD DETAILS**

620 **Immunofluorescence/immunocytochemistry and imaging**

621 Cells were stained using standard immunofluorescence techniques as previously described²¹.
622 Briefly cells were fixed using 4% paraformaldehyde (PFA; Sigma-Aldrich) for 10 minutes, washed
623 three times (5 minutes each) with phosphate buffered saline (PBS; Sigma-Aldrich) 0.1% Tween
624 20 solution (PBS-T; Sigma-Aldrich), followed by permeabilization using 0.3% Triton X-100
625 (USB Corporation) for 10 minutes. Further, cells were washed with PBS-T and blocked with 3%
626 bovine serum albumin (BSA; Sigma-Aldrich) for 1 hour and incubated with primary and secondary
627 antibodies (see Key Resource Table) for 1 hour each at room temperature. The nuclei were
628 counterstained with DAPI (Sigma-Aldrich), and mounted onto glass slides using FluorSaveTM
629 (Merck Millipore). The images were captured at 512 x 512 pixels using a confocal laser scanning
630 microscope (Fluoview 3000 Olympus, Japan), and were acquired using either 40X (1.3 NA) or
631 60X (1.4 NA) oil immersion objectives and diode lasers 405 nm, 488 nm and 561 nm. Z step size
632 was set at 0.5 μ m with 1 airy unit of pinhole diameter.

633 **Western blot and ELISA**

634 Cell pellets were lysed using lysis buffer (50 mM Tris-HCL, pH 7.4, 2 mM EDTA, 0.1% SDS, 1%
635 Triton-X 100, 0.5% Na-deoxycholate, 150 mM NaCl, protease and phosphatase inhibitor
636 cocktails), followed by protein estimation performed using BCA assay kit (ThermoFisher
637 Scientific). 20 μ g/sample protein was loaded on a precast NuPAGE 4-12% Bis-Tris Protein gel
638 (ThermoFisher Scientific) followed by protein transfer onto a 0.45 μ m nitrocellulose membrane
639 (GE Healthcare). Blocking was done using 1:1 TBST (0.2 M Trizma base, 0.15 M NaCl, 0.1%
640 Tween-20) and Odyssey blocking buffer (Li-COR Bioscience, USA) for 1 hr at room temperature.
641 The membranes were incubated with primary antibody FMRP (1:1000, AbCam, Cambridge, UK)
642 and β -actin (1:6000, Merck), in blocking buffer overnight at 4°C. Next day blot was washed with

643 TBST and incubated in secondary antibodies with IRDye 680RD and IRDye 800CW (Li-COR
644 Bioscience, USA) respectively for 1 hr at room temperature. Following further washing, the blot
645 was dried and imaged using Li-COR Odyssey FC infrared system. S100 β levels were measured in
646 ACM using human S100 β DuoSet ELISA (R&D Systems, Minneapolis) in combination with the
647 respective ancillary reagent kit (#DY008, R&D Systems) according to the manufacturer's
648 instructions. Briefly, wells were coated with capture antibody (4 μ g/ml in PBS) overnight at room
649 temperature (RT); followed by washing (0.05% Tween 20 in PBS) and blocked in 1% BSA in
650 PBS. A further 100 μ l of sample or standard in reagent diluent (0.1% BSA, 0.05% Tween 20 in
651 Tris-buffered Saline (TBS)) was added, followed by detection antibody (100 ng/ml in reagent
652 diluent, 2 hours at RT each) and incubated with Streptavidin-HRP B (20 min at RT), followed by
653 washing and addition of substrate solution (20 min at RT). The final reaction was stopped using
654 stop solution and optical density was determined at 450 nm with the correction of 540 nm.

655 **Electrophysiology**

656 Whole-cell patch clamp recordings were performed using similar protocols as described previously
657 ^{21,55,57}. All the recordings were performed at room temperature. Briefly, coverslips containing
658 human cortical neurons and astrocytes co-cultured for 8 weeks in vitro were transferred to the
659 recording chamber perfused with an external recording solution comprising of (in mM): NaCl
660 152, KCl 2.8, HEPES 10, CaCl₂ 2, glucose 10, pH 7.3 – 7.4 (300 – 320 mOsm) with a flow rate of
661 approximately 1.35 ml per minute. Recordings were performed using a MultiClamp 700B
662 amplifier (Molecular Devices, San Jose, CA). Data were sampled at 10 kHz, and digitized, via a
663 Digidata 1550, at 20 kHz. Patch-pipettes, fabricated from thick-walled borosilicate glass, were
664 filled with an internal recording solution containing (in mM): K-gluconate 155, MgCl₂ 2, HEPES
665 10, Na-PiCreatine 10, Mg₂-ATP 2, and Na₃-GTP 0.3, pH 7.3 (280 - 290 mOsm) and had resistances
666 of 3-4 M Ω . Spontaneous network recordings were performed in the current-clamp mode with
667 bridge balance mode and pipette capacitance neutralized. Membrane voltage was held at -70 mV.
668 The duration of action potential bursts was measured from the start of the first action potential to
669 the time of the last action potential with bursts being defined as two or more action potentials
670 occurring during a period of depolarization. Stimulation protocols were generated using pClamp
671 10.5 software and subsequent offline analysis was conducted using Clampfit 11.1 software.

672 S100 β (Sigma-Aldrich) and veratridine (Hello Bio) baseline bursts were recorded for 10
673 min. Veratridine (0.5 μ M) and S100 β (5 μ M) were bath applied and bursts were further recorded
674 from the same cell for 20 min. The last 10 min were taken for analysis. For experiments examining

675 the effects of S100 β antibody application on the bursting profile, an initial baseline (10 min) of
676 activity was rec followed by bath application of S100 β antibody (1:100). Bursts of action potentials
677 were then recorded (for a further 10 min) and their durations and frequencies compared to pre-
678 antibody recordings.

679 To record the persistent sodium current (I_{NaP}), neurons were clamped at -80 mV, then
680 stepped to -100 mV and a depolarizing voltage ramp to -20 mV (20 mV/s) was applied³¹. I_{NaP}
681 was isolated by subtracting the current recorded in the presence of tetrodotoxin (TTX; 1 μ M, Hello
682 Bio) from current recorded immediately prior in the same neuron. S100 β and veratridine were bath
683 applied to assess their effects on the I_{NaP} in the presence of TTX.

684 **Quantification and Statistical Analysis**

685 All values are expressed as mean \pm standard error of the mean (SEM) and each data set was
686 assessed for normality. Students t-tests were used for data sets that passed the normality test, and
687 Mann-Whitney and Wilcoxon tests were used for data sets that failed the normality test. Paired
688 statistical tests were used for data sets before and after drug application. Two –way repeated
689 measures ANOVA, followed by post-hoc Sidak’s test were used for I_{NaP} current – voltage
690 relationships. GraphPad Prism 7 (GraphPad software Inc., La Jolla, CA, RRID: SCR_002798) was
691 used for all the statistical tests. In all the experiments, ‘n’ denotes the number of cells while ‘N’
692 represents number of de novo preparations of batches of culture from which ‘n’ is obtained. For
693 all the pharmacology experiments, only one coverslip per drug application was used.

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