

# 1 Title: The SARS-CoV-2 spike (S) and the orthoreovirus p15 cause neuronal 2 and glial fusion

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## 18    **Abstract**

19    Numerous enveloped viruses use specialized surface molecules called fusogens to enter host  
20    cells<sup>1</sup>. During virus replication, these fusogens decorate the host cells membrane enabling them  
21    the ability to fuse with neighboring cells, forming syncytia that the viruses use to propagate  
22    while evading the immune system. Many of these viruses, including the severe acute respiratory  
23    syndrome coronavirus 2 (SARS-CoV-2), infect the brain and may cause serious neurological  
24    symptoms through mechanisms which remain poorly understood<sup>2-4</sup>. Here we show that  
25    expression of either the SARS-CoV-2 spike (S) protein or p15 protein from the baboon  
26    orthoreovirus is sufficient to induce fusion between interconnected neurons, as well as between  
27    neurons and glial cells. This phenomenon is observed across species, from nematodes to  
28    mammals, including human embryonic stem cells-derived neurons and brain organoids. We  
29    show that fusion events are progressive, can occur between distant neurites, and lead to the  
30    formation of multicellular syncytia. Finally, we reveal that in addition to intracellular molecules,  
31    fusion events allow diffusion and movement of large organelles such as mitochondria between  
32    fused neurons. Our results provide important mechanistic insights into how SARS-CoV-2 and  
33    other viruses could affect the nervous system circuitries causing neurological symptoms.

34

## 35    **Main**

36    Viruses from diverse families, such as rabies virus, herpes simplex virus, dengue virus,  
37    orthoreovirus, and the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), can  
38    infect neurons<sup>3,5</sup>. Viral brain infections are characterized by multiple neurological symptoms,  
39    including headache, fever, confusion, epileptic seizures, and loss of taste or smell. In more  
40    severe cases, viral brain infections can lead to encephalitis and meningitis, and potentially  
41    irreversible neuronal deficits such as paralysis and death. Most clinical symptoms originate from

42 the death of infected neurons <sup>4</sup>. However, some viruses do not kill their host cells, and the  
43 chronic neurological sequelae of these infections cannot be explained by the loss of infected  
44 neurons <sup>2</sup>. Other neuropathological mechanisms must therefore underlie the progression of these  
45 viral infections leading to brain dysfunction. In non-neuronal tissues, viruses use their fusogens  
46 to fuse with host membranes and enter cells <sup>1</sup>. Once expressed inside the host cell, the viral  
47 fusogens redecorate the cell membrane conferring the ability to fuse with neighboring cells. This  
48 results in the formation of multinucleated syncytia, which allow viral propagation ‘from within’,  
49 without the need for virion release into the extracellular space <sup>6,7</sup>. As defined over 100 years ago  
50 by Ramon y Cajal, neurons are individual units that do not base their development or  
51 communication on cellular fusion, with the preservation of their individuality being critical for  
52 the correct function of the nervous system. It is currently unknown whether the presence of viral  
53 fusogens can cause neuronal fusion and the formation of syncytia, thereby altering the neuronal  
54 circuitry and function.

55 To address this question, we began by using the fusion-associated small transmembrane (FAST)  
56 fusogen p15, isolated from the baboon orthoreovirus (BRV), which infects the brain of these  
57 primates and causes meningoencephalomyelitis <sup>8,9</sup>. p15 is the only viral protein required by the  
58 BRV to form a syncytium <sup>10</sup>, with no receptor protein on the host cell being needed to facilitate  
59 fusion <sup>11,12</sup>. To safely mimic the result of viral neuronal infection, we transfected p15 in  
60 embryonic mouse primary hippocampal neurons, and visualized the presence of fusion through  
61 the transfer of different intracellular fluorophores between neurons. Immediately after isolation,  
62 a population of neurons was co-transfected by electroporation with a plasmid containing p15 and  
63 another containing GFP; a second population of neurons was co-transfected with a plasmid  
64 containing mCherry and the empty control vector. The two neuronal populations were then  
65 plated together and maintained in culture for 7 days (7 DIV). Our results revealed that the

66 expression of p15 was sufficient to induce neuronal fusion, as detected by the presence of  
67 neurons containing both the GFP and mCherry fluorophores (Fig. 1A first row, and B), a  
68 phenotype that was never observed when the control vector was co-transfected in the absence of  
69 p15 (Fig. 1a third row, and b). To determine whether the fluorophore diffusion was caused by  
70 the fusogenic properties of p15, we generated an inactive version of this fusogen, p15 $\Delta$ 21-22, in  
71 which two residues of the N-terminal of the transmembrane domain were truncated<sup>13</sup>.  
72 Importantly, expression of this inactive fusogen completely abolished the fusion of neurons (Fig.  
73 1a second row, and b).  
74 Neuronal fusion implies a temporary or permanent diffusion of cytoplasmic material between  
75 cells<sup>14</sup>. To confirm that this was the case, we co-transfected neurons with p15 and the  
76 photoconvertible fluorescent protein Kaede, which shifts from green to red fluorescence upon  
77 illumination with ultraviolet (UV) light (350 – 400 nm). After identification of interconnected  
78 adjacent green fluorescent neurons, we photoconverted the green Kaede fluorophore by applying  
79 brief pulses of UV light in a small region of one neuron (donor). Newly generated red  
80 photoconverted Kaede molecules rapidly diffused to the adjacent neuron (acceptor) (Extended  
81 Data Fig. 1a and Supplementary video 1). The diffusion of red Kaede was measured as a  
82 decrease in the red fluorescence within the donor neurons (Extended Data Fig. 1b), with a  
83 concomitant increase in the red fluorescence in the acceptor neurons (Extended Data Figs. 1c, d),  
84 thereby conclusively demonstrating the existence of an active cytoplasmic bridge between p15-  
85 fused neurons. In the absence of fusion, red Kaede remained confined within the photoconverted  
86 neuron (Extended Data Fig. 1a, and 1e-g). We then asked whether the fusion bridges allowed  
87 exchange of cellular components larger than fluorescent proteins. To address this question, we  
88 used the photo-activatable mitochondrial marker mito-mPA-GFP, which allows to visualize

89 mitochondria movement. Indeed, in fused neurons, we observed diffusion of mitochondria from  
90 donor to acceptor neurons (Extended Data Fig. 2, Supplementary Video 2).

91 We next asked whether the neuronal fusion induced by p15 was restricted to two adjacent  
92 neurons, or if it was a propagating event that generated syncytia with a larger number of  
93 interconnected neurons. To address this possibility, we co-transfected neurons with p15 and  
94 GFP and monitored the appearance of syncytia over a period of 7 days. The number of neuronal  
95 syncytia increased over time, appearing as clusters of interconnected GFP-positive neurons that  
96 progressively incorporated more cells (Fig. 1c-e). Moreover, through live confocal imaging  
97 performed over a period of 50 hours, we observed the progressive appearance of GFP in  
98 surrounding neurons, revealing the occurrence of fusion events (Supplementary Video 3).

99 Fusion was observed between the somas of adjacent neurons as well as between the processes  
100 (*i.e.* dendrites and axons) of distant neurons, resulting in fusion bridges with variable lengths that  
101 could extend over hundreds of microns (Extended Data Fig. 3).

102 To determine if the p15-mediated neuronal fusion capacity was also conserved *in vivo*, we  
103 generated transgenic *Caenorhabditis elegans* strains in which p15 and GFP were expressed  
104 simultaneously under the control of the *mec-4* promoter (*Pmec-4::p15* and *Pmec-4::GFP*), which  
105 is active in the six mechanosensory neurons (ALM left and right, AVM, PVM and PLM left and  
106 right; Fig. 2a panel i and v). Similar to what was observed in mammalian neurons in culture, any  
107 fusion event with nearby neurons or tissues would be detectable by diffusion of GFP from the  
108 mechanosensory neurons to other cells. As predicted, we observed the appearance of additional  
109 GFP-positive cells in the head, mid body and tail of the animal, a phenomenon never observed in  
110 non-transgenic siblings or wild-type animals (Fig. 2a, b). Based on its stereotypic location and  
111 morphology, we identified ALN as the most prevalent additional GFP-positive neurons. ALN  
112 are a pair of sensory neurons located in the tail of the animal that extend their axons in close

113 association with the axons of the ipsilateral ALM mechanosensory neurons <sup>15</sup>. Other frequently  
114 GFP-positive neurons were the pair of interneurons LUA, which are located in the tail of the  
115 animal and extend anterior neurites in close proximity with those of the PLM neurons, and the  
116 mechanosensory neurons PVD, positioned in the mid-body of the animal with two extensively  
117 branched dendrites and a long axon <sup>15</sup>. Despite over 90% of the GFP-positive cells being  
118 neurons, we also identified fluorescence in hypodermal cells, which form a tissue in which the  
119 PLM and ALM axons are embedded and has a glial-like function <sup>16,17</sup> (Fig. 2a, panel iv). Similar  
120 to mammalian neurons, the expression of the inactive fusogen p15 $\Delta$ 21-22 within  
121 mechanosensory neurons did not result in fusion with neurons or hypodermal cells (Fig. 2b).  
122 Taken together, these results indicate that the *in vivo* expression of p15 can drive the fusing of  
123 neurons with other neurons or glial cells that are located in close proximity, suggesting a possible  
124 pathomechanism of neuronal malfunction caused by orthoreovirus infection.  
125 Orthoreoviruses can be transmitted from bats to humans <sup>18</sup>, and SARS-CoV-2 virus is derived  
126 from bat coronaviruses <sup>19</sup>. SARS-CoV-2 primarily causes a respiratory illness, but increasing  
127 evidence reveal also brain infection <sup>20-22</sup>. Unlike p15, the spike S protein that functions as the  
128 fusogen of the SARS-CoV-2 virus must bind to a receptor protein, the human angiotensin-  
129 converting enzyme 2 (hACE2), located on the surface of the host cell <sup>23</sup>, and uses neuropilin-1 as  
130 a co-factor to enhance infectivity <sup>24</sup>. Among other tissues, hACE2 is expressed in neuronal and  
131 glial cells in the human central nervous system <sup>25</sup>. Mouse neurons express mACE2 <sup>26</sup>, which  
132 shares 81.86% interspecies homology with the human protein but lacks key residues for spike S  
133 binding <sup>23</sup>. For this reason, to study the neuronal fusion properties of the spike S protein, both  
134 spike S and hACE2 must be expressed in murine neurons. Using a similar approach as that  
135 described for p15, we independently electroporated two neuronal populations, one with a  
136 plasmid expressing GFP plus a plasmid containing a codon-optimized spike S protein, and the

137 other with a plasmid expressing mCherry plus a plasmid containing the hACE2 receptor. The  
138 two populations were then plated together and cultured for 7 days. The expression of the  
139 fusogen spike S and its receptor hACE2 in adjacent cells resulted in the fusion of these neurons  
140 and the mixture of the fluorophores (Fig. 3a first row, and b). Unlike p15, the presence of the  
141 specific receptor was required to initiate cellular fusion, as expression of spike S or hACE2 alone  
142 did not generate any fusion events (Fig. 3a second and third rows, and b). To determine whether  
143 the fusion of neurons was caused by the fusogenic properties of spike S, we used two fusion-  
144 inactive versions of this protein, the spike S-2P and the spike S-6P (HexaPro). First, we  
145 generated the spike S-2P variant containing two consecutive proline substitutions in the C-  
146 terminal S2 subunit<sup>27</sup>. These two mutations retain the spike S in a prefusion conformation,  
147 blocking its fusion capacity<sup>28</sup>. Interestingly, this mutant form has been chosen for the creation  
148 of most available vaccines against SARS-CoV-2<sup>29</sup>. The spike S-6P contains four additional  
149 proline substitutions (F817P, A892P, A899P, A942P) that further stabilize the prefusion  
150 conformation, and increase protein expression and ability to withstand heat stress<sup>30</sup>. Our results  
151 indicate that none of these two versions of spike S induce neuronal fusion (Fig. 3c and d). In  
152 addition to neuron-neuron fusion, we observed neuron-glia and glia-glia fusion events if fusogen  
153 and receptor were expressed in these cell types (Extended Data Fig. 4). Neurons could fuse  
154 through their soma or via contacting neurites, forming fusion bridges of lengths that could reach  
155 over 100 microns (Extended Data Fig. 5). The functionality of these fusion bridges was  
156 evaluated using the exchange of photoconverted red Kaede fluorophores between fused neurons.  
157 Similar to what we observed with p15, photoconversion of Kaede in a small region of one fused  
158 neuron (donor) resulted in the fast diffusion of red Kaede molecules towards the other fused  
159 neuron (acceptor) (Extended Data Fig. 6). These results demonstrate that fusion through spike S  
160 and hACE2 also generates functional bridges that allow the diffusion of proteins. Importantly,

161 the fused neurons retained their morphology, extended processes, and remained viable for the  
162 duration of the experiment.

163 Finally, we tested whether p15 and spike S have the potential to cause neuronal fusion in human  
164 embryonic stem cell (hESC)-derived neurons and brain organoids. Human neurons and  
165 organoids were transfected with a fluorophore (mCherry or GFP) with either p15, spike S or the  
166 inactive spike S-6P, and compared with their respective controls expressing fluorescence  
167 markers only. Remarkably, we observed that both p15 and full-length spike S fusogens caused  
168 extensive neuronal fusion in both human-derived systems (Fig. 4 and Extended Data Fig. 7).

169 Importantly, the spike S-6P mutant also failed to induce cell fusion in cultured human neurons  
170 (Fig. 4b and c) and brain organoids (Fig. 4e).

171 Fused neurons can result in compromised neuronal circuitry and altered animal behavior, as  
172 previously shown for *C. elegans* chemosensory neurons that ectopically express endogenous  
173 fusogens<sup>31</sup>. Our results demonstrate that neurons expressing viral fusogens acquire the capacity  
174 to fuse, potentially compromising their functional circuit properties while remaining viable. This  
175 previously uncharacterized event could explain at least some, if not most, of the neurological  
176 consequences of viral infections of the nervous system. Moreover, most of the current  
177 immunization approaches for COVID-19 are based on expressing the spike S protein in the host  
178 cells as an epitope to trigger the immune system response<sup>29</sup>. These nucleic acid-based vaccines  
179 deliver the antigen encoded as mRNA, such as the Pfizer-BioNTech BNT162b2 and the  
180 Moderna mRNA-1273 vaccines<sup>32</sup>, or as adenovirus-enclosed DNA, such as the Oxford-  
181 AstraZeneca ChAdOx1 nCoV-19/AZD1222<sup>33</sup> and Johnson & Johnson Ad26.COV2.S<sup>28</sup>  
182 vaccines. The current versions of the Moderna, Pfizer-BioNTech and Johnson & Johnson  
183 vaccines encode the full-length spike S (spike S-2P) with two mutations that stabilize the pre-  
184 fusion conformation and inactivate its fusogenicity<sup>28,34,35</sup>. Our findings demonstrate that it is

185 critical to consider the fusogenic potential when designing any future vaccines in which viral  
186 fusogens are to be expressed in mammalian cells.

187

## 188 **Methods**

189 *Molecular biology*

190 Standard molecular biology methods were used. The p15 DNA sequence was obtained from the  
191 National Center for Biotechnology Information (<http://www.ncbi.nlm.nih.gov/gene/>). The  
192 plasmid was then designed using the software ‘A Plasmid Editor’ and the insert was generated  
193 by Integrated DNA Technologies. The *Pmec-4::p15* plasmid was constructed by subcloning p15  
194 between Msc I and Nhe I. The *CMV::p15* plasmid was generated by subcloning *p15* into the  
195 pmaxCloning<sup>TM</sup> vector (Lonza, # VDC-1040) between Hind III and Not I. The  
196 *CMV::p15Δ21/22* plasmid was generated as previously described<sup>13</sup> by deletion of the amino  
197 acids 21 and 22 of the N-terminus of the transmembrane domain. The *CMV::SARS-CoV-2-S-2P*  
198 plasmid was generated by introducing two prolines at the 986 (K986P) and the 987 (V987P)  
199 positions of the *SARS-CoV-2-S* gene of the pCMV14-3X-Flag-SARS-CoV-2 S plasmid.  
200 Mutations were done using the QuikChangeII Site-Directed Mutagenesis Kit (*p15Δ21/22*  
201 forward primer 5'-CCACCGCCAAATGCTTTGTTGAAAGCAGTTCTACTG-3'; *p15Δ21/22*  
202 reverse primer 5'-CAGTAGAACTGCTTCAACAAAAGCATTGGCGGTGG-3'; *SARS-CoV-*  
203 *2-S-2P* forward plasmid 5'-CCTGAGTCGCCTGATCCGCCGGAAGCTGAAGTTC-3';  
204 *SARS-CoV-2-S-2P* reverse plasmid 5'-  
205 GAACTTCAGCTTCCGGCGGATCAAGGCGACTCAGG-3'). Positive clones were confirmed  
206 by Sanger sequencing. The Kaede-N1 plasmid was a gift from Michael Davidson (Addgene  
207 plasmid # 54726; <http://n2t.net/addgene:54726>; RRID:Addgene\_54726)<sup>36</sup>. The mito-mPA-GFP  
208 plasmid was a gift from Richard Youle (Addgene plasmid # 23348;

209 http://n2t.net/addgene:23348; RRID:Addgene\_23348) <sup>37</sup>. The pCMV14-3X-Flag-SARS-CoV-2  
210 S plasmid was a gift from Zhaohui Qian (Addgene plasmid # 145780;  
211 http://n2t.net/addgene:145780; RRID:Addgene\_145780) <sup>38</sup>. The pcDNA3.1-SARS2-Spike  
212 plasmid was a gift from Fang Li (Addgene plasmid # 145032 ; http://n2t.net/addgene:145032 ;  
213 RRID:Addgene\_145032) <sup>39</sup>. The SARS-CoV-2 S-6P plasmid was a gift from Jason McLellan  
214 (Addgene plasmid # 154754 ; http://n2t.net/addgene:154754 ; RRID:Addgene\_154754) <sup>30</sup>. The  
215 pcDNA3.1-hACE2 plasmid was a gift from Fang Li (Addgene plasmid # 145033;  
216 http://n2t.net/addgene:145033; RRID:Addgene\_145033) <sup>39</sup>.

217

218 *Animal ethics and mouse strains*

219 All experimental procedures using animals were conducted under the guidelines of the  
220 Australian Code of Practice for the Care and Use of Animals for Scientific purposes, and were  
221 approved by the University of Queensland Animal Ethics Committee (2019/AE000243).

222 Wildtype (WT, C57Bl/6 background) mice were maintained on a 12 h light/dark cycle and  
223 housed in a PC2 facility with ad libitum access to food and water.

224

225 *Murine neuronal culture*

226 Primary hippocampal neurons were obtained from mice at embryonic day E16. Isolated  
227 hippocampi were prepared as previously described <sup>40,41</sup>. Briefly, 250,000 neurons were plated  
228 onto poly-L-lysine-coated 35 mm glass-bottom dishes (In Vitro Scientific) in Neurobasal  
229 medium (Gibco) supplemented with 5% fetal bovine serum (Hyclone), 2% B27, 2 mM  
230 GlutaMAX and 50 U/mL penicillin/streptomycin (Invitrogen). The medium was changed to  
231 serum-free/antibiotic-free medium 24 h post-seeding, and half the medium was changed every  
232 week.

233 *Human embryonic stem cell-derived neurons*

234 Human embryonic stem cell (hESC)-derived cortical neurons were differentiated from hESCs  
235 (H9, WIC-WA09-MB-001; WiCell) using a modification of a previously published protocol <sup>42</sup>.  
236 Briefly, the hESCs were maintained on Matrigel (#354230; Corning) in Essential 8 medium  
237 (#A1517001; Life Technologies). hESCs were dissociated with Accutase (#AT-104-500;  
238 Innovate Cell Technologies) at 37 °C for 1 min and seeded on AggreWell 800 (#34815;  
239 StemCell Technologies) in Essential 8 medium with ROCK inhibitor Y-27632 (10 µM, #72308,  
240 STEMCELL Technologies). After 24 h, spheroids were collected and transferred into ultra-low-  
241 attachment plates (#CLS3471; Sigma-Aldrich) with Essential 6 medium (#A1516401; Life  
242 Technologies) containing dorsomorphin (2.5 µM, #P5499, Sigma-Aldrich), SB-431542 (10 µM,  
243 #1614, Tocris) and cultured until day 13. The spheroids were then collected and placed into  
244 Matrigel (Corning)-coated 6 well-plates with DMEM/F12 (#11330057; Thermo Fisher  
245 Scientific), supplemented with 1X N2 (#17502048; Thermo Fisher Scientific). The produced  
246 neural progenitor cells (NPCs) were passaged until day 30, and then replated on poly-L-lysine  
247 (10 µg/ml #P2636; Sigma-Aldrich) and laminin (20 µg/ml #23017015; Thermo Fisher  
248 Scientific)-coated plates and maintained in Neurobasal medium (#21103049; Thermo Fisher  
249 Scientific) supplemented with B-27 (Thermo Fisher Scientific), Glutamax (#35050061; Thermo  
250 Fisher Scientific), brain-derived neurotrophic factor (BDNF, #78133; StemCell Technologies),  
251 ascorbic acid (200 µM, #72132; StemCell Technologies), GDNF (20 ng/ml, #78139.1, StemCell  
252 Technologies) and cAMP (1 µM, #A6885; Sigma-Aldrich) for at least 2 weeks.

253

254 *Brain organoids*

255 hESC-derived brain organoids were produced as previously reported <sup>43</sup>. Briefly, hESCs were  
256 incubated with Accutase (#AT-104-500; Innovate Cell Technologies) at 37 °C for 1 min.

257 Dissociated single cells were seeded on AggreWell 800 (#34815, StemCell Technologies). 1000  
258 single cells were added per well in Essential 8 medium supplemented with 10µM ROCK  
259 inhibitor Y-27632, and centrifuged at 100 xg for 3 min. Cultures were incubated at 37 °C and  
260 5% CO<sub>2</sub>. After 24 h, the organoids were transferred into ultra-low-attachment 60φ dishes  
261 (#CLS3261, Sigma-Aldrich) in Essential 6 medium supplemented with 2.5 µM dorsomorphin, 10  
262 µM SB-431542 and 2.5 µM XAV-939 (#3748, Tocris). The medium was changed every day for  
263 5 days. On day 6, the medium was changed into Neurobasal A (#10888; Thermo Fisher  
264 Scientific) containing GlutaMax (#35050; Thermo Fisher Scientific) and B-27 without vitamin A  
265 (#12587; Thermo Fisher Scientific). The medium was supplemented with 20 ng/ml epidermal  
266 growth factor (EGF, #236-EG; R&D Systems) and 20 ng/ml basic fibroblast growth factor  
267 (bFGF, # 233-FB; R&D Systems) until day 24, when it was switched to Neurobasal A medium  
268 containing GlutaMax, B-27 without vitamin A, and supplemented with 20 ng/ml BDNF and 20  
269 ng/ml Neurotrophin-3 (NT3, #78074; STEMCELL Technologies) until day 43. Finally, the  
270 medium was changed to Neurobasal A containing GlutaMax and B-27 without vitamin A.

271

272 *Electroporation and transfection*

273 When required, murine neurons were electroporated using the Invitrogen Neon<sup>TM</sup> transfection  
274 system (#MPK1025, ThermoFisher Scientific) following the manufacturer's instructions.  
275 Briefly, immediately after dissection, isolated neurons were washed twice in Ca<sup>2+</sup>-Mg<sup>2+</sup>-free PBS  
276 and resuspended in buffer R containing 1 to 2 µg of the DNA to electroporate. The conditions for  
277 the electroporation were voltage 1500 V, width 10 ms, and 3 pulses. Alternatively, the neurons  
278 were transfected after 7-12 DIV using the Lipofectamine 2000 (#11668027, Invitrogen) reagent  
279 as previously described<sup>40</sup>. Following the manufacturer's instructions, human neurons were  
280 transfected with Lipofectamine 3000 (#L3000015, Thermo Fisher Scientific) and 1.6 µg of

281 plasmid DNA at 40 - 50 DIV; human brain organoids were transfected with Lipofectamine 3000  
282 and 2.4 µg of plasmid DNA at 43 - 50 DIV. The transfection medium was replaced after 24 h.

283

284 *Caenorhabditis elegans* strain maintenance, crosses and manipulation

285 Standard techniques were used for *C. elegans* maintenance, genetic crosses and manipulations<sup>44</sup>.  
286 Experiments were performed on hermaphrodite animals grown at room temperature (~22 °C) on  
287 nematode growth medium plates, seeded with OP50 *Escherichia coli*. Previously generated  
288 transgenes used in this study include: *vdEx1266* [*Pmec-4::p15* 5ng/µl]; *vdEx1268* [*Pmec-4::p15*  
289 5ng/µl]. Transgenic strains generated during this study were obtained by standard  
290 microinjection techniques<sup>45</sup>, and include: *vdEx1266* [*Pmec-4::p15* 5 ng/µl]; *vdEx1268* [*Pmec-*  
291 *4::p15* 5 ng/µl]; *vdEx1417* [*Pmec-4::p15Δ21/22*; 15 ng/µl]; *vdEx1487* [*Pmec-4::p15Δ21/22*; 5  
292 ng/µl]; *vdEx1489* [*Pmec-4::p15Δ21/22*; 5 ng/µl]. All injection mixes had a total concentration  
293 of 100 ng/µl and contained the transgene of interest, empty pSM plasmid as a filler, and a co-  
294 injection marker for the identification of transgenic animals. As the cell-cycle transfer of  
295 extrachromosomal arrays (*vdEx*) is unstable, some animals lose the transgene. This provides an  
296 internal control for each transgenic strain, with these controls being referred to as ‘non-  
297 transgenic siblings’ or *transgene (-)*.

298

299 *Immunofluorescence staining*

300 Murine neurons were fixed with a solution of 4% paraformaldehyde (PFA) in PBS for 10 min.  
301 Neurons were rinsed in PBS, and permeabilized with a solution of 0.1% Triton-X-100 in PBS for  
302 10 min. They were then washed with PBS and incubated in blocking solution (5% horse serum,  
303 1% BSA in PBS) for 1 h at room temperature. After blocking, the neurons were incubated with  
304 the primary antibodies to GFP (#AB16901, Merck Millipore), mCherry (#ab167453, Abcam)

305 and anti-MAP2 (#188004, Synaptic Systems), diluted in primary antibody solution (1% BSA in  
306 PBS) overnight at 4 °C. They were then washed with PBS and incubated with the secondary  
307 antibodies Alexa Fluor 488 goat anti-chicken (#A-11039, ThermoFisher Scientific), Alexa Fluor  
308 555 goat anti-rabbit (#A32732, ThermoFisher Scientific), Alexa Fluor 647 goat anti-guinea pig  
309 (#A21450, ThermoFisher Scientific) and DAPI (#62248, ThermoFisher Scientific) in the  
310 secondary antibody solution (5% horse serum in PBS) for 1 h at room temperature. Finally, the  
311 neurons were washed and mounted in Vectashield® Plus antifade mounting medium (#H-2000,  
312 Vector Laboratories). The staining protocol was slightly modified for human neurons, with  
313 fixing for 5 min, blocking with 5% (w/v) BSA (#A9418, Sigma-Aldrich) and staining for MAP2  
314 only. Organoids were fixed with 4% PFA at room temperature or 30 min at room temperature.  
315 PBS-rinsed organoids were permeabilized with 0.5% (v/v) Triton X-100 (#X100; Sigma-  
316 Aldrich) in PBS for 30 min, followed by blocking in 5% (w/v) BSA (#A7906; Sigma-Aldrich)  
317 for 5 h. The organoids were then incubated with primary antibodies to MAP2 (#ab5392, Abcam)  
318 at 4 °C for 3 days. Following PBS washing, they were incubated with Alexa Fluor-conjugated  
319 secondary antibodies (#A11039 or #A21437, Thermo Fisher Scientific) at 4 °C for 3 days.  
320 Nuclei were counterstained with DAPI (#D1306; Invitrogen) for 30 min and washed with PBS  
321 before imaging.

322

323 *Confocal microscopy of fixed samples*

324 The imaging for immunofluorescence microscopy was carried out on either a Zeiss Plan  
325 Apochromat 40x/1.2 NA water-immersion objective on a confocal/two-photon laser-scanning  
326 microscope (LSM 710; Carl Zeiss) built around an Axio Observer Z1 body (Carl Zeiss),  
327 equipped with two internal gallium arsenide phosphide (GaAsP) photomultiplier tubes (PMTs)  
328 and three normal PMTs for epi- (descanned) detection, or using a Zeiss Plan Apochromat

329 40x/1.2 NA water-immersion objective on a confocal/two-photon laser-scanning microscope  
330 (LSM 710; Carl Zeiss) and confocal microscope (LSM 880, Carl Zeiss) built around an Axio  
331 Observer Z1 body (Carl Zeiss), equipped with two internal gallium arsenide phosphide (GaAsP)  
332 photomultiplier tubes (PMTs) and three normal PMTs for epi- (descanned) detection. Both  
333 systems were controlled by Zeiss Zen Black software. Images were further processed and  
334 analyzed with FIJI-ImageJ<sup>46</sup>.

335

336 *Live imaging*

337 *C. elegans* animals were immobilized using 0.05% tetramisole hydrochloride on 3.5% agar pads.  
338 The animals were imaged with a Zeiss Axio Imager Z1 microscope equipped with a  
339 Photometrics camera (Cool Snap HQ2), and analyzed using Metamorph software (Molecular  
340 Devices) and FIJI-ImageJ. Cytoplasmic GFP was visualized with 470/80 nm excitation and  
341 525/50 nm emission filters. Image acquisition was performed using SlideBook 6.0 (3I, Inc) and  
342 processed using FIJI-ImageJ.

343 For live-cell imaging of Kaede diffusion in mammalian neurons, fusogen/empty vector-Kaede-  
344 transfected neurons were bathed in imaging buffer (145 mM NaCl, 5.6 mM KCl, 2.2 mM CaCl<sub>2</sub>,  
345 0.5 mM MgCl<sub>2</sub>, 5.6 mM D-glucose, 0.5 mM ascorbic acid, 0.1% BSA, 15 mM HEPES, pH 7.4).

346 Neurons were visualized at 37 °C on a Zeiss Plan Apochromat 40x/1.2 NA water-immersion  
347 objective on a confocal/two-photon laser-scanning microscope (LSM 710; Carl Zeiss). For  
348 transfected neurons, a UV pulse was applied on a 5 μm x 5 μm region of interest (ROI).

349 Photoconversion resulted in the emergence of a spot of red fluorescence that rapidly diffused  
350 through the soma and proximal dendrites of the neurons. Simultaneous green and red images  
351 were collected every 785 ms; 5 images were acquired before photoconversion and 50 images  
352 after photoconversion. Photoconversion and diffusion of the fluorophore were performed on

353 neurons separated by 25  $\mu\text{m}$  to 100  $\mu\text{m}$ . In control conditions, UV light was applied first outside  
354 the neuron, 50  $\mu\text{m}$  away from the soma, and then inside the soma.

355 Similarly, for live-cell imaging of mito-mPA-GFP diffusion on mammalian neurons, a UV pulse  
356 was applied on a 10  $\mu\text{m}$  x 10  $\mu\text{m}$  ROI. Photoconversion resulted in the emergence of green  
357 mitochondria that slowly moved from one neuron to the other along the fusion bridge. Green  
358 images were collected every 5 min; 72 images were acquired before photoconversion and one  
359 last image was acquired 13 h later.

360

361 *Neuron-neuron, neuron-glia and glia-glia fusion quantification in mammalian neuronal cultures*  
362 Cell-cell fusion was quantified as the percentage of pair of transfected neurons with their somas  
363 within a radius  $\leq 200 \mu\text{m}$ , and that contain simultaneously GFP and mCherry. Neuron-neuron  
364 fusion is identified by two fused cells positive for MAP2 staining; neuron-glia fusion is  
365 identified by one of the two fused cells is positive for MAP2; glia-glia fusion is identified by  
366 none of the fused cells positive for MAP2. Neuronal syncytia were quantified as the percentage  
367 of interconnected neurons within a distance  $\leq 200 \mu\text{m}$ .

368

369 *Statistical analysis*

370 Results were analyzed statistically using GraphPadPrism software (GraphPad Software, Inc).  
371 The D'Agostino and Pearson test was used to test for normality. The unpaired two-tailed non-  
372 parametric Mann-Whitney U test was used for comparison of two groups, when the data were  
373 not normally distributed. For datasets comparing more than two groups, we performed one-way  
374 ANOVA Kruskal-Wallis test followed by Dunn's *post hoc* test for multiple comparisons, one-  
375 way ANOVA Brown-Forsythe and Welch tests followed by the Games-Howell's *post hoc* test  
376 for multiple comparisons, or two-way ANOVA followed by Geisser-Greenhouse correction and

377 the Šidák *post hoc* test for multiple comparisons. Statistical comparisons were performed on a  
378 per-dish or a per-neuron basis. 2-4 technical replicate dishes were imaged and 2-3 independent  
379 cultures were used per condition. Each mouse dissection provided neurons from a pool of at  
380 least 6 different embryos. Values are represented as the mean  $\pm$  SD or mean  $\pm$  SEM. The tests  
381 used are indicated in the respective figure legends. A *p*-value below 0.05 was accepted as  
382 significant.

383

384 **References**

- 385 1 Martens, S. & McMahon, H. T. Mechanisms of membrane fusion: disparate players and  
386 common principles. *Nat Rev Mol Cell Biol* **9**, 543-556, doi:10.1038/nrm2417 (2008).
- 387 2 van den Pol, A. N. Viral infection leading to brain dysfunction: more prevalent than  
388 appreciated? *Neuron* **64**, 17-20, doi:10.1016/j.neuron.2009.09.023 (2009).
- 389 3 Song, E. *et al.* Neuroinvasion of SARS-CoV-2 in human and mouse brain. *J Exp Med*  
390 **218**, doi:10.1084/jem.20202135 (2021).
- 391 4 Griffin, D. E. & Hardwick, J. M. Perspective: virus infections and the death of neurons.  
392 *Trends Microbiol* **7**, 155-160, doi:10.1016/s0966-842x(99)01470-5 (1999).
- 393 5 John, C. C. *et al.* Global research priorities for infections that affect the nervous system.  
394 *Nature* **527**, S178-186, doi:10.1038/nature16033 (2015).
- 395 6 Compton, A. A. & Schwartz, O. They might be giants: does syncytium formation sink or  
396 spread HIV infection? *PLoS Pathog* **13**, e1006099, doi:10.1371/journal.ppat.1006099  
397 (2017).
- 398 7 Frankel, S. S. *et al.* Replication of HIV-1 in dendritic cell-derived syncytia at the mucosal  
399 surface of the adenoid. *Science* **272**, 115-117, doi:10.1126/science.272.5258.115 (1996).

400 8 Duncan, R., Murphy, F. A. & Mirkovic, R. R. Characterization of a novel syncytium-  
401 inducing baboon reovirus. *Virology* **212**, 752-756, doi:10.1006/viro.1995.1536 (1995).

402 9 Kumar, S. *et al.* Reovirus-associated meningoencephalomyelitis in baboons. *Vet Pathol*  
403 **51**, 641-650, doi:10.1177/0300985813497487 (2014).

404 10 Dawe, S. & Duncan, R. The S4 genome segment of baboon reovirus is bicistronic and  
405 encodes a novel fusion-associated small transmembrane protein. *J Virol* **76**, 2131-2140,  
406 doi:10.1128/jvi.76.5.2131-2140.2002 (2002).

407 11 Chan, K. M. C. *et al.* Evolutionarily related small viral fusogens hijack distinct but  
408 modular actin nucleation pathways to drive cell-cell fusion. *Proc Natl Acad Sci U S A*  
409 **118**, doi:10.1073/pnas.2007526118 (2021).

410 12 Top, D., Read, J. A., Dawe, S. J., Syvitski, R. T. & Duncan, R. Cell-cell membrane  
411 fusion induced by p15 fusion-associated small transmembrane (FAST) protein requires a  
412 novel fusion peptide motif containing a myristoylated polyproline type II helix. *J Biol  
413 Chem* **287**, 3403-3414, doi:10.1074/jbc.M111.305268 (2012).

414 13 Clancy, E. K. & Duncan, R. Helix-destabilizing, beta-branched, and polar residues in the  
415 baboon reovirus p15 transmembrane domain influence the modularity of FAST proteins.  
416 *J Virol* **85**, 4707-4719, doi:10.1128/JVI.02223-10 (2011).

417 14 Neumann, B., Nguyen, K. C., Hall, D. H., Ben-Yakar, A. & Hilliard, M. A. Axonal  
418 regeneration proceeds through specific axonal fusion in transected *C. elegans* neurons.  
419 *Dev Dyn* **240**, 1365-1372, doi:10.1002/dvdy.22606 (2011).

420 15 Altun, Z. F., Chen, B., Wang, Z. W. & Hall, D. H. High resolution map of  
421 *Caenorhabditis elegans* gap junction proteins. *Dev Dyn* **244**, 903,  
422 doi:10.1002/dvdy.24287 (2015).

423 16 Chalfie, M. *et al.* The neural circuit for touch sensitivity in *Caenorhabditis elegans*. *J  
424 Neurosci* **5**, 956-964 (1985).

425 17 Coakley, S., Ritchie, F. K., Galbraith, K. M. & Hilliard, M. A. Epidermal control of  
426 axonal attachment via beta-spectrin and the GTPase-activating protein TBC-10 prevents  
427 axonal degeneration. *Nat Commun* **11**, 133, doi:10.1038/s41467-019-13795-x (2020).

428 18 Wang, L. F. & Anderson, D. E. Viruses in bats and potential spillover to animals and  
429 humans. *Curr Opin Virol* **34**, 79-89, doi:10.1016/j.coviro.2018.12.007 (2019).

430 19 Burki, T. The origin of SARS-CoV-2. *Lancet Infect Dis* **20**, 1018-1019,  
431 doi:10.1016/S1473-3099(20)30641-1 (2020).

432 20 Meinhardt, J. *et al.* Olfactory transmucosal SARS-CoV-2 invasion as a port of central  
433 nervous system entry in individuals with COVID-19. *Nat Neurosci* **24**, 168-175,  
434 doi:10.1038/s41593-020-00758-5 (2021).

435 21 Kase, Y. & Okano, H. Neurological pathogenesis of SARS-CoV-2 (COVID-19): from  
436 virological features to clinical symptoms. *Inflamm Regen* **41**, 15, doi:10.1186/s41232-  
437 021-00165-8 (2021).

438 22 Zhang, B. Z. *et al.* SARS-CoV-2 infects human neural progenitor cells and brain  
439 organoids. *Cell Res* **30**, 928-931, doi:10.1038/s41422-020-0390-x (2020).

440 23 Yan, R. *et al.* Structural basis for the recognition of SARS-CoV-2 by full-length human  
441 ACE2. *Science* **367**, 1444-1448, doi:10.1126/science.abb2762 (2020).

442 24 Cantuti-Castelvetro, L. *et al.* Neuropilin-1 facilitates SARS-CoV-2 cell entry and  
443 infectivity. *Science* **370**, 856-860, doi:10.1126/science.abd2985 (2020).

444 25 Khan, S. & Gomes, J. Neuropathogenesis of SARS-CoV-2 infection. *eLife* **9**,  
445 doi:10.7554/eLife.59136 (2020).

446 26 Doobay, M. F. *et al.* Differential expression of neuronal ACE2 in transgenic mice with  
447 overexpression of the brain renin-angiotensin system. *Am J Physiol Regul Integr Comp  
448 Physiol* **292**, R373-381, doi:10.1152/ajpregu.00292.2006 (2007).

449 27 Wrapp, D. *et al.* Cryo-EM structure of the 2019-nCoV spike in the prefusion  
450 conformation. *Science* **367**, 1260-1263, doi:10.1126/science.abb2507 (2020).

451 28 Bos, R. *et al.* Ad26 vector-based COVID-19 vaccine encoding a prefusion-stabilized  
452 SARS-CoV-2 Spike immunogen induces potent humoral and cellular immune responses.  
453 *NPJ Vaccines* **5**, 91, doi:10.1038/s41541-020-00243-x (2020).

454 29 Krammer, F. SARS-CoV-2 vaccines in development. *Nature* **586**, 516-527,  
455 doi:10.1038/s41586-020-2798-3 (2020).

456 30 Hsieh, C. L. *et al.* Structure-based design of prefusion-stabilized SARS-CoV-2 spikes.  
457 *Science* **369**, 1501-1505, doi:10.1126/science.abd0826 (2020).

458 31 Giordano-Santini, R. *et al.* Fusogen-mediated neuron-neuron fusion disrupts neural  
459 circuit connectivity and alters animal behavior. *Proc Natl Acad Sci U S A* **117**, 23054-  
460 23065, doi:10.1073/pnas.1919063117 (2020).

461 32 Polack, F. P. *et al.* Safety and efficacy of the BNT162b2 mRNA COVID-19 vaccine. *N  
462 Engl J Med* **383**, 2603-2615, doi:10.1056/NEJMoa2034577 (2020).

463 33 Watanabe, Y. *et al.* Native-like SARS-CoV-2 spike glycoprotein expressed by ChAdOx1  
464 nCoV-19/AZD1222 vaccine. *ACS Cent Sci* **7**, 594-602, doi:10.1021/acscentsci.1c00080  
465 (2021).

466 34 Corbett, K. S. *et al.* SARS-CoV-2 mRNA vaccine design enabled by prototype pathogen  
467 preparedness. *Nature* **586**, 567-571, doi:10.1038/s41586-020-2622-0 (2020).

468 35 Xia, X. Domains and functions of spike protein in SARS-CoV-2 in the context of vaccine  
469 design. *Viruses* **13**, doi:10.3390/v13010109 (2021).

470 36 Kremers, G. J., Hazelwood, K. L., Murphy, C. S., Davidson, M. W. & Piston, D. W.  
471 Photoconversion in orange and red fluorescent proteins. *Nat Methods* **6**, 355-358,  
472 doi:10.1038/nmeth.1319 (2009).

473 37 Karbowski, M. *et al.* Quantitation of mitochondrial dynamics by photolabeling of  
474 individual organelles shows that mitochondrial fusion is blocked during the Bax  
475 activation phase of apoptosis. *J Cell Biol* **164**, 493-499, doi:10.1083/jcb.200309082  
476 (2004).

477 38 Ou, X. *et al.* Characterization of spike glycoprotein of SARS-CoV-2 on virus entry and  
478 its immune cross-reactivity with SARS-CoV. *Nat Commun* **11**, 1620,  
479 doi:10.1038/s41467-020-15562-9 (2020).

480 39 Shang, J. *et al.* Structural basis of receptor recognition by SARS-CoV-2. *Nature* **581**,  
481 221-224, doi:10.1038/s41586-020-2179-y (2020).

482 40 Joensuu, M. *et al.* Visualizing endocytic recycling and trafficking in live neurons by  
483 subdiffractional tracking of internalized molecules. *Nat Protoc* **12**, 2590-2622,  
484 doi:10.1038/nprot.2017.116 (2017).

485 41 Fath, T., Ke, Y. D., Gunning, P., Gotz, J. & Ittner, L. M. Primary support cultures of  
486 hippocampal and substantia nigra neurons. *Nat Protoc* **4**, 78-85,  
487 doi:10.1038/nprot.2008.199 (2009).

488 42 Guttikonda, S. R. *et al.* Fully defined human pluripotent stem cell-derived microglia and  
489 tri-culture system model C3 production in Alzheimer's disease. *Nat Neurosci* **24**, 343-  
490 354, doi:10.1038/s41593-020-00796-z (2021).

491 43 Yoon, S. J. *et al.* Reliability of human cortical organoid generation. *Nat Methods* **16**, 75-  
492 78, doi:10.1038/s41592-018-0255-0 (2019).

493 44 Brenner, S. The genetics of *Caenorhabditis elegans*. *Genetics* **77**, 71-94 (1974).

494 45 Mello, C. C., Kramer, J. M., Stinchcomb, D. & Ambros, V. Efficient gene transfer in *C.*  
495 *elegans*: extrachromosomal maintenance and integration of transforming sequences.  
496 *EMBO J* **10**, 3959-3970 (1991).

497 46 Schneider, C. A., Rasband, W. S. & Eliceiri, K. W. NIH Image to ImageJ: 25 years of  
498 image analysis. *Nat Methods* **9**, 671-675, doi:10.1038/nmeth.2089 (2012).

499

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515

516 **Author contributions**

517 Conceptualization: RMM, RGS, MAH

518 Methodology: RMM, RGS, GB, YDK, LI, MAH  
519 Investigation: RMM, RGS, EK, ANC, MAR, ER  
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521 Funding acquisition: LI, YDK, FAM, MAH  
522 Project administration: LI, YDK, MAH  
523 Supervision: LI, YDK, MAH  
524 Writing – original draft: RMM, RGS, MAH  
525 Writing – review & editing: RMM, RGS, EK, GB, FAM, YDK, LM, MAH  
526

## 527 **Ethics declarations**

528 Competing interests  
529 The authors declare that they have no competing interest.

530

531

## 532 **Additional information**

533 Supplementary Information is available for this paper:  
534 Supplementary video 1  
535 Supplementary video 2  
536 Supplementary video 3

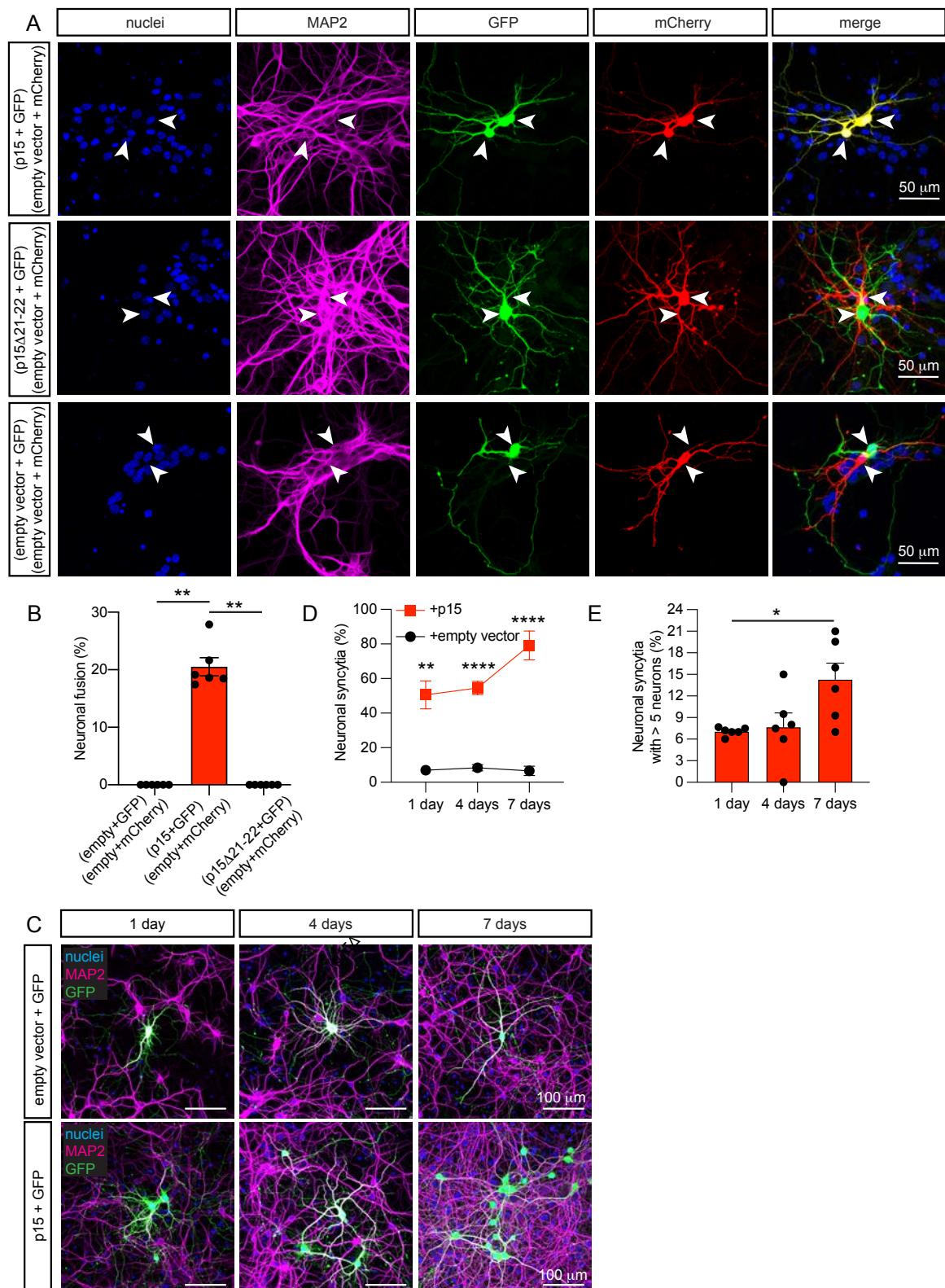
537

538 All data are available in the main text or the supplementary materials. Correspondence and  
539 requests for materials should be addressed to Massimo A. Hilliard.

540

541 **Figures**

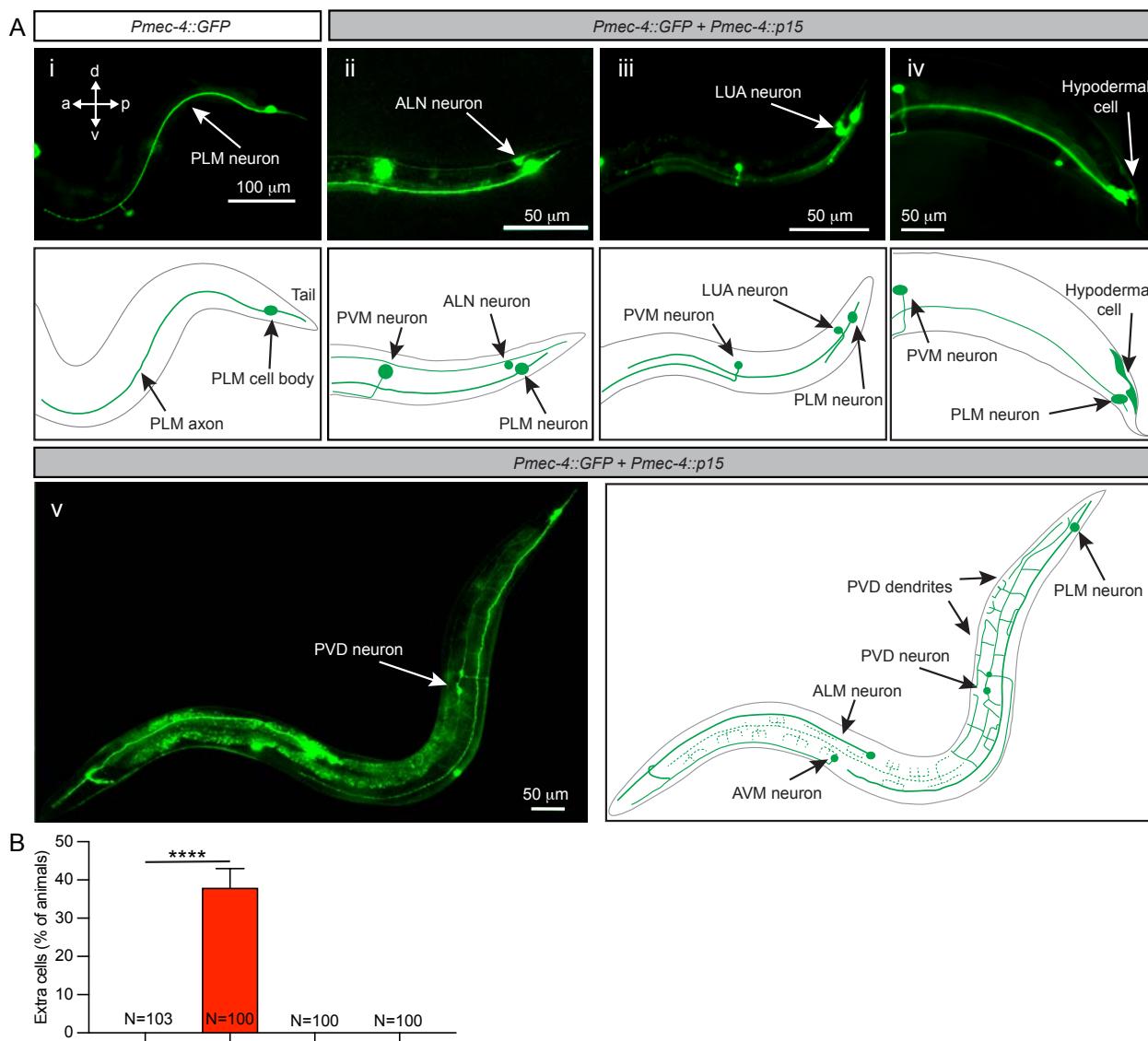
542 **Fig. 1: Expression of p15 induces fusion of murine neurons in culture.**



543 **a**, Representative images of fused neurons (upper row panels) identifiable with GFP (green) and  
544 mCherry (red) fluorescence appearing simultaneously in adjacent neurons (yellow in the merge  
545 panel), or non-fused control neurons (middle and lower row panels) with green and red  
546 fluorescence in adjacent neurons. Two populations of hippocampal neurons expressing either  
547 p15 and GFP, or empty vector and mCherry were cultured together for 7 days (7 DIV). In  
548 control conditions, p15 was substituted by the non-fusogenic mutant p15 $\Delta$ 21-22, or by the empty  
549 vector. Immunocytochemistry for nuclei (blue), neuronal MAP2 (magenta), GFP (green) and  
550 mCherry (red). **b**, Quantification of neuronal fusion as the percentage of transfected neurons  
551 that fuse (yellow) when two neurons are in proximity ( $\leq 200$   $\mu$ m). **c**, Representative images of  
552 neurons illustrating the propagation of fusion over time (upper panels). Hippocampal neurons  
553 were co-transfected at 7-10 DIV with p15 and GFP (or empty vector and GFP in control, lower  
554 panels), and were cultured for 1 day, 4 days or 7 days. Immunocytochemistry for nuclei (blue),  
555 MAP2 (magenta) and GFP (green). **d**, Quantification of neuronal syncytia as the percentage of  
556 interconnected neurons within a distance of  $\leq 200$   $\mu$ m. **e**, Quantification of the average number  
557 of interconnected neurons per syncytium containing more than 5 neurons. Data in **b** are  
558 displayed as mean  $\pm$  SEM,  $n > 150$  neurons analyzed in 6 independent dishes from  $> 2$  cultures,  
559 One-way ANOVA Kruskal-Wallis test followed by Dunn's *post hoc* test in **e** comparing all  
560 groups to empty vector control. Data in **d** and **e** are displayed as mean  $\pm$  SEM,  $n > 350$  neurons  
561 analyzed in  $> 4$  independent dishes from 4 cultures. Two-way ANOVA in **d** followed by  
562 Geisser-Greenhouse correction and the Šidák *post hoc* test comparing treatments (+ empty vector  
563 vs + p15) within each condition (days in culture). One-way ANOVA Kruskal-Wallis test  
564 followed by Dunn's *post hoc* test in **e** comparing all groups to 1 day.  $*p < 0.05$ ,  $**p < 0.01$ ,  
565  $***p < 0.0001$ .

566

567 **Fig. 2: Expression of p15 induces neuronal fusion *in vivo* in *C. elegans* neurons.**

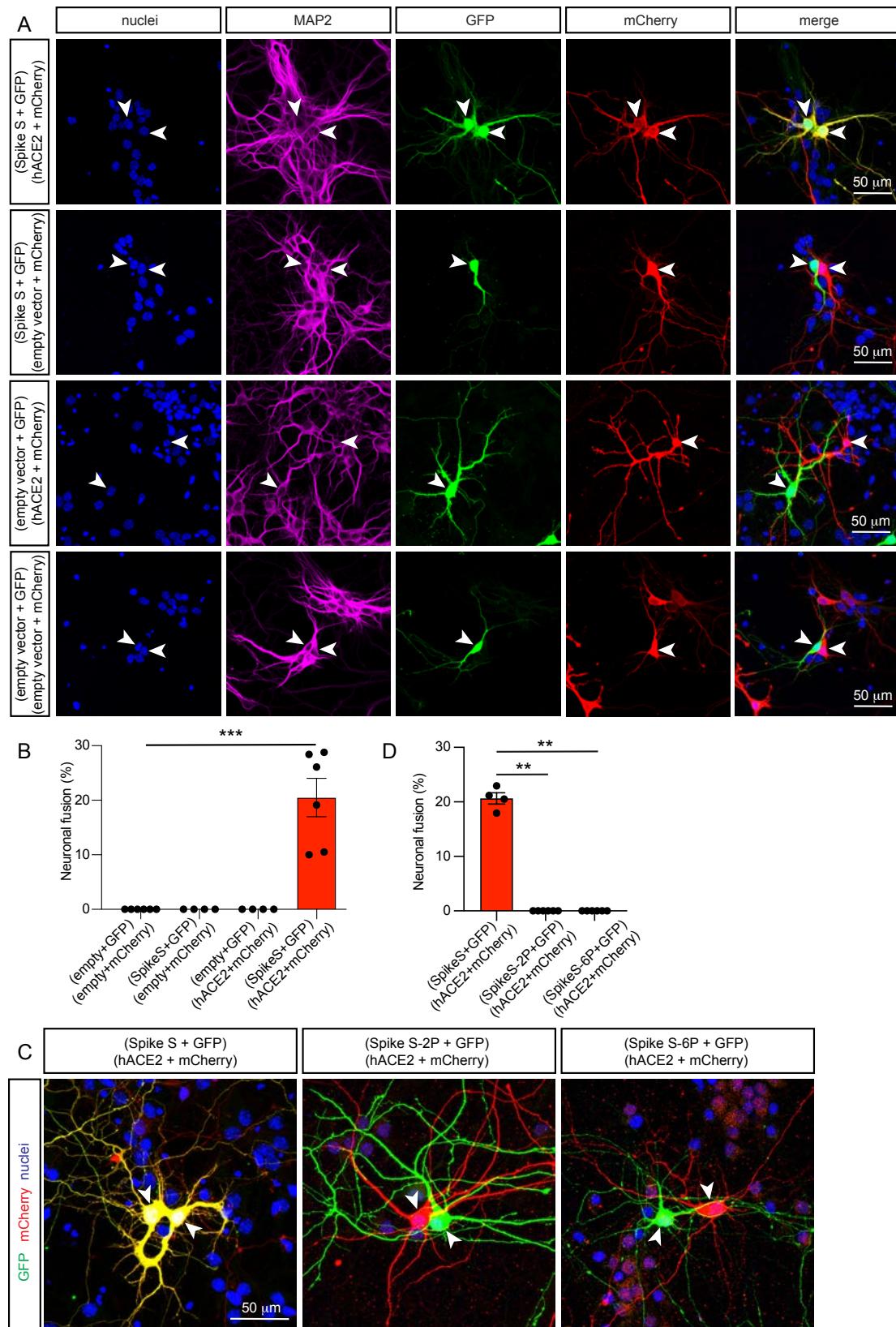


568

569 **a**, Representative images of animals expressing GFP in the six mechanosensory neurons (*Pmec-*  
570 *4::GFP*) in control conditions (i.e. no fusogen present, panel i), or when co-expressing p15  
571 within the same neurons (*Pmec-4::GFP + Pmec-4::p15*) (panels ii, iii, iv, v). The anterior (a),  
572 posterior (p), dorsal (d) and ventral (v) axes are maintained through all the images and the focus  
573 is on the posterior lateral mechanosensory neuron (PLM). Representative images of the  
574 appearance of additional cells in animals expressing active p15: ALN neurons (ii), LUA neurons  
575 (iii), hypodermal (glial-like) cells (iv), and PVD neurons (v). **b**, Quantification of the

576 percentage of animals presenting extra cells in non-transgenic siblings (- transgene), animals  
577 expressing p15, or animals expressing inactive p15 (p15 $\Delta$ 21-22). Data in **b** are displayed as  
578 mean  $\pm$  SD, with the number of animals analyzed per condition (N) listed in the bar graph. One-  
579 way ANOVA Brown-Forsythe and Welch tests in **b** followed by Games-Howell's *post hoc* test,  
580 comparing each mutant with its non-transgenic siblings (- transgene). \*\*\* $p < 0.0001$ .  
581

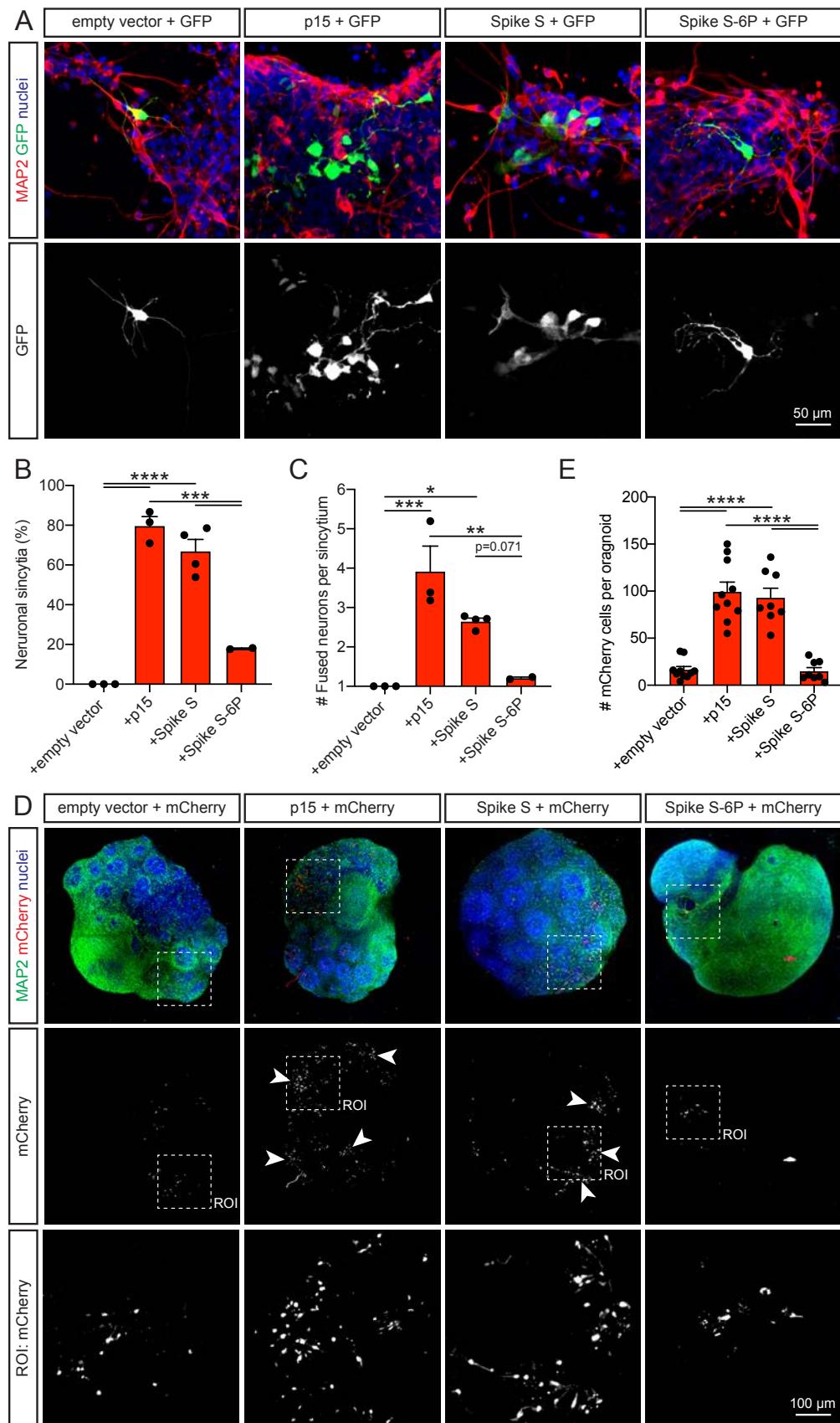
582 **Fig. 3: Expression of spike S and its receptor hACE2 induce fusion of murine neurons in**  
 583 **culture.**



584 **a**, Representative images of fused neurons (first row), or non-fused control neurons (other  
585 rows). Two populations of hippocampal neurons expressing a combination of two plasmids as  
586 indicated on the left (spike S and GFP, hACE2 and mCherry, empty vector and GFP, or empty  
587 vector and mCherry) were cultured together for 7 days (7 DIV). Immunocytochemistry for  
588 nuclei (blue), MAP2 (magenta), GFP (green) and mCherry (red). Neuronal fusion only occurred  
589 (first row) when one population of neurons was transfected with spike S and GFP, and the other  
590 with hACE2 and mCherry, as visualized by the presence of GFP and mCherry in the same  
591 neurons (yellow in the merge panel). **b**, Quantification of neuronal fusion as the percentage of  
592 neurons that fuse (yellow) when two neurons are in proximity ( $\leq 200 \mu\text{m}$ ). **c**, Representative  
593 images of fused neurons (first panel), or non-fused neurons (second and third panels). Two  
594 populations of hippocampal neurons expressing a combination of two plasmids as indicated  
595 above the images (spike S and GFP, hACE2 and mCherry, spike S-2P and GFP, or spike S-6P  
596 and GFP) were cultured together for 7 days (7 DIV). Immunocytochemistry for nuclei (blue),  
597 MAP2 (magenta), GFP (green) and mCherry (red). Neuronal fusion only occurred (first panel)  
598 when the full-length WT spike S protein was transfected, as visualized by the presence of GFP  
599 and mCherry in the same neurons (yellow in the panel), and not when any of the non-fusogenic  
600 mutants (spike S-2P, spike S-6P) were used (second and third panels). **d**, Quantification of  
601 neuronal fusion as the percentage of neurons that fuse (yellow) when two neurons are in  
602 proximity ( $\leq 200 \mu\text{m}$ ). Data in **b** and **d** were displayed as mean  $\pm$  SEM,  $n > 200$  neurons  
603 analyzed in 4-6 independent dishes from 2 dissections, one-way ANOVA Kruskal-Wallis test in  
604 **e** followed by Dunn's *post hoc* test comparing all groups to the group without spike S or hACE2.  
605  $^{**}p < 0.01$ ,  $^{***}p < 0.001$ .

606

607 **Fig. 4: p15 and spike S induce fusion in human neurons and brain organoids.**



608 **a**, Representative images of 2D-cultured neurons illustrating fusion of cells into syncytia.  
609 Human neurons were co-transfected at 40-50 DIV with GFP and either p15, spike S or spike S-  
610 6P (or empty vector in controls), then cultured for 7 days. Immunocytochemistry for nuclei  
611 (blue), MAP2 (red) and GFP (green/white). **b**, Quantification of neuronal syncytia as the  
612 percentage of interconnected neurons within a distance of  $\leq 200$   $\mu\text{m}$ . **c**, Quantification of the  
613 average number of interconnected neurons per syncytium containing more than one neuron. **d**,  
614 Representative images of 3D neuronal organoids illustrating fusion of cells into syncytia.  
615 Organoids were co-transfected at 43-50 DIV with mCherry and either p15, spike S or spike S-6P  
616 (or empty vector in controls), then cultured for 6 days. Immunocytochemistry for nuclei (blue),  
617 MAP2 (green) and mCherry (red/white). Regions of interest (ROIs) show higher magnification  
618 at positions indicated by broken lines. Arrowheads indicate clusters of fused neurons. **e**,  
619 Quantification of the average number of mCherry-positive cells per organoid section 6 days after  
620 transfection. Data in **b**, **c** and **e** are displayed as mean  $\pm$  SEM, averages of  $n > 30$  neurons  
621 analyzed in independent experiments for **b** and **c**, and  $n > 8$  organoids analyzed in independent  
622 experiments for **e**. One-way ANOVA followed by Tukey *post hoc* test in **b** and **c**.  $*p < 0.05$ ,  $**p$   
623  $< 0.01$ ,  $***p < 0.001$ ,  $****p < 0.0001$ .