

1 **The necroptosis machinery mediates axonal degeneration in a**  
2 **model of Parkinson disease**

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4 **Maritza Oñate<sup>1,2,3</sup>, Alejandra Catenaccio<sup>1,2</sup>, Natalia Salvadores<sup>1,2</sup>, Cristian Saquel<sup>1,2</sup>,**  
5 **Alexis Martinez<sup>2,4,5</sup>, Ines Moreno-Gonzalez<sup>6,7</sup>, Nazaret Gamez<sup>6,7</sup>, Paulina Soto<sup>2,4,5</sup>,**  
6 **Claudio Soto<sup>6</sup>, Claudio Hetz<sup>2,4,5,8,9\*</sup> and Felipe A. Court<sup>1,2,9,\*</sup>**

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8 <sup>1</sup>Center for Integrative Biology, Faculty of Sciences, Universidad Mayor, Santiago, Chile

9 <sup>2</sup>FONDAP Geroscience Center for Brain Health and Metabolism, Santiago, Chile.

10 <sup>3</sup>Department of Physiology, Faculty of Biological Sciences, Pontificia Universidad Católica  
11 de Chile, Santiago, Chile.

12 <sup>4</sup>Biomedical Neuroscience Institute (BNI), Faculty of Medicine, University of Chile, Santiago,  
13 Chile.

14 <sup>5</sup>Program of Cellular and Molecular Biology, Institute of Biomedical Sciences, University of  
15 Chile, Santiago, Chile.

16 <sup>6</sup>The Mitchell Center for Alzheimer's Disease, Department of Neurology, McGovern Medical  
17 School, The University of Texas Health Science Center at Houston, Houston, Texas.

18 <sup>7</sup>Networking Research Center on Neurodegenerative Diseases (CIBERNED), Department  
19 of Cell Biology, Faculty of Sciences, University of Malaga, Malaga, Spain.

20 <sup>8</sup>Department of Immunology and Infectious diseases, Harvard School of Public Health,  
21 Boston MA, USA.

22 <sup>9</sup>Buck Institute for Research on Aging, Novato, CA 94945, USA.

23

24 **\*Address correspondence to:**

25 Felipe A. Court: Center for Integrative Biology, Faculty of Sciences, Universidad Mayor.

26 Camino la Piramide 5750, Santiago, Chile, P.O.BOX 8580745, Tel: +56-2-2328-1340; email:  
27 [felipe.court@umayor.cl](mailto:felipe.court@umayor.cl).  
28 Claudio Hetz, Program of Cellular and Molecular Biology, Second floor, sector B, Institute  
29 of Biomedical Sciences, University of Chile. Independencia 1027, Santiago, Chile, P.O.BOX  
30 70086, Tel: +56-2-2978-6506; email: [chetz@med.uchile.cl](mailto:chetz@med.uchile.cl) or [chetz@hsph.harvard.edu](mailto:chetz@hsph.harvard.edu)

31 **Abstract**

32 Parkinson's disease (PD) is the second most common neurodegenerative condition,  
33 characterized by motor impairment due to the progressive degeneration of dopaminergic  
34 neurons in the substantia nigra and depletion of dopamine release in the striatum.  
35 Accumulating evidence suggest that degeneration of axons is an early event in the disease,  
36 involving destruction programs that are independent of the survival of the cell soma.  
37 Necroptosis, a programmed cell death process, is emerging as a mediator of neuronal loss  
38 in models of neurodegenerative diseases. Here, we demonstrate activation of necroptosis  
39 in postmortem brain tissue from PD patients and in a toxin-based mouse model of the  
40 disease. Inhibition of key components of the necroptotic pathway resulted in a significant  
41 delay of 6-hydroxydopamine dependent axonal degeneration of dopaminergic and cortical  
42 neurons *in vitro*. Genetic ablation of necroptosis mediators MLKL and RIPK3, as well as  
43 pharmacological inhibition of RIPK1 *in vivo*, decreased dopaminergic neuron degeneration,  
44 improving motor performance. Together, these findings suggest that axonal degeneration in  
45 PD is mediated by the necroptosis machinery, a process here referred to as *necroaxoptosis*,  
46 a druggable pathway to target dopaminergic neuronal loss.

47

48 **Running title:** Necroptosis activation in Parkinson's disease

49

50 **Keywords:** Parkinson's disease / necroptosis / axonal degeneration /  
51 neurodegeneration / nec-1s / necroaxoptosis

## 52 **Introduction**

53            Parkinson's disease (PD) is the second most common neurodegenerative disease.  
54    The increase in life span and changes in life style predicts that PD will double over the next  
55    generation (1). The characteristic motor symptoms of PD include resting tremor,  
56    bradykinesia, rigidity of the limbs and abnormal gait (2), associated to the loss of  
57    dopaminergic neurons in the substantia nigra pars compacta (SNpc) and reduction of  
58    dopamine release in the striatum (3). Several genetic alterations have been identified in rare  
59    familial PD cases (4), whereas more than 90% of the cases are considered idiopathic.

60            A characteristic feature of PD is a progressive "dying back" process of neuronal loss.  
61    PD pathology initiates in the striatal terminals and proceeds in a retrograde fashion to the  
62    somas in the SNpc (6–9). Studies using genetic manipulation demonstrated that apoptosis  
63    inhibition prevents cell death of mesencephalic neurons, without a significant impact in the  
64    loss of axons and dopamine depletion in the striatum (10–12). Axonal degeneration is a  
65    regulated process independent of cell soma survival, executed by mechanisms that  
66    disassemble axonal structures that are then engulfed by glial cells and circulating  
67    macrophages (13). Among the mechanisms of axonal destruction, key components include  
68    calcium release from the endoplasmic reticulum, mitochondrial dysfunction, and ROS  
69    production (14). We have recently demonstrated that axonal degeneration triggered by  
70    mechanical or toxic insults is commanded by a regulated program that involves components  
71    of the necroptosis machinery (15,16).

72            Necroptosis is a programmed cell death mechanism associated with cell death in  
73    several pathological conditions such as ischemia-reperfusion injury in the heart, liver injury,  
74    viral infection, cancer and neurodegeneration (17,18). Necroptosis can be initiated by a  
75    variety of stimuli (19), including TNF- $\alpha$ , Fas, TRAIL, interferons and activation of TLRs  
76    (review in (20)). TNF- $\alpha$ -induced necroptosis is the best characterized necroptotic initiator to

77 date. Upon TNFR1 activation by TNF- $\alpha$ , complex I is formed by the recruitment of the kinase  
78 RIPK1, cIAPs1/2 and adapter proteins TRADD and TRAF1/2 (21). In this complex, RIPK1  
79 is poly-ubiquitylated by cIAPs (22) activating stress pathways (i.e. MAPK and NF- $\kappa$ B),  
80 impacting cell survival and inflammation (23). Nevertheless, dissociation of complex I leads  
81 to the formation of a cytosolic pro-cell death machinery (complex II) or the necrosome  
82 complex if caspases are inhibited or absent (24). Upon necrosome formation, RIPK1 is  
83 activated and autophosphorylated, activating RIPK3 which binds and phosphorylates the  
84 pseudokinase MLKL, the most downstream effector of necroptosis (20). Phosphorylated  
85 MLKL oligomerizes and translocates to the plasma membrane, forming pores to directly  
86 execute a necrotic form of cell death (24,25). Other studies suggested that MLKL interacts  
87 with the cation channel TRPM7, leading to abnormal calcium influx leading to cell swelling  
88 and plasma membrane rupture (29). Additionally, it has been demonstrated that MLKL is  
89 required for necrosome translocation to the mitochondria to enhance aerobic respiration and  
90 mitochondrial ROS production, leading to a metabolic collapse (30).

91         Recent reports associated the activation of necroptosis to several neurodegenerative  
92 conditions (reviewed in (31)). Genetic or pharmacological inhibition of necroptosis exert  
93 neuroprotective effects in models of brain damage, including ischemia (32–34), traumatic  
94 injury (35), viral infections (36), retinal damage (37–39) and spinal cord injury (40). Recent  
95 advances in the field have demonstrated the therapeutic potential of inhibiting necroptosis  
96 in several neurodegenerative diseases including amyotrophic lateral sclerosis (ALS) (41,42)  
97 multiple sclerosis (MS) (43) and Alzheimer's disease (AD) (44). Analysis of postmortem  
98 brain tissue derived from PD patients indicated increased levels of RIPK1, RIPK3 and MLKL  
99 at the SNpc (45). Inhibition of RIPK1 using the small molecule nec-1s protected  
100 dopaminergic neurons on a pharmacological model of PD *in vivo* (45) and *in vitro* (46).

101 Nevertheless, whether necroptosis activation is functionally associated to motor dysfunction  
102 and denervation in PD has not been explored.

103 In this study, we investigated the contribution of the necroptotic pathway to axonal  
104 degeneration in dopaminergic neurons in the context of PD. Analysis of human PD  
105 postmortem brain tissue and mouse models of the disease indicated the activation of key  
106 components of the necroptosis machinery in dopaminergic neurons of the SNpc. Functional  
107 assessment indicated that targeting RIPK1 or MLKL significantly reduced axonal  
108 degeneration in dopaminergic primary neurons. Genetic ablation of RIPK3 and MLKL, or  
109 administration of nec-1s to a mouse model of PD attenuated axonal degeneration,  
110 translating into an improvement of motor performance. Our results demonstrate a novel  
111 function of the necroptosis machinery in controlling the mechanisms of axonal destruction  
112 in PD and suggest that strategies to inhibit necroptosis may have important therapeutic  
113 benefits to attenuate neurodegeneration in PD.

114 **Results**

115 **1- Pharmacological inhibition of the necroptosis machinery delays 6-OHDA-induced**  
116 **neurite degeneration *in vitro*.**

117 We studied the involvement of necroptosis in PD using primary neuronal cultures from the  
118 embryonic mesencephalon, as this region give rise to the SNpc during brain development.  
119 6-hydroxydopamine (6-OHDA) was used as a relevant neurotoxic PD insult. First, we  
120 evaluated the expression and activation of the necroptotic component MLKL. pMLKL  
121 expression was studied by immunofluorescence in addition with Acetylated Tubulin staining  
122 to visualize neurons. We observed that 6-OHDA triggers a clear modification of the pMLKL  
123 staining pattern in neurons, from diffuse in control conditions to a punctate pattern in neurites  
124 treated with 6-OHDA (Fig. 1A-D), suggesting the formation of MLKL oligomers in  
125 degenerating neurons. Moreover, neurons were treated with 6-OHDA for 3 and 6 hours and  
126 phosphorylation of MLKL (pMLKL) was evaluated by western blot. Low basal expression of  
127 pMLKL was detected in control conditions after vehicle treatment, which were increased by  
128 3-fold after 6 hours of 6-OHDA treatment (Fig. 1E), suggesting activation of this necroptotic  
129 component after 6-OHDA treatment.

130 Then, we manipulated the necroptotic pathway by inhibiting RIPK1 using nec-1s (47).  
131 Mesencephalic neuronal cultures were exposed to 6-OHDA or vehicle as control, in the  
132 presence or absence of nec-1s followed by morphological assessment by immunostaining.  
133 6-OHDA induced the degeneration of neurites characterized by fragmentation and neurite  
134 beading, a phenomenon that was completely prevented by nec-1s treatment (Fig. 2A).  
135 Quantitative assessment of degeneration (See Supp. Fig. 1A) confirmed the protective  
136 effects achieved by nec-1s over neurites (Fig. 2B). Further classification of neurites into  
137 intact, beaded or fragmented indicated almost complete protection of their integrity (Fig. 2C).

138 Of note, nec-1s alone did not alter neurite morphology of mesencephalic neurons (Fig. 2A-  
139 C).

140 Since RIPK1 can trigger apoptotic cell death under certain conditions (48), we  
141 studied the participation of MLKL, in 6-OHDA-dependent neurodegeneration. To this end,  
142 pharmacological inhibition of MLKL was evaluated in mesencephalic neuronal cultures using  
143 GW806742x (GW80), which binds to MLKL blocking its translocation to the plasma  
144 membrane (49). Although GW80 treatment induce a slight, but not significant protection in  
145 6-OHDA treated neurons (Fig. 2D-E and Supp. Fig. 1B), neurite classification analysis  
146 revealed a significant protection of GW80 over 6-OHDA-dependent neurite degeneration  
147 process (Fig. 2F).

148 In addition to dopaminergic neurons, PD has been also associated to  
149 neurodegeneration of olfactory, cortical and autonomic peripheral neurons (50). Therefore,  
150 we performed pharmacological inhibition of RIPK1 and MLKL in cortical neuronal cultures  
151 exposed to 6-OHDA. Morphologically, treatment with nec-1s or GW80 resulted in a  
152 significant protection of neurite degeneration of cortical neurons exposed to 6-OHDA (Fig.  
153 2G-J).

154 6-OHDA is known to induce cell death in neurons, indeed treatment with 6-OHDA in  
155 mesencephalic and cortical neurons leads to an increase of neurons with condensed nuclei,  
156 a morphological reporter of apoptosis, but not a decrease in neuronal density (Supp. Fig. 2).  
157 Nuclei condensation was not protected by RIPK1 or MLKL inhibition using nec-1s or GW80,  
158 respectively, suggesting the co-activation of two cell death mechanisms in different neuronal  
159 compartments (Supp. Fig. 2), as we have previously demonstrated for other pro-  
160 degenerative stimuli (15).

161           Taken together, our results indicate that treatment with 6-OHDA triggers necroptotic  
162 components activation. Moreover, pharmacological inhibition of two key components of the  
163 necroptotic pathway reduces neurodegeneration in mesencephalic and cortical neurons  
164 cultures.

165

## 166 **2- Activation of necroptosis markers in the brain of PD patients.**

167 We next assessed the possible activation of necroptosis markers in the brain of PD patients.  
168 We analyzed the phosphorylation levels of MLKL, a measure of its activation, in postmortem  
169 samples derived from PD patients and age-matched healthy controls (HC). Analysis of  
170 MLKL phosphorylation at Ser358 and neuromelanin pigment positive cells (marker of  
171 dopaminergic neurons) in the SNpc indicated extensive MLKL phosphorylation in PD cases  
172 (Supp. Fig. 3A and B, see inset and arrows in PD samples). As expected, control tissue  
173 presented marked neuromelanin staining, which was considerably lower in samples from  
174 PD patients (Supp. Fig. 3A, arrowheads), suggesting loss of dopaminergic neurons. We  
175 then assessed the expression of pMLKL in different cellular populations. To this end, double  
176 immunofluorescence analysis was performed in HC or PD samples using antibodies against  
177 tyrosine hydroxylase (TH), GFAP and Iba1, to stain dopaminergic neurons, astrocytes and  
178 microglia, respectively, together with pMLKL (Fig. 3A-C). In addition to the marked increase  
179 in pMLKL in PD patients, colocalization analysis reveal that around 50% of the pMLKL  
180 signal was localized in TH-positive neurons, 20% in astrocytes and only 5% in microglia  
181 (Fig. 3D). In addition, a strong increase in TH/pMLKL double positive cells was detected in  
182 brain tissue derived from PD patients compared to healthy control subjects (Fig. 3E).

183

184 **3- MLKL activation and necrosome formation in the nigrostriatal pathway after**  
185 **exposure to 6-OHDA.**

186 To establish whether the necroptotic pathway contributes to dopaminergic neuron  
187 degeneration in experimental PD, we first analyzed the levels of activation of critical  
188 molecular mediators in animals injected with 6-OHDA. To this end, animals were exposed  
189 to 6-OHDA in the striatum and then biochemical and histological analysis was performed in  
190 the nigrostriatal circuit (Fig. 4A and B). Western blot analysis indicated elevated levels of  
191 phosphorylated MLKL in the striatum only after 3 days of 6-OHDA injection and not at 7 days  
192 post-treatment (Fig. 4C). Moreover, a transient upregulation of total MLKL levels was  
193 observed at 3 days post-6-OHDA injection (Fig. 4C). Similar results were observed for  
194 pMLKL in the nigrostriatal pathway (Fig. 4D). Unexpectedly, when the SNpc of the same  
195 animals was analyzed, no changes in pMLKL and total MLKL was observed after the 6-  
196 OHDA challenge (Fig. 4E).

197 We then determined if the activation of MLKL occurs on a cell-autonomous manner  
198 in dopaminergic neurons. Analyses of the distribution of pMLKL together with tyrosine  
199 hydroxylase (TH) using co-immunofluorescence revealed a significant increase in pMLKL in  
200 dopaminergic neurons at the striatum, nigrostriatal pathway and SNpc (Fig. 4F-H, upper  
201 panels and 4I). Similar results were obtained when phosphorylated RIPK3 were analyzed in  
202 the same regions (Fig. 4F-H, lower panels). Importantly, the increase in pMLKL and pRIPK3  
203 observed after the 6-OHDA challenge was not present in the respective MLKL-deficient  
204 (MLKL<sup>-/-</sup>) or RIPK3-deficient mice (RIPK3<sup>-/-</sup>, Supp. Fig. 4).

205 Activation of MLKL by RIPK3 is dependent on the formation of a RIPK1-RIPK3-MLKL  
206 necrosome complex (28). Therefore, formation of the necrosome was evaluated in the  
207 striatum 3 days after 6-OHDA injection by immunoprecipitation. Pull down of RIPK1 revealed

208 an increase in pMLKL-RIPK1 interaction in the 6-OHDA injected hemisphere compared to  
209 the contralateral side (Fig. 4J). Together, these results demonstrate a progressive and  
210 retrograde activation of the necroptosis machinery in a relevant experimental model of PD.

211

212 **4- The necroptosis machinery contributes to axonal degeneration and neuronal loss**  
213 **on an animal model of PD.**

214 To study the possible participation of the necroptosis signaling pathway to axonal  
215 degeneration *in vivo*, we set up the experimental conditions to dissociate the process of  
216 early axonal degeneration from the loss of somas at the SNpc on a toxicological model of  
217 PD. We established a methodology to evaluate the nigrostriatal circuit in the mouse brain  
218 since most of the studies in PD are focused on striatal denervation of axonal terminals in the  
219 striatum (CPu) and neuronal cell loss in the SNpc. Serial coronal sections of the entire  
220 nigrostriatal circuit of animals unilaterally injected with 6-OHDA at the CPu were obtained at  
221 3 and 7 days post-surgery (Fig. 5A). Striatal denervation was calculated by measuring the  
222 optical density of TH in CPu sections comparing non-injected with injected hemisphere (Fig.  
223 5B). A significant and progressive decrease was observed at 3 and 7 days after 6-OHDA  
224 treatment. (Fig. 5C). Axonal degeneration was analyzed along rostro-caudal axis by  
225 evaluating axonal tract lengths in both hemispheres (Fig. 5D). In the non-injected  
226 hemisphere, axonal tract lengths showed no differences along sections, however a  
227 progressive decrease was detected in injected hemispheres 3 and 7 days after 6-OHDA  
228 injection (Supp. Fig. 5A-B). Estimation of the percentage of axonal loss at 3 and 7 days post  
229 6-OHDA injection demonstrated a spatial and temporal progression of axonal degeneration  
230 (Fig. 5E). Finally, the loss of dopaminergic neuronal somas was estimated in the SNpc by  
231 quantification of TH-positive neurons (Fig. 5F). No changes were observed across the SNpc

232 at 3 days post 6-OHDA injection, whereas 7 days of treatment resulted in significant  
233 neuronal loss (Fig. 5G). These results were also validated when the spatial distribution of  
234 TH neurons was quantified (Supp. Fig. 5C). Injection of vehicle had no effect at the different  
235 regions analyzed compared to the non-injected hemisphere (Supp. Fig. 6A-F). Together,  
236 these results showed that the 6-OHDA model is suitable to study axonal degeneration in the  
237 absence of evident neuronal loss.

238 We next performed loss-of-function studies to define the involvement the necroptosis  
239 machinery in axonal neurodegeneration and motor impairment triggered by 6-OHDA. To this  
240 end, we genetically disrupted the expression of MLKL, as the participation of this protein  
241 defines a canonical necroptotic process. MLKL<sup>-/-</sup> and their littermate control animals  
242 (MLKL<sup>+/+</sup>) were injected with 6-OHDA in the right striatum and after 7 days. Optical density  
243 analysis of the striatum showed no differences in striatal denervation in MLKL<sup>-/-</sup> animals  
244 when compared to control mice (Fig. 6A-B). In sharp contrast, ablation of MLKL expression  
245 significantly protected dopaminergic axonal tracks in animals challenged with 6-OHDA (Fig.  
246 6C-D and Supp. Fig. 7A-B). At the SNpc, the number of TH-positive somas were protected  
247 in MLKL<sup>-/-</sup> mice against 6-OHDA, observing a 21% of loss when compared to a reduction of  
248 34% in littermate control animals (Fig. 6E-F and Supp. Fig. 7C). To validate these results  
249 we genetically targeted the expression of RIPK3. RIPK3<sup>-/-</sup> and their littermate control animals  
250 (RIPK3<sup>+/+</sup>) were injected with 6-OHDA in the right striatum and 7 days after, serial coronal  
251 sections of the entire circuit were obtained to evaluate axonal degeneration. Again, optical  
252 density analysis of the striatum showed no differences in striatal denervation in RIPK3<sup>-/-</sup>  
253 animals when compared to control mice (Fig. 6G-H). In sharp contrast, ablation of RIP3K  
254 expression significantly protected dopaminergic axonal tracks after 6-OHDA (Fig. 6I-J and  
255 Supp. Fig. 7D-E). Together, these results suggest that MLKL and RIPK3 contributes to the  
256 dying back degeneration of axons observed in the 6-OHDA model.

257

258 **5- Ablation of MLKL and RIPK3 expression improve motor performance on a PD**  
259 **model.**

260 Given that both MLKL and RIPK3 deficiency reduced axonal neurodegeneration induced by  
261 6-OHDA *in vivo*, we determined if these effects translated in the recovery of the motor  
262 capacity. The cylinder test was performed to measure forepaw akinesia after unilateral 6-  
263 OHDA lesion. A slight tendency, but no differences were found in RIPK3<sup>-/-</sup> however, MLKL<sup>-</sup>  
264 <sup>-</sup> mice showed a significant improvement in forepaw akinesia at 7 days after injection (Fig.  
265 7A-B). To monitor motor coordination, we performed the rotarod test in the same animals  
266 before surgery, and at 3 and 7 days after 6-OHDA injection. No basal alterations in motor  
267 performance were detected in both MLKL and RIPK3 null mice (Fig. 7C-D). Remarkably, at  
268 3 days post injection, a reduced decay in performance of about 40% was observed in MLKL<sup>-</sup>  
269 <sup>-</sup> mice compared to control animals (Fig. 7C). Similarly, improved motor control was  
270 observed in RIPK3 knockout (Fig. 7D). Importantly, vehicle injection had no effect in motor  
271 performance measured by the rotarod (Supp. Fig. 6G). Together, these results indicate that  
272 the necroptosis machinery mediates in part the neurodegeneration cascade observed in our  
273 toxicological model of PD, resulting in improved motor activity.

274

275 **6- Pharmacological inhibition of RIPK1 decreases dopaminergic neuron degeneration**  
276 **and reduces motor impairment of experimental PD.**

277 Our previous results indicates a functional role of the necroptosis machinery in the  
278 degeneration of dopaminergic axons in *in vivo*. To increase the translational potential of  
279 these findings, we tested the consequences of pharmacologically inhibiting the necroptosis  
280 machinery in experimental PD. To this end, we intraperitoneally administrated nec-1s daily

281 for three days before and after exposing animals to the 6-OHDA challenge at the striatum.  
282 7 days after 6-OHDA injection morphological analysis of the brain was performed.  
283 Consistent with our genetic studies, treatment of animals with nec-1s have no effect in 6-  
284 OHDA-dependent striatal denervation (Fig. 8A-B). When the nigrostriatal axonal tract was  
285 visualized in the same animals, a significant protection was observed after nec-1s  
286 administration (Fig. 8C-D and Supp. Fig. 8A-B). Furthermore, a significant decrease in  
287 forepaw akinesia was found in nec-1s treated mice at 3 days post 6-OHDA injection (Fig.  
288 8E). Finally, motor performance was improved in animals injected with nec-1s (Fig. 8F).  
289 Together, these results indicate that pharmacological intervention of the necroptosis  
290 machinery results in significant neuroprotection of experimental PD.

291 **Discussion**

292 PD is a chronic neurodegenerative condition characterized by the degeneration of  
293 nigrostriatal dopaminergic neurons located in the SNpc. Axonal loss is emerging as a critical  
294 pathological event in PD, which occurs in early stages of the disease and precedes somatic  
295 neuronal death (7,51–54). In addition, axonal degeneration is a common feature of other  
296 diseases including AD, ALS, and MS, representing an interesting transversal target for  
297 therapeutic interventions. However, the exact mechanisms by which axons degenerate in  
298 PD remains unknown. Here, we unveiled a functional role for RIPK1, RIPK3 and MLKL as  
299 part of the molecular machinery that executes axonal degeneration in dopaminergic neurons  
300 upon 6-OHDA. Dopaminergic neurons have the largest volume occupied by dendrites and  
301 axons, where the volume of the soma is virtually a minor portion of the whole cell (55).  
302 Because of the compartmentalized nature of neurons, the idea that the necroptosis  
303 machinery controls the “death” of the axon may have an evolutionary origin where  
304 components of necrosome further specialized in this alternative disassembling pathway.  
305 Based in our findings, we would like to propose the concept of “necroaxoptosis” as a novel  
306 mechanism of axonal degeneration mediated by components of the necroptosis machinery.

307 Axonal loss in models of PD has been usually quantified using relatively simple  
308 measurements (51,56). Here, we developed a novel histological method to study of the  
309 whole nigrostriatal pathway along the antero-posterior axis. Remarkably, measurements of  
310 axonal length in every section was sufficient to depict the retrograde and progressive effect  
311 of the 6-OHDA injection along the tract. In addition, biochemical analysis demonstrated a  
312 coincident activation of the necroptosis pathway.

313 Activation of necroptosis has been described in several neurodegenerative  
314 conditions (57,58). Importantly, interaction between RIPK1, RIPK3 and MLKL is detected  
315 in postmortem samples from MS and AD patients. In our toxicological PD model, we

316 observed an early increase in the interaction between RIPK1 and pMLKL in the striatum,  
317 suggesting the formation of the necrosome complex. Whether necroaxoptosis contributes  
318 to axonal degeneration in other diseases where necroptosis is important to induce the death  
319 of the cell soma remains to be determined.

320           Previous evidence in neurodegenerative conditions indicates that necroptosis can  
321 be activated in glial cells, including astrocytes and oligodendrocytes (41–43). Our *in vitro*  
322 and *in vivo* data suggested that activation of necroptosis takes place in neurons after  
323 treatment with 6-OHDA. Remarkably, our study in human samples showed that pMLKL is  
324 mainly expressed in neurons, which is also in agreement with a cell-autonomous activation  
325 of necroptosis in the axonal compartment initiated by different pro-degenerative stimuli  
326 (15,16). Multiple pathological processes may activate necroptosis in PD, including  
327 inflammation and oxidative stress. It's known that  $\alpha$ -synuclein released from degenerating  
328 neurons stimulates microglia, leading to microglia activation and TNF- $\alpha$  secretion (59,60).  
329 In agreement with this idea, MPTP-induced striatal dysfunction is reduced in mice lacking  
330 TNF- $\alpha$  expression, whereas the loss of dopaminergic neurons at the SNpc is not affected  
331 (61). In other neurodegenerative conditions, including MS and ALS, sustained inflammation  
332 has been associated to TNF $\alpha$ -mediated necroptosis (31,62). Similarly, in a mouse model of  
333 ALS, oligodendrocytes showed greater sensitivity to TNF $\alpha$ -induced cell death, and  
334 microglia showed increased phosphorylation of RIPK1 and increased expression of a  
335 number of inflammatory genes (41). Therefore, in diverse neurodegenerative conditions,  
336 activation of RIPK1 plays a major role in microglia-dependent inflammatory activation,  
337 leading to a cell non-autonomous induction of necroptosis in oligodendrocytes and as a  
338 consequence, neurodegeneration (62). Whereas the occurrence of necroptosis in  
339 oligodendrocytes across the axon surface contributes to necroaxoptosis remains to be  
340 determined.

341           Together, evidence suggest that the necroptotic machinery operates on a cell-  
342 autonomous manner in neurons to execute axonal degeneration possibly involving ROS  
343 production as a triggering event (15), a pathological event involved in PD (63–66).  
344 Additionally, a cell non-autonomous mechanism may contribute to this degeneration process  
345 in dopaminergic neurons where TNF $\alpha$  might be secreted by microglia. Since the  
346 mechanisms associated to axonal loss are different from those involved in canonical  
347 programmed cell death and the destruction of the cell body, necroaxoptosis emerges as a  
348 novel and potential mechanism for axonal degeneration. Thus, inhibition of necroaxoptosis  
349 appears to be a promising therapeutic target for functional and structural preservation of  
350 axons and terminals in PD and in other neurodegenerative conditions.

## 351 **Materials and Methods**

### 352 **Neuronal primary cultures**

353 Mesencephalic neuronal cultures were obtained from embryonic E14.5 C57Bl/6 mice and  
354 cortical neuronal cultures were obtained from embryonic E18.5 Sprague-Dawley rats. The  
355 protocol used for both type of primary culture was the same, except the brain structure  
356 dissected. Briefly, meninges were removed from each brain and ventral mesencephalon or  
357 cortex from both hemispheres were dissected, trypsinized and plated onto 0.1 mg/mL poly-  
358 L-lysine coverslips or plastic dishes. Neurons were grown in Neurobasal media  
359 supplemented with B27 and L-glutamine. After 7 days *in vitro* (DIV), neuronal cultures were  
360 treated with 40  $\mu$ M of 6-OHDA (in 0.2% ascorbic acid) or vehicle. For inhibitory treatments,  
361 cells were exposed to 30  $\mu$ M of necrostatin-1s (nec-1s, Biovision) or 0.5  $\mu$ M of GW806742x  
362 (GW80, AdipoGen). For control conditions, cells were incubated with fresh Neurobasal  
363 supplemented medium (control), vehicle or inhibitors alone.

### 364 **Puncta quantification analysis**

365 Quantification of pMLKL puncta in *in vitro* experiments was measure as the number of  
366 pMLKL-positive particles per neuron (based on DAPI and Acetylated Tubulin staining). All  
367 images were acquired and processed simultaneously using Image J software. Binary masks  
368 of pMLKL staining images were obtained to analyze size fragment of particles. Particles with  
369 a size area between 0.2 and 5  $\mu$ m<sup>2</sup> were classified as pMLKL-positive puncta.

### 370 **Neurite integrity index**

371 Quantification of neurite integrity in *in vitro* experiments was calculated as the ratio between  
372 staining area of intact neurites and total staining area (area of intact neurites + area of  
373 fragmented neurites) of Acetylated Tubulin immunofluorecense. Non-neuronal staining was  
374 discarded by co-localization of Acetylated Tubulin and Neurofilament heavy chain staining.  
375 All images were acquired and processed simultaneously using Image J software. Binary  
376 masks of each image were obtained to analyze size fragment of particles. Particles with a

377 size area equal or lower than 25  $\mu\text{m}^2$  and with a circularity index higher than 0.3 were  
378 classified as degenerated neurite fragments. Particles with a size area higher than 25  $\mu\text{m}^2$   
379 with a circularity index lower than 0.3 were classified as intact neurite.

### 380 **Human tissue staining**

381 Human brain tissue (substantia nigra pars compacta) from PD patients (n=3) and control  
382 samples (HC, n=3) was obtained from Banner Sun Health Research Institute. After blocking  
383 the endogenous peroxidase activity with 3%  $\text{H}_2\text{O}_2$ -10% methanol for 20 min, brain sections  
384 were heated at 80°C for 30 min in 50mM citrate buffer pH 6.0 for antigen retrieval prior  
385 overnight incubation with anti-phospho-MLKL (S358) mouse monoclonal antibody (1:200:  
386 Signalway antibody). Primary antibody was detected by incubating 1 hour with sheep anti-  
387 mouse HRP-linked secondary antibody (General Electric), and peroxidase reaction was  
388 visualized using DAB Kit (Vector) following the manufacturer's instructions. Finally, all  
389 sections were dehydrated in graded ethanol, cleared in xylene, and cover-slipped with DPX  
390 mounting medium. 3 samples from each individual were examined under a bright field  
391 microscope (DMI6000B, Leica Microsystems) and representative photomicrographs were  
392 taken with a digital camera (DFC310 FX Leica). Immunoreactivity percentage was defined  
393 as the percentage of area stained with anti-pMLKL related to the substantia nigra analyzed  
394 in different coronal sections (3 sections/subject). pMLKL-immunopositive signal was  
395 converted into 8-bit gray scale and identified by a threshold intensity to quantify the area  
396 labeled per total area analyzed. For double immunofluorescence labeling, brain sections  
397 were treated with autofluorescence eliminator reagent (2160, Millipore Sigma) and  
398 incubated with anti-phospho-MLKL (S358) mouse monoclonal antibody (1:200: Signalway  
399 antibody) for 48 hours, followed by overnight incubation with secondary primary antibody:  
400 rabbit polyclonal anti-TH (1:1000, Millipore Sigma), anti-GFAP (1:1000, Abcam), or anti-Iba1  
401 (1:1000, Wako Chemicals). Sections were sequentially incubated for 1 hour with goat anti-  
402 rabbit Alexa 488 and donkey anti-mouse Alexa 594 secondary antibodies (1:500,

403 Invitrogen), rinsed and covered using FluorSave Reagent (Merck Millipore) for  
404 immunofluorescent evaluation under a confocal laser microscope (Nikon A1R). Human  
405 samples were manipulated following the universal precautions for working with human  
406 specimens and as directed by the Institutional Review Board of the University of Texas  
407 Health Science Center at Houston (HSC-MS-14-0608).

#### 408 **Experimental animals**

409 Adult (12-16 weeks old) C57BL/6 mice, MLKL knockout and RIPK3 knockout mice were  
410 used. MLKL and RIPK3 knockout mice were kindly provided by Dr. Douglass Green (St.  
411 Jude Children's Research Hospital, Memphis, TN, USA) and have been described  
412 previously (67,68). Animals were kept under standard conditions of light and temperature  
413 and were feed with food and water *ad libitum* in the Animal Facility of the Sciences Faculty  
414 of the Mayor University. The research protocol n° 08-2016 was approved by the Animal  
415 Care and Use Scientific Ethic Committee of the Mayor University.

#### 416 **Toxicological model of PD**

417 Mice were anesthetized with isoflurane and placed in a stereotaxic frame (David Kopf  
418 Instruments, USA). A single unilateral injection was performed in the right striatum at the  
419 following coordinates: anteroposterior (AP): +0.07 cm, medio-lateral (ML): -0.17 cm and  
420 dorso-ventral (DV): -0.31 cm, relative to bregma (according to (69)) as previously described  
421 (70). 2 µl of a solution of 8 µg of 6-OHDA (4 µg/µl in 0.2% ascorbic acid) was injected at a  
422 rate of 0.5 µl/min. As control, we injected 2 µl of vehicle solution (0.2% ascorbic acid) using  
423 the same protocol. Animals were euthanized by overdose of anesthesia at different days  
424 post injection for further analysis.

#### 425 **Histological analysis**

426 Mice were deeply anesthetized with isoflurane and intracardially perfused with isotonic  
427 saline followed by 4% paraformaldehyde. Brains were dissected, post-fixed overnight in 4%  
428 paraformaldehyde at 4 °C and then incubated in 30% sucrose. Tissue was cryoprotected in

429 optimal cutting temperature compound (OCT, Tissue-Tek) at -20 °C and serial coronal  
430 sections of 25 µm thick containing the nigrostriatal circuit (from rostral striatum to ventral  
431 midbrain) were obtained using a cryostat (Leica, CM1860). Injected hemisphere was  
432 marked for identification. Serial free-floating sections were processed for  
433 immunohistochemistry as previously described (70). Briefly, slices were quenched with  
434 0.3% H<sub>2</sub>O<sub>2</sub> for 30 min, blocked with 0.5% bovine serum albumin and 0.2% triton X-100 for  
435 2 hours and incubated with primary antibody (rabbit anti-Tyrosine hydroxylase, 1:500  
436 Millipore) overnight at 4 °C. Then, sections were washed with 0.1 M PBS and incubated  
437 with secondary biotinylated antibody (goat anti-rabbit, 1:500 Vector Laboratories) for 2  
438 hours at RT. After washing, slices were incubated with avidin-biotin-peroxidase complex  
439 (Vector Laboratories) for 1 hour at RT followed by 0.1 M PBS washes and developed with  
440 3,3-diaminobenzidine (DAB, Sigma-Aldrich). Finally, sections were mounted on glass slides  
441 with Entellan medium (Merck).

#### 442 **Densitometry analysis of dopaminergic striatal innervation**

443 Dopaminergic terminals at the striatum (CPu) were analyzed every 100 µm of the entire  
444 area. Sections were scanned using an Epson Perfection V600 Photo scanner and striatal  
445 innervation was quantified by measuring the optical density of TH-immunoreactivity in the  
446 striatum using ImageJ software (NIH, USA). Results were expressed as the integrated  
447 density of the entire region and as the percentage of TH immunoreactivity loss compared to  
448 control hemisphere as previously described (70).

#### 449 **Quantification of dopaminergic axons**

450 Dopamine axonal tract at the nigrostriatal pathway (NSt) images were obtained in a Nikon  
451 Eclipse E200 microscope. The length of the TH-positive axonal tract was calculated in each  
452 sections every 100 µm. Spatial distribution of axonal length was calculated in each section  
453 along the antero-posterior axis in injected and contralateral non-injected hemisphere.  
454 Percentage of axonal loss was calculated in the injected hemisphere compared with the

455 contralateral hemisphere in each section. Measurements were performed using Image J  
456 software.

#### 457 **Dopaminergic neuron cell counting**

458 Estimation of the number of TH-positive neurons at the SNpc was performed in serial  
459 sections of the entire midbrain every 100  $\mu\text{m}$  as previously reported (70). The percentage of  
460 TH-immunoreactive neurons relative to the contralateral (non-injected) side was determined  
461 by counting the total number of TH-positive neurons in the entire SNpc. Spatial distribution  
462 analysis of each section was performed along antero-posterior axis. Measurements were  
463 performed using Image J software.

#### 464 **Tissue processing and co-immunoprecipitation**

465 For biochemical analysis, brains were extracted and washed in ice-cold 0.1 M PBS and  
466 then sectioned in a stainless steel brain matrix (coronal slices, 1 mm spacing). 2 mm  
467 sections from striatum, nigrostriatal axonal pathway and mesencephalon (containing the  
468 entire SNpc) were homogenized in 100  $\mu\text{l}$  of RIPA buffer (50 mM Tris-HCl, 150 mM NaCl,  
469 1% NP-40, 0.5% Sodium deoxycholate, 0.1% SDS and 5 mM EDTA, pH 7.6) containing  
470 protease (1 mM PMSF and protein inhibitor cocktail, PIC) and phosphatase inhibitors (1  
471 mM NaF and 50 mM  $\text{Na}_3\text{VO}_4$ ). Protein concentration was estimated using the Pierce BCA  
472 Protein Assay Kit (Thermo Scientific). For cell culture experiments, cells were collected in  
473 0.1 M PBS and then homogenated in RIPA buffer for protein quantification as previously  
474 described. For immunoprecipitation experiments, 100  $\mu\text{g}$  of brain protein lysates were  
475 incubated with 4  $\mu\text{g}$  of anti-RIPK1 (Cell signaling) for 48 hours while rotating at 4  $^{\circ}\text{C}$ . After  
476 incubation, 50  $\mu\text{l}$  of Protein A magnetic beads (Invitrogen) were added to each sample and  
477 were incubated at 4  $^{\circ}\text{C}$  for 3 h while rotating. Following magnetic separation, beads were  
478 mixed with loading buffer and boiled at 90  $^{\circ}\text{C}$  for 5 min. Samples were loaded onto a 10%  
479 SDS/PAGE gel and Western blot was performed for pMLKL antibody as followed described.

#### 480 **Western blot**

481 Brain and cell culture lysates containing 50 µg of protein were loaded into 10% SDS/PAGE  
482 gels and transferred onto methanol-activated PVDF membranes (Thermo Scientific).  
483 Membranes with transferred proteins were blocked/permeabilized for 1 hour in 5% BSA in  
484 TBS and incubated with different antibodies (mouse anti-pMLKL, 1:1000 Millipore, rabbit  
485 anti-MLKL, 1:1000 Abcam, and rabbit anti-Hsp90, 1:1000 Santa Cruz) overnight at 4 °C.  
486 Membranes were incubated with HRP-secondary antibodies (goat anti-mouse HRP and  
487 goat anti-rabbit HRP, 1:3000 Biorad) for 1 hour at RT and revealed using ECL (Invitrogen)  
488 and Chemidoc™ MP Imaging System (Biorad). For densitometry analysis of the bands,  
489 Image J software was used.

#### 490 **Culture cells immunofluorescence**

491 For immunocytochemical analysis, neurons were fixed with 4% paraformaldehyde for 15  
492 min. Then, cells were blocked-permeabilized in 5% gelatin from cold water fish skin and  
493 0.1% triton X-100 for 1 h followed by incubation with primary antibodies (mouse anti-  
494 acetylated tubulin, 1:1000 Sigma-Aldrich, rabbit anti-Neurofilament heavy chain, 1:500  
495 Sigma-Aldrich and rabbit anti-pMLKL, 1:250 Abcam) overnight at 4 °C. After washing with  
496 0.1 M PBS, cells were incubated with secondary antibodies (goat anti-mouse Alexa Fluor  
497 488, goat anti-mouse Alexa Fluor 546, goat anti-rabbit Alexa Fluor 546, and goat anti-rabbit  
498 Alexa Fluor 488, 1:1000 Thermo-Fisher) for 2 hours at RT, then washed with 0.1 M PBS  
499 and coverslips were mounted on slides with Fluoromont-G (Electron Microscopy Sciences)  
500 solution with DAPI nuclear staining (Thermo-Fisher). Images were obtained using a Leica  
501 DMI8 microscope.

#### 502 **Mouse brain immunofluorescence**

503 For immunofluorescence analysis, 25 µm coronal sections from the striatum, nigrostriatal  
504 pathway and substantia nigra pars compacta were used. Antigen retrieval was performed  
505 with boiling Sodium Citrate 10mM, pH 6 for 10 min and then, sections were  
506 blocked/permeabilized in 5% BSA, 0,3% Triton X-100 in 0.1M TBS for 2 hours at RT,

507 following by primary antibody incubation overnight at 4°C (rabbit anti-pRIPK3, 1:200  
508 Abcam; mouse anti-TH, 1:200 Millipore; mouse anti-pMLKL, 1:200 Millipore; rabbit anti-TH,  
509 1:200 Millipore, in blocking/permeabilizing solution). Sections were washed in 0.1M TBS  
510 and then incubated in secondary antibodies for 2 hours at RT in TBS (goat anti-rabbit Alexa  
511 Fluor 488, goat anti-mouse Alexa Fluor 546, goat anti-mouse Alexa Fluor 488 and goat anti-  
512 rabbit Alexa Fluor 546, 1:400 Thermo-Fisher). Finally, sections were washed in 0.1M TBS  
513 and mounted in Mowiol (Sigma).

#### 514 ***In vivo* necrostatin-1s treatment**

515 For nec-1s (Biovision, CA, USA) preparation the compound was dissolved in DMSO (50%  
516 w/v) and then transferred into 35% PEG solution as previously described (43). C57Bl/6 mice  
517 were treated for 3 days with nec-1s (8 mg/Kg i.p.) or vehicle before surgery and then, daily  
518 injected for 7 days after 6-OHDA injection.

#### 519 **Behavioral tests in mice**

520 Behavioral tests were performed in injected animals before surgery (baseline) and at 3 and  
521 7 days post-injection for analysis of functional motor and coordination performance. The  
522 cylinder test was used to assess asymmetric forelimb use as previously reported (70). Mice  
523 were placed in a transparent glass cylinder of 20 cm diameter for 5 min and was videotaped  
524 during the test. The number of 'ipsilateral' (right touches) and 'contralateral touches' (left  
525 touches) was quantified and represented as the percentage of 'contralateral touches' of all  
526 movements observed. For the rotarod test, mice were placed into a rotating rod (Model  
527 LE8500, Panlab SL) and the time until mice fell was measured (latency fall). Animals were  
528 subjected to 4 trials per day using the accelerated speed test protocol, consisted in  
529 increasing speed trial starting with 4 rpm up to 40 rpm within 120 s. Animals waited ~5 min  
530 between each trial to avoid fatigue.

#### 531 **Statistical analysis**

532 All experiments were analyzed in a blinded manner. Data are shown as mean  $\pm$  SEM.  
533 Statistical analysis were performed using Student's t-test or one-way ANOVA, two-way  
534 ANOVA test or two-way repeated measures ANOVA test, followed by Bonferroni's *post hoc*  
535 test or with Mann Whitney non-parametric test, using GraphPad Prism 5.0 software.

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550

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552 M.O, A.C., C.S.\*, A.M, N.G. analyzed data. M.O., I.M-G., C.S.\*\*, C.H., F.A.C., designed  
553 experiments and wrote the manuscript.

554 \*Cristian Saquel, \*\*Claudio Soto.

555

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750

751 **Figure Legends**

752

753 **Figure 1. pMLKL activation in cultured neurons after 6-OHDA treatment. (A-C)**

754 Mesencephalic primary cultures were treated with 6-OHDA or vehicle for 1 hour. Untreated  
755 cultures were used as control. Cells were immunostained against Acetylated Tubulin (AcT,

756 red) and phospho-MLKL (pMLKL, green). Nuclei were stained with DAPI (blue). **(D)**

757 Quantification of the number of pMLKL-positive puncta per neuron was estimated in each

758 condition. **(E, left)** Cortical primary cultures were treated with 6-OHDA for 3 or 6 hours.

759 pMLKL expression was measured by western blot. Hsp90 was used as loading control. **(E,**

760 **right)** Densitometric analysis was performed in each condition for pMLKL and normalized

761 against Hsp90. Scale bar, 30  $\mu\text{m}$ ; insets, 15  $\mu\text{m}$ . Data are shown as mean  $\pm$  SEM. Statistical

762 differences were obtained using one-way ANOVA followed by Bonferroni's *post hoc* test. \*

763  $p < 0.05$ , \*\*  $p < 0.01$ .  $n = 3$  per group.

764

765 **Figure 2. Pharmacological inhibition of RIPK1 and MLKL delays neurite degeneration**

766 ***in vitro*. (A-C)** Mesencephalic neuronal cultures were treated for 6 hours with 6-OHDA alone

767 or together with the RIPK1 inhibitor nec-1s, or vehicle. **(D-F)** Similar cultures were treated

768 with 6-OHDA alone or together with the MLKL inhibitor GW80, or vehicle. Untreated cultures

769 were used as control. Cells were immunostained for acetylated tubulin (AcT, green) and

770 neurofilament heavy chain (NF-H, red). Nuclei were stained using DAPI (blue). In each

771 condition, the neurite integrity index **(B and E)** or a classification of neurite morphology **(C**

772 **and F)** was calculated. **(G-H)** Cortical neuronal cultures were treated for 6 hours with 6-

773 OHDA alone or together with vehicle or the RIPK1 inhibitor nec-1s. **(I-J)** Similar cultures

774 were treated with 6-OHDA alone or together with vehicle or the MLKL inhibitor GW80 . Cells

775 were stained for acetylated tubulin (green), neurofilament heavy chain (NF-H) and DAPI

776 (blue). **(H, J)** Neurite integrity index calculated for each condition. Scale bars, 50  $\mu\text{m}$ . Data

777 are shown as mean  $\pm$  SEM. Statistical differences were obtained using one-way ANOVA  
778 followed by Bonferroni's *post hoc* test. \*  $p < 0.05$ , \*\*  $p < 0.01$ ; \*\*\*  $p < 0.001$  compared to  
779 control, vehicle and nec-1s or GW80 conditions. #  $p < 0.05$ ; ##  $p < 0.01$ , ###  $p < 0.001$   
780 compared to 6-OHDA condition.  $n = 3$  per group.

781

782 **Figure 3. Activation of pMLKL in postmortem samples of human PD brains. (A)**

783 Human samples of substantia nigra from healthy control patients (HC) and Parkinson's  
784 disease patients (PD) were co-immunostained against pMLKL (red) and TH (green), (B)  
785 GFAP (green), and (C) Iba-1 (green). Nuclei were stained using DAPI (blue). Arrows  
786 indicate double positive cells. (D) Percentage of colocalization of pMLKL with different cell  
787 markers TH, GFAP and Iba-1 was measured in PD samples. (E) Quantification of  
788 TH/pMLKL double positive cells in each condition. Scale bars, 25  $\mu\text{m}$ . Data are shown as  
789 mean  $\pm$  SEM. Statistical differences were obtained using one-way ANOVA followed by  
790 Bonferroni's *post hoc* test in (D) and student's t-test in (E). \*  $p < 0.05$ ; \*\*\*  $p < 0.001$ .  $n = 3$   
791 per condition.

792

793 **Figure 4. Activation of necroptosis in the nigrostriatal pathway after 6-OHDA**

794 **treatment *in vivo*. (A)** Wild-type mice were injected with 6-OHDA in the right striatum and  
795 analyzed at 3 or 7 days post-injection. (B) Coronal sections of 2 mm thickness were obtained  
796 from the (a) striatum, (b) axonal tract and (c) substantia nigra, and the non-injected and  
797 injected regions were divided for western blot analysis (indicated in red boxes). pMLKL and  
798 MLKL protein expression was evaluated in (C) striatum, (D) nigrostriatal pathway and (E)  
799 and substantia nigra. Hsp90 was used as loading control. Coronal sections of (F) striatum,  
800 (G) nigrostriatal pathway and (H) substantia nigra from mice injected with 6-OHDA were  
801 immunostained at 3 and 7 days post injection. Schemes show the analyzed area in each  
802 brain region. (Upper panels) Sections were immunostained for TH (red) and pMLKL (green)

803 and **(Lower panels)** TH (red) and pRIPK3 (green). Colocalization in each condition is  
804 pointed with arrows. **(I)** TH-positive neurons also immunoreactive for pMLKL or pRIPK3  
805 were counted and normalized to the total dopaminergic TH-positive neurons in substantia  
806 nigra. **(J)** Proteins extracted from 3 days-injected striatum and contralateral hemisphere  
807 were immunoprecipitated with an antibody against RIPK1 and probed for pMLKL. Relative  
808 pMLKL levels were calculated by densitometry. Scale bars, 50  $\mu\text{m}$ . Data are shown as mean  
809  $\pm$  SEM. Statistical differences were obtained using one-way ANOVA followed by  
810 Bonferroni's *post hoc* test in (C), (D), (E), and (I) and by student's t-test in (J). \*  $p < 0.05$ ; \*\*  
811  $p < 0.01$ , \*\*\*  $p < 0.001$ .  $n = 3$  animals per condition.

812

813 **Figure 5. 6-OHDA treatment induces a progressive and retrograde degeneration of**  
814 **nigrostriatal neurons. (A)** Wild-type mice were injected with 6-OHDA in the right striatum.  
815 The contralateral hemisphere was kept non-injected as a control. Serial coronal sections of  
816 the entire nigrostriatal circuit were obtained at 3 and 7 days post 6-OHDA injection and  
817 immunostained for TH. **(B, left)** Representative scheme of striatal region analyzed. **(B,**  
818 **right)** The striatum of non-injected and injected hemispheres are demarked with blue and  
819 green dashed lines, respectively. Striatal area analyzed (CPu) at 3 and 7 days after injection.  
820 Scale bar, 1 mm. **(C)** Striatal denervation was calculated as the total integrated optical  
821 density in each condition. **(D, left)** Scheme of Nigrostriatal (NSt) pathway region analyzed.  
822 **(D, right)** NSt at 3 and 7 days post-injection. Axonal tracts are indicated using dashed lines  
823 for each hemisphere at 3 and 7 days post 6-OHDA injection. Scale bar, 500  $\mu\text{m}$ . **(E)** Spatial  
824 distribution of axonal loss at 3 and 7 days post 6-OHDA injection. **(F, left)** Scheme of the  
825 substantia nigra pars compacta (SNpc) region analyzed. **(F, right)** SNpc at 3 and 7 days  
826 post 6-OHDA injection. Scale bar, 500  $\mu\text{m}$ . **(G)** Quantification of the total number of TH-  
827 positive cells in the entire SNpc at 3 and 7 days post 6-OHDA injection. Data are shown as  
828 mean  $\pm$  SEM. Statistical differences were obtained using one-way ANOVA in (C) and (G)

829 and two-way ANOVA in (E) followed by Bonferroni's *post hoc* test. \*  $p < 0.05$ ; \*\*  $p < 0.01$ ;  
830 \*\*\*  $p < 0.001$ . n = 10 animals per group.

831

832 **Figure 6. MLKL and RIPK3 deficiency delays neurodegeneration after 6-OHDA**  
833 **injection.** MLKL<sup>-/-</sup>, RIPK3<sup>-/-</sup> and corresponding WT sibling mice were injected with 6-OHDA  
834 in the right striatum (CPu) and the contralateral hemisphere was kept non-injected as a  
835 control. Serial coronal sections of the entire nigrostriatal circuit were obtained 7 days after  
836 6-OHDA injection and immunostained for TH. **(A, left)** Scheme of striatal region analyzed.  
837 **(A, right)** Representative striatal coronal sections from MLKL<sup>-/-</sup> and MLKL<sup>+/+</sup> mice  
838 unilaterally injected with 6-OHDA in the right striatum. Scale bar, 1 mm. **(B, left)** Striatal  
839 denervation was calculated as total integrated optical density in non-injected and injected  
840 hemisphere. **(B, right)** Percentage of TH loss staining was estimated from integrated  
841 density. **(C, left)** Scheme of nigrostriatal pathway region analyzed. **(C, right)**  
842 Representative images from nigrostriatal axons (NSt) from MLKL<sup>-/-</sup> and MLKL<sup>+/+</sup> mice  
843 injected and immunostained for TH. Scale bar, 500  $\mu$ m. **(D)** Spatial distribution of axonal  
844 loss in each genotype. **(E, left)** Scheme of substantia nigra region analyzed. **(E, right)** 6-  
845 OHDA induced neuronal loss in MLKL<sup>-/-</sup> and MLKL<sup>+/+</sup> mice analyzed in the substantia nigra  
846 pars compacta (SNpc). Scale bar, 500  $\mu$ m. **(F)** Quantification of total number of TH-positive  
847 cells in the entire SNpc represented as the percentage of neuronal loss in each genotype.  
848 **(G, left)** Scheme of striatal region analyzed. **(G, right)** Representative striatal coronal  
849 sections from RIPK3<sup>-/-</sup> and RIPK3<sup>+/+</sup> mice injected with 6-OHDA in the right striatum. Scale  
850 bar, 1 mm. **(H, left)** Striatal Integrated density **(H, right)** and percentage of TH loss  
851 calculated in each condition. **(I, left)** Scheme of nigrostriatal pathway region analyzed. **(I,**  
852 **right)** Representative coronal sections from nigrostriatal pathway in RIPK3<sup>-/-</sup> and RIPK3<sup>+/+</sup>  
853 injected mice. Scale bar, 500  $\mu$ m. **(J)** Spatial distribution of axonal loss in each genotype.  
854 Data are shown as mean  $\pm$  SEM. Statistical differences were analyzed using two-way

855 ANOVA followed by Bonferroni's *post hoc* test in (B and H, for Integrated density  
856 measurements, D and J), and by student's t-test in (B, F and H, for percentage of loss). \*  $p$   
857  $< 0.05$ ; \*\*  $p < 0.01$ .  $n = 8$  animals per group.

858

859 **Figure 7. MLKL and RIPK3 ablation improves motor behavior after 6-OHDA injection.**

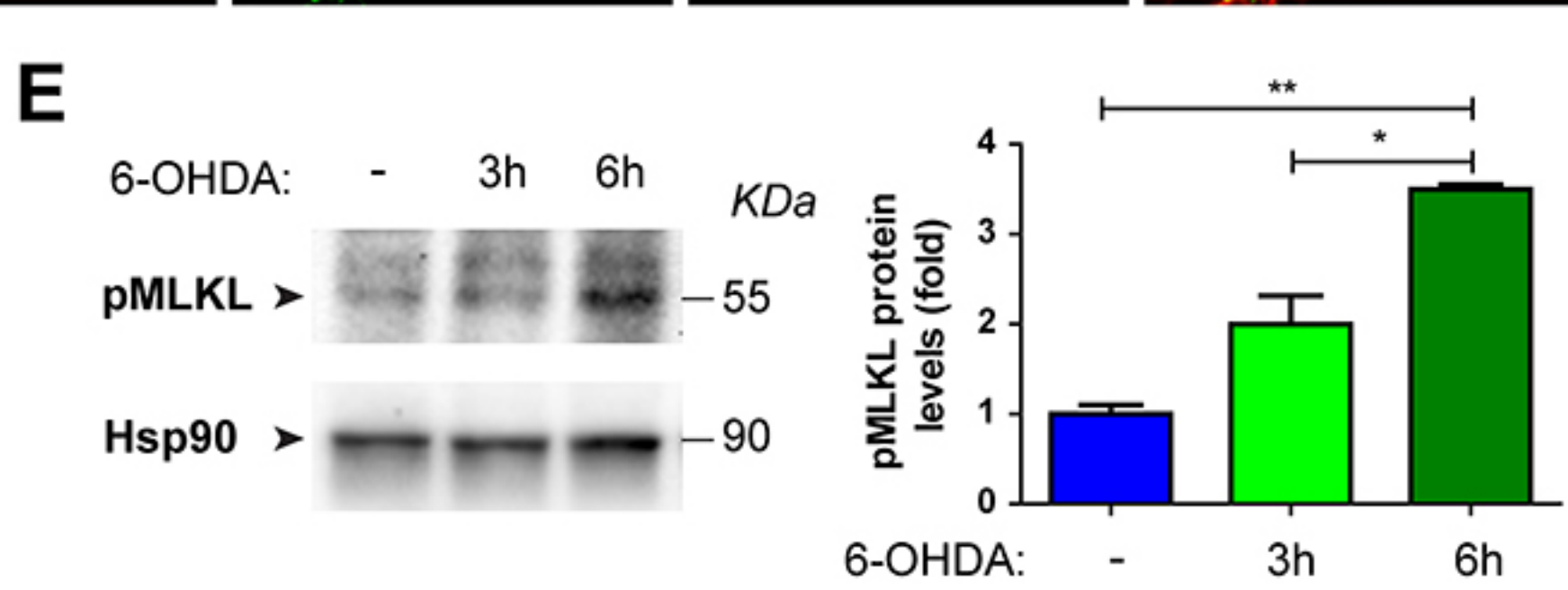
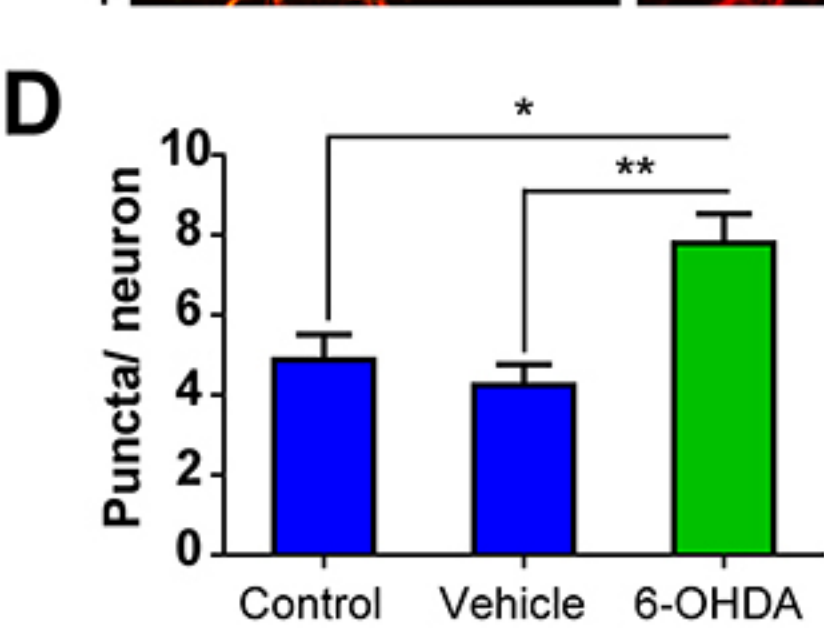
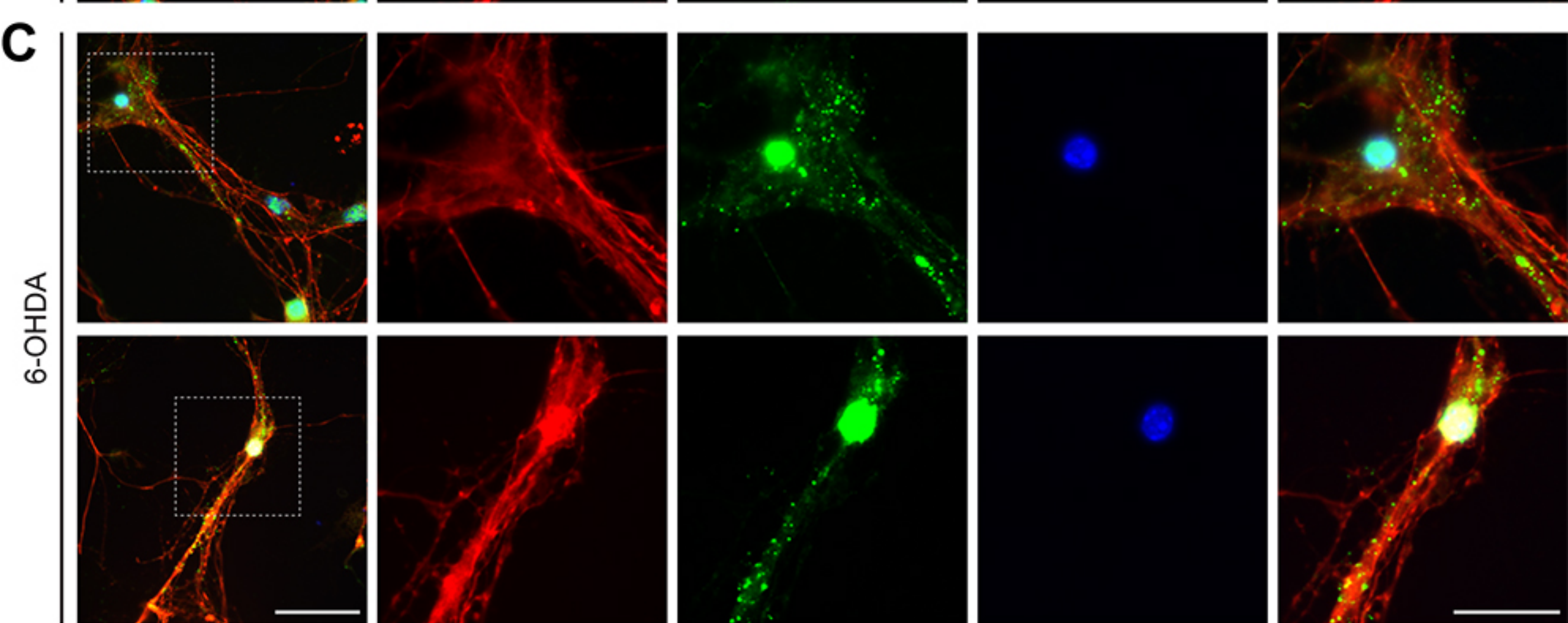
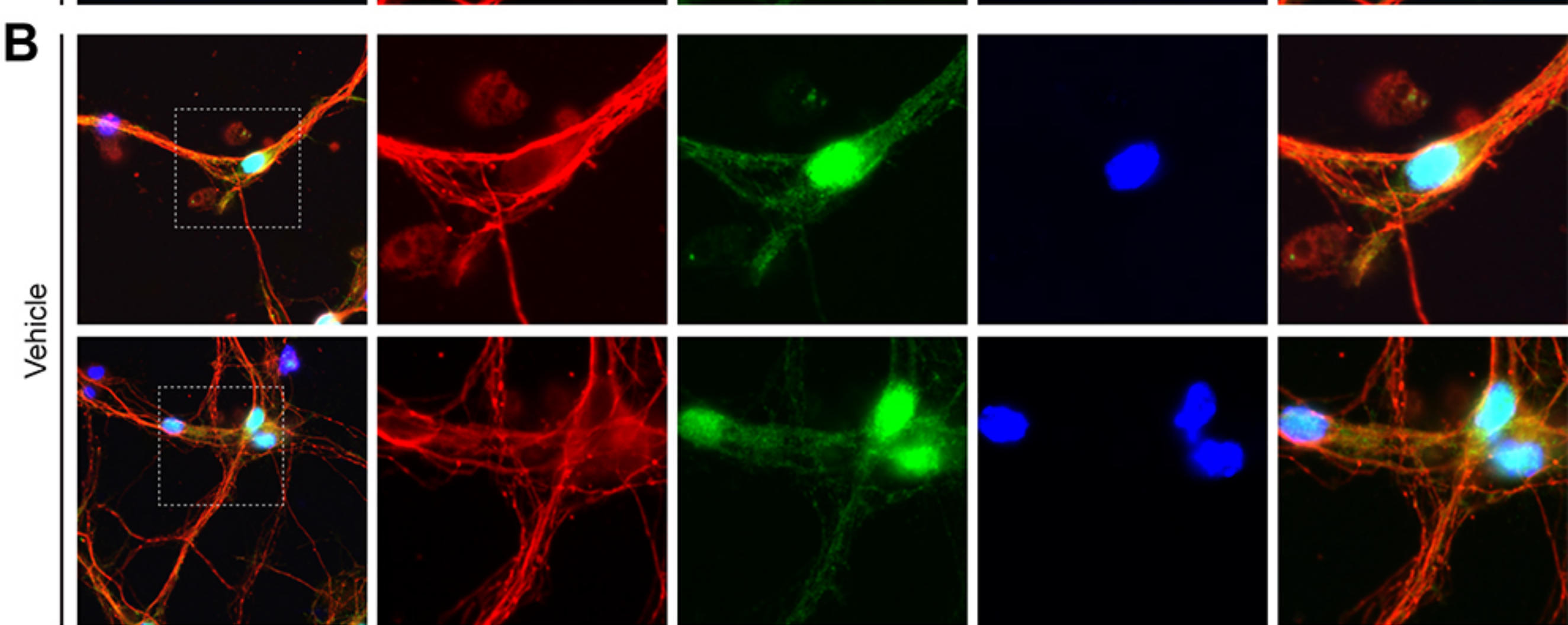
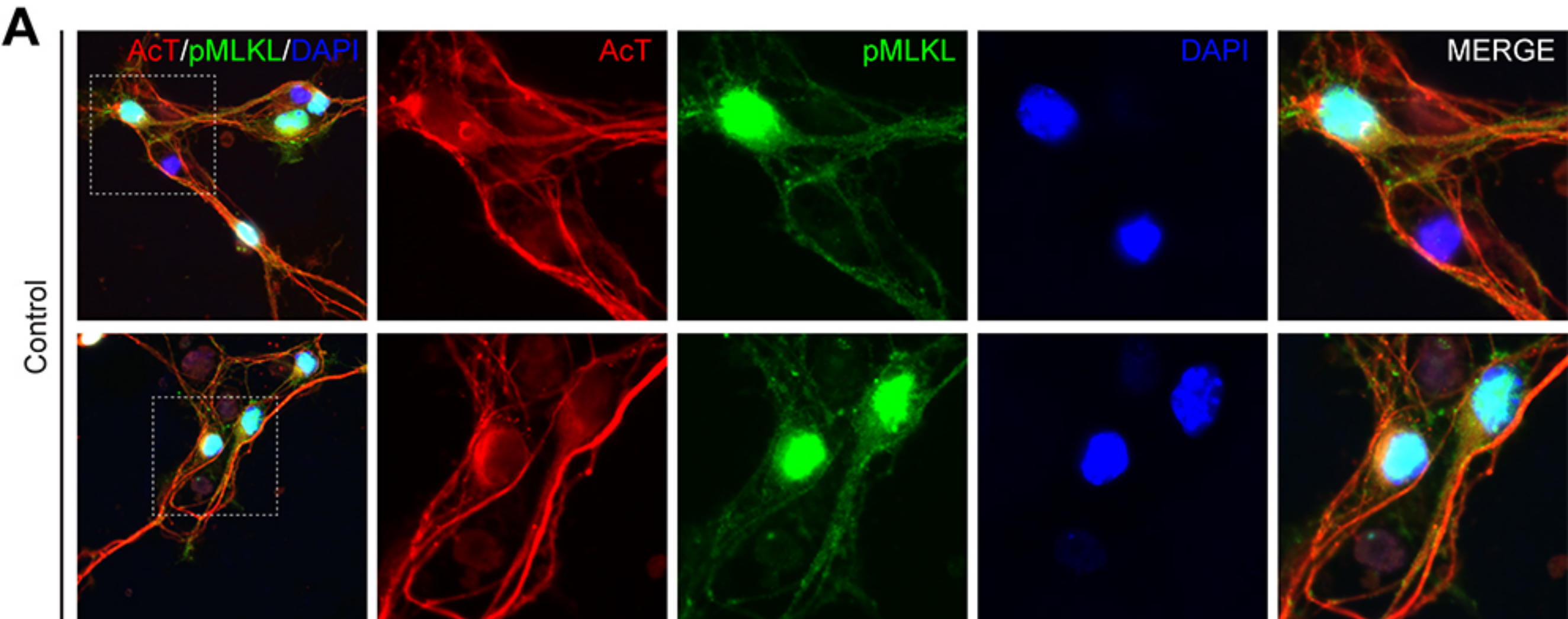
860 MLKL<sup>-/-</sup>, RIPK3<sup>-/-</sup> and WT littermate mice were injected with 6-OHDA in the right striatum.  
861 Contralateral hemisphere was kept non-injected as a control. **(A, B)** Forepaw akinesia after  
862 injection was performed using the cylinder test. Percentage of touches from the paw  
863 contralateral to the injection side was measured in MLKL<sup>-/-</sup> and RIPK3<sup>-/-</sup>, respectively. **(C,**  
864 **D)** Motor performance was tested using the rotarod test by measuring the latency to fall in  
865 an accelerated protocol in each genotype. Data are shown as mean  $\pm$  SEM. Statistical  
866 differences were analyzed using two-way repeated measures ANOVA followed by  
867 Bonferroni's *post hoc* test. \*  $p < 0.05$ .  $n = 8$  animals per group.

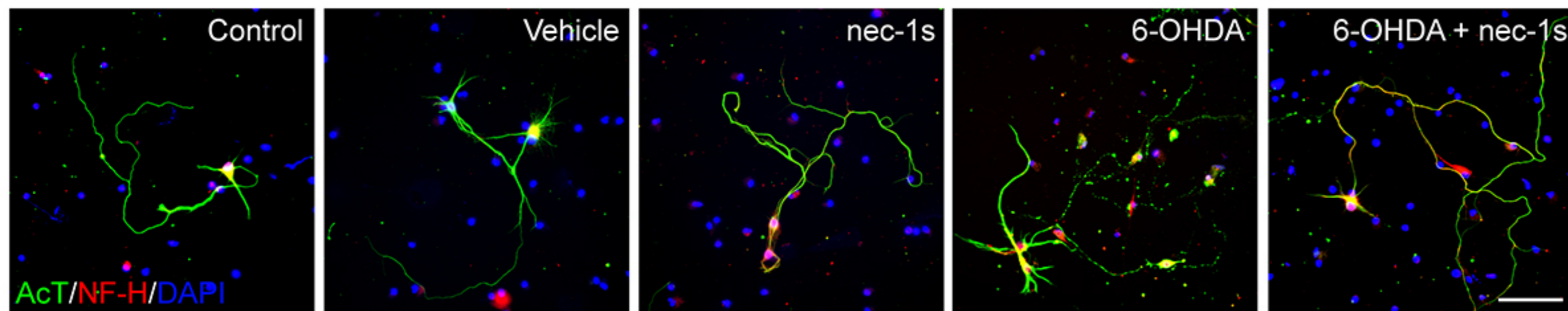
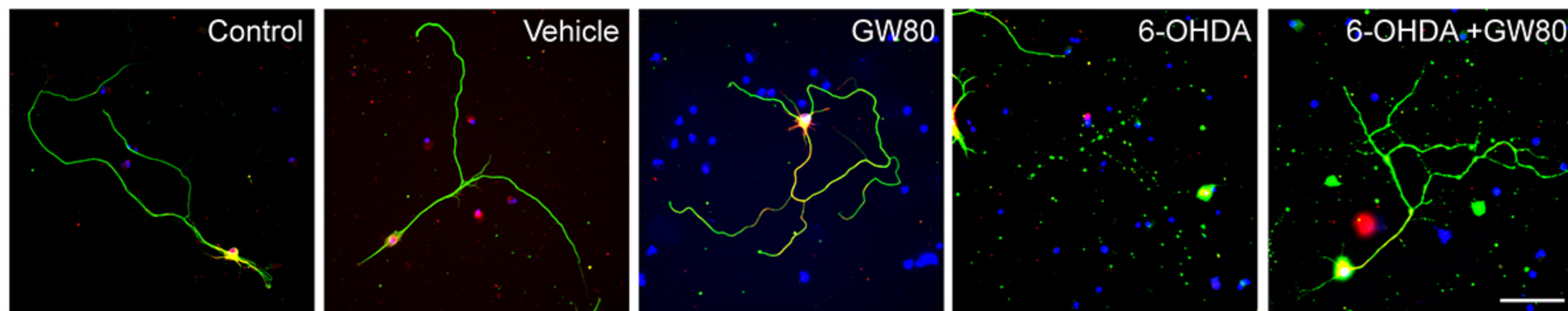
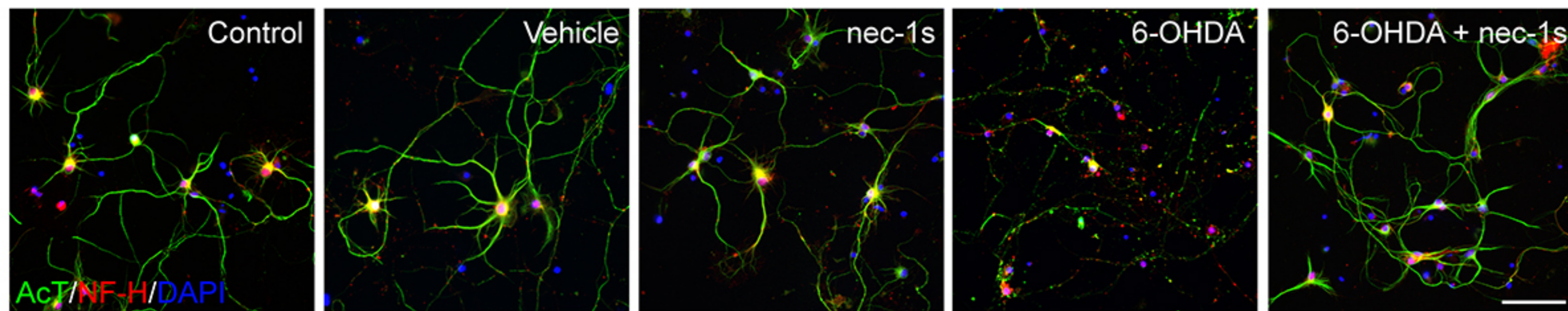
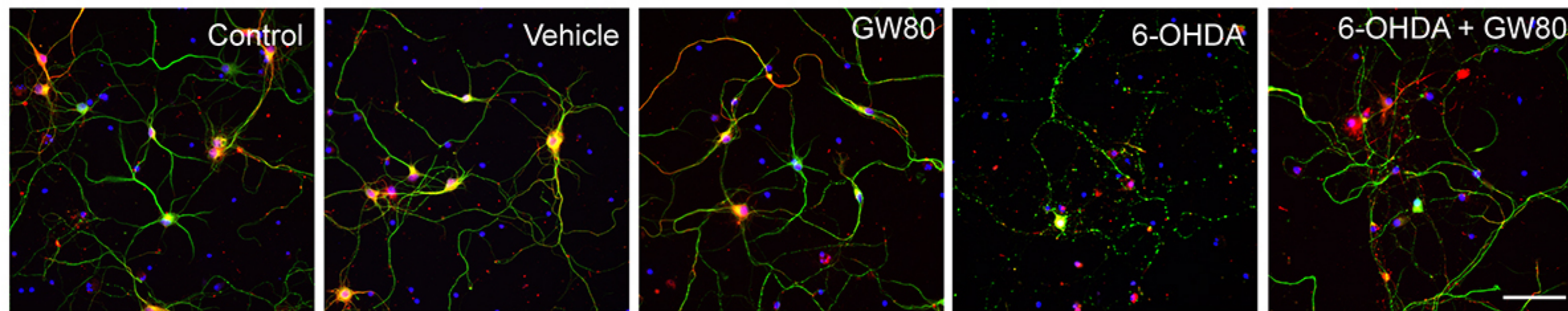
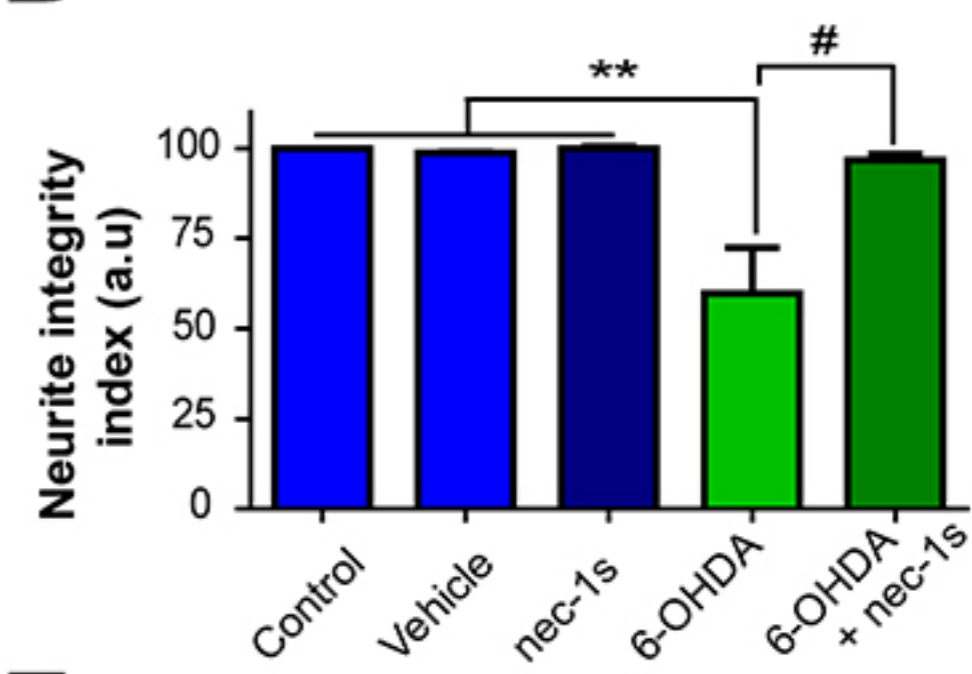
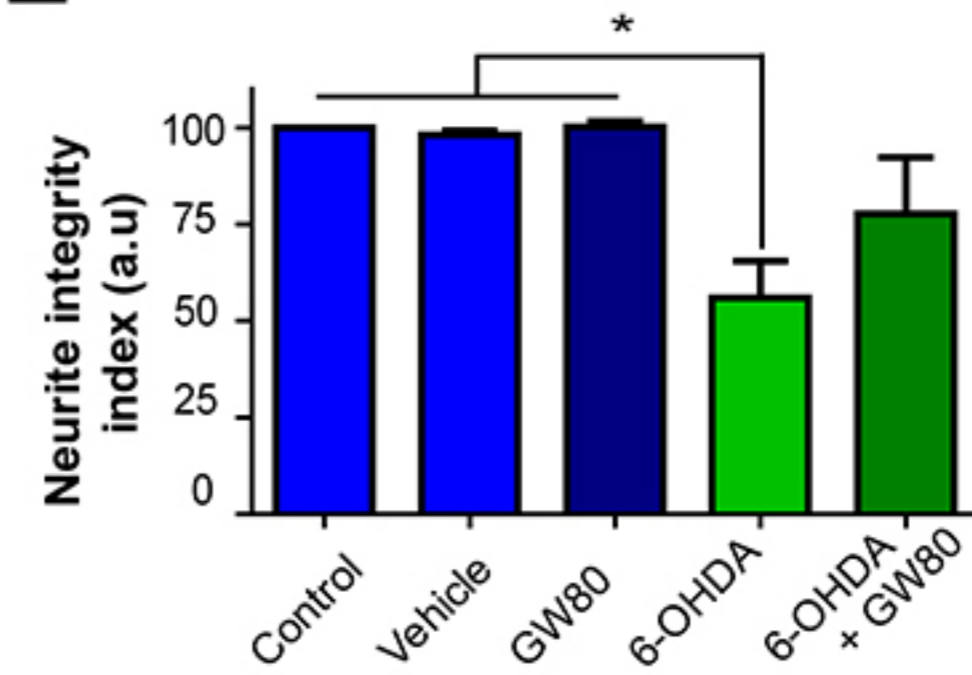
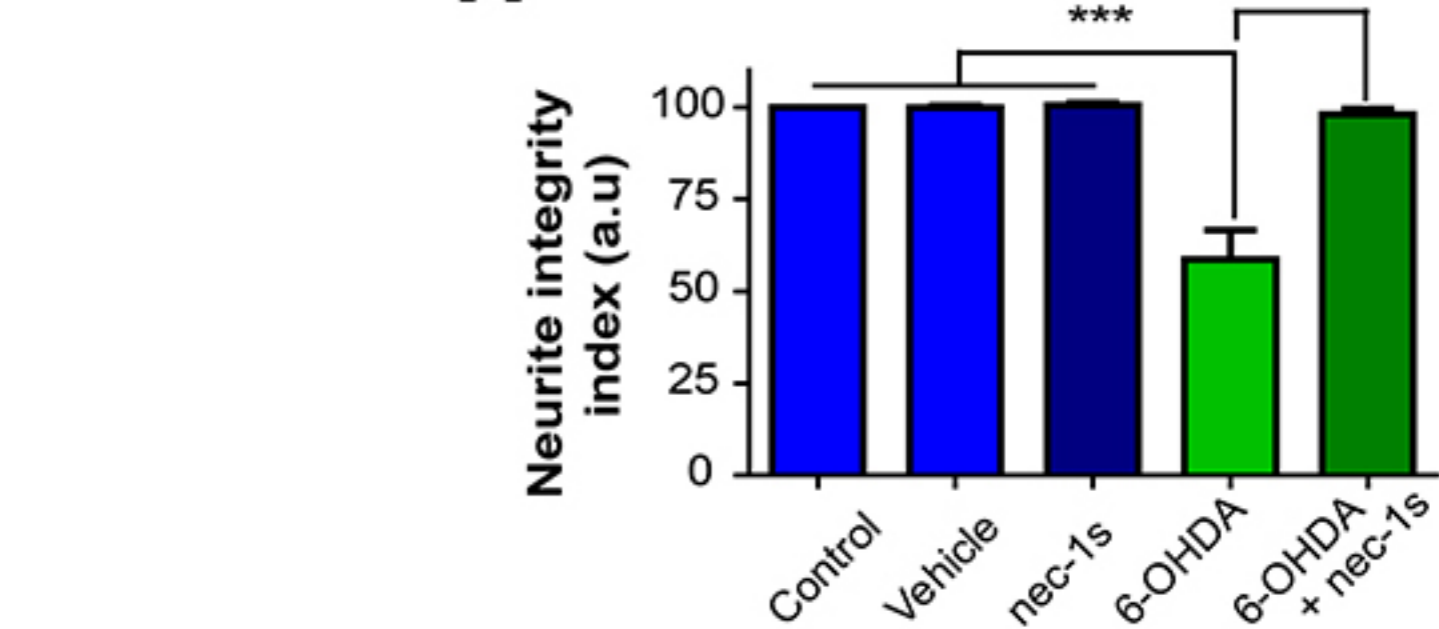
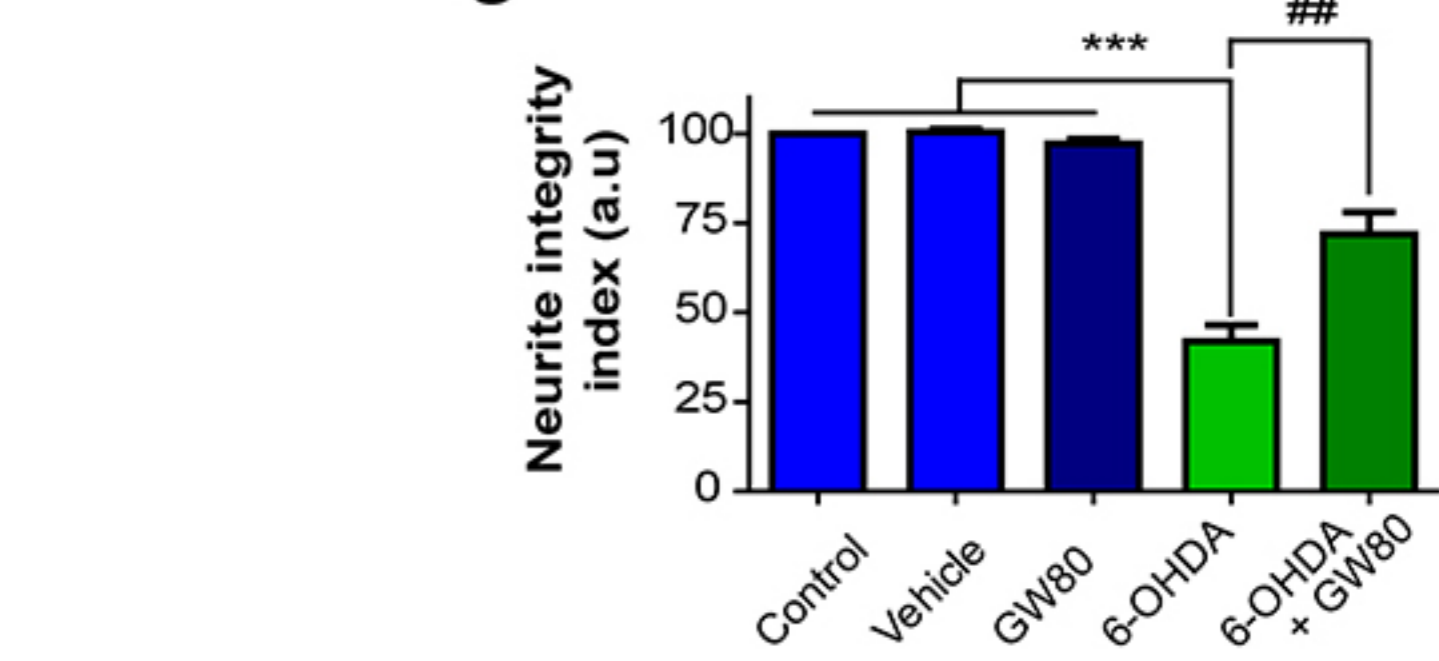
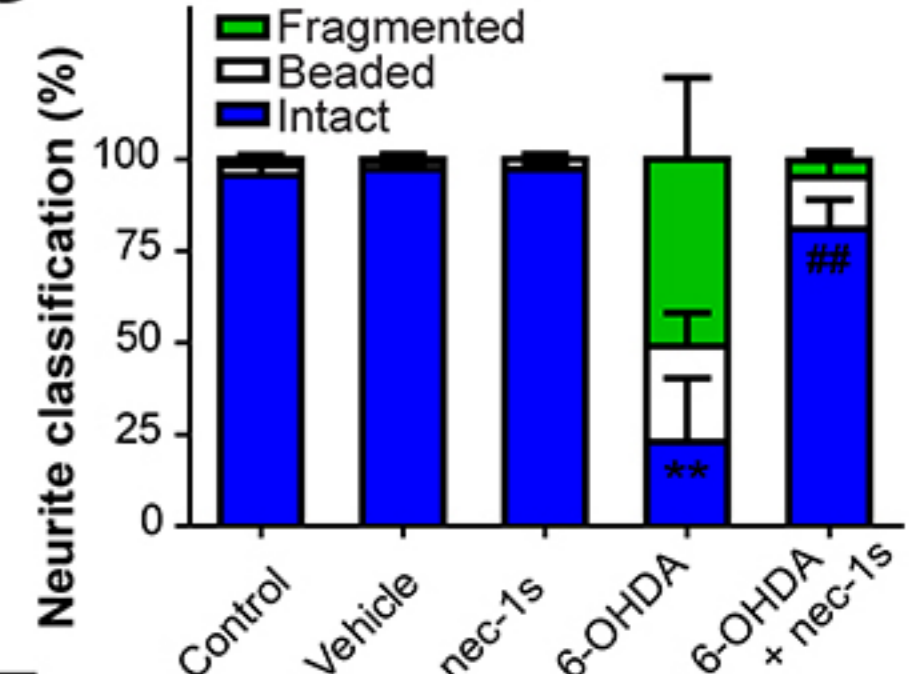
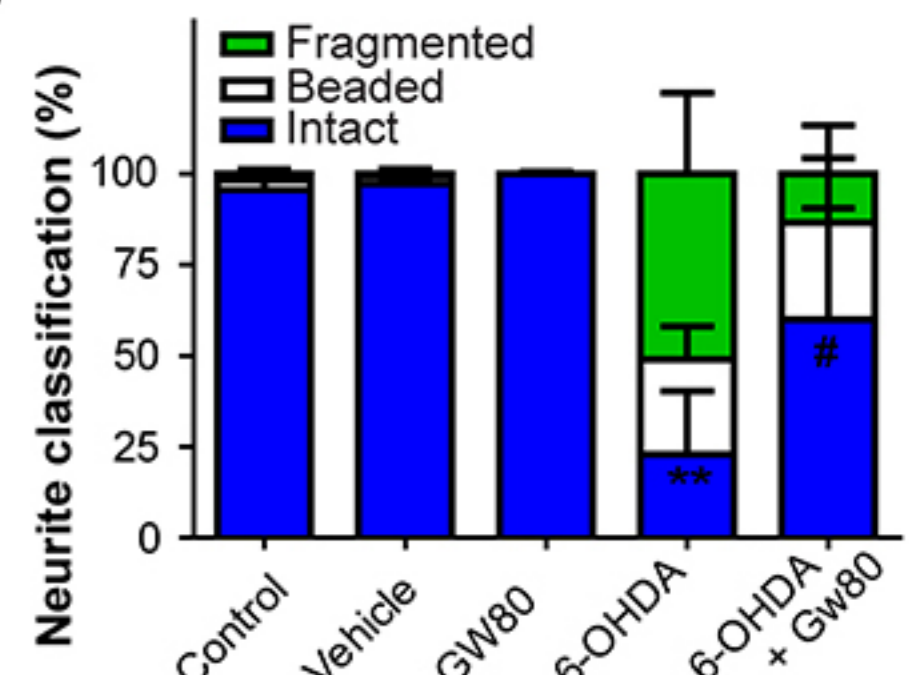
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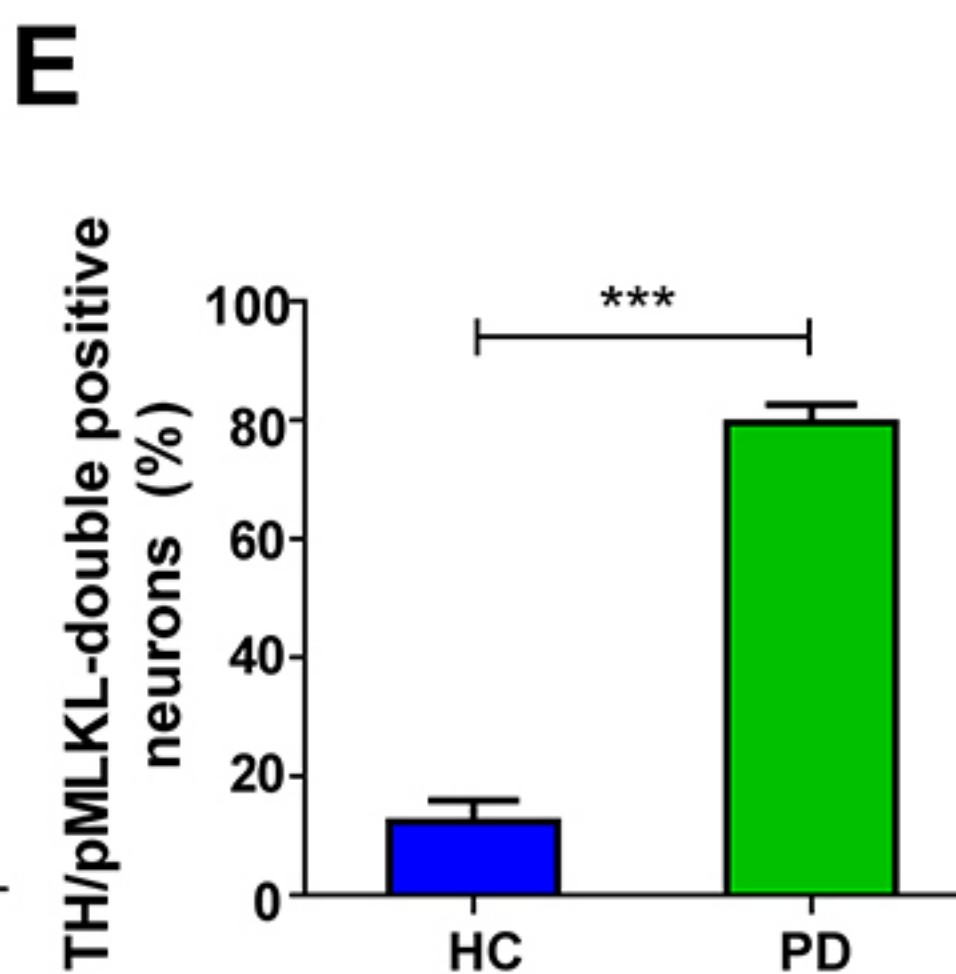
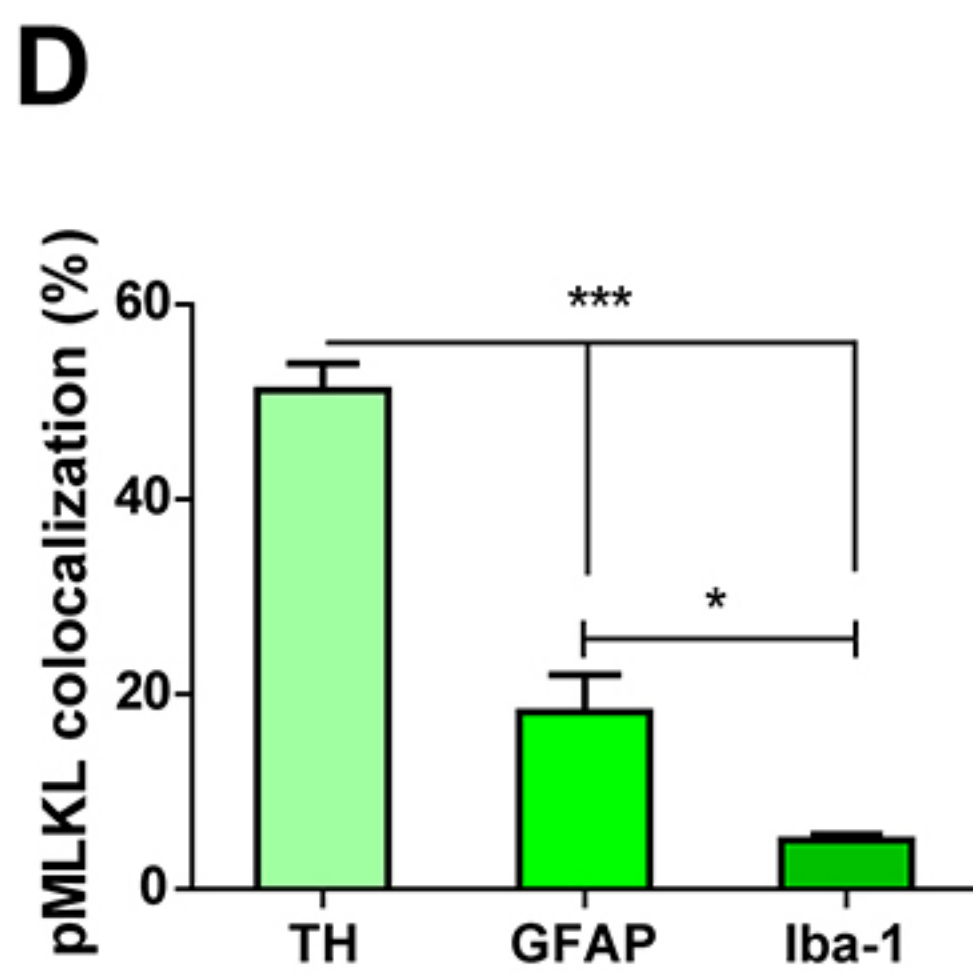
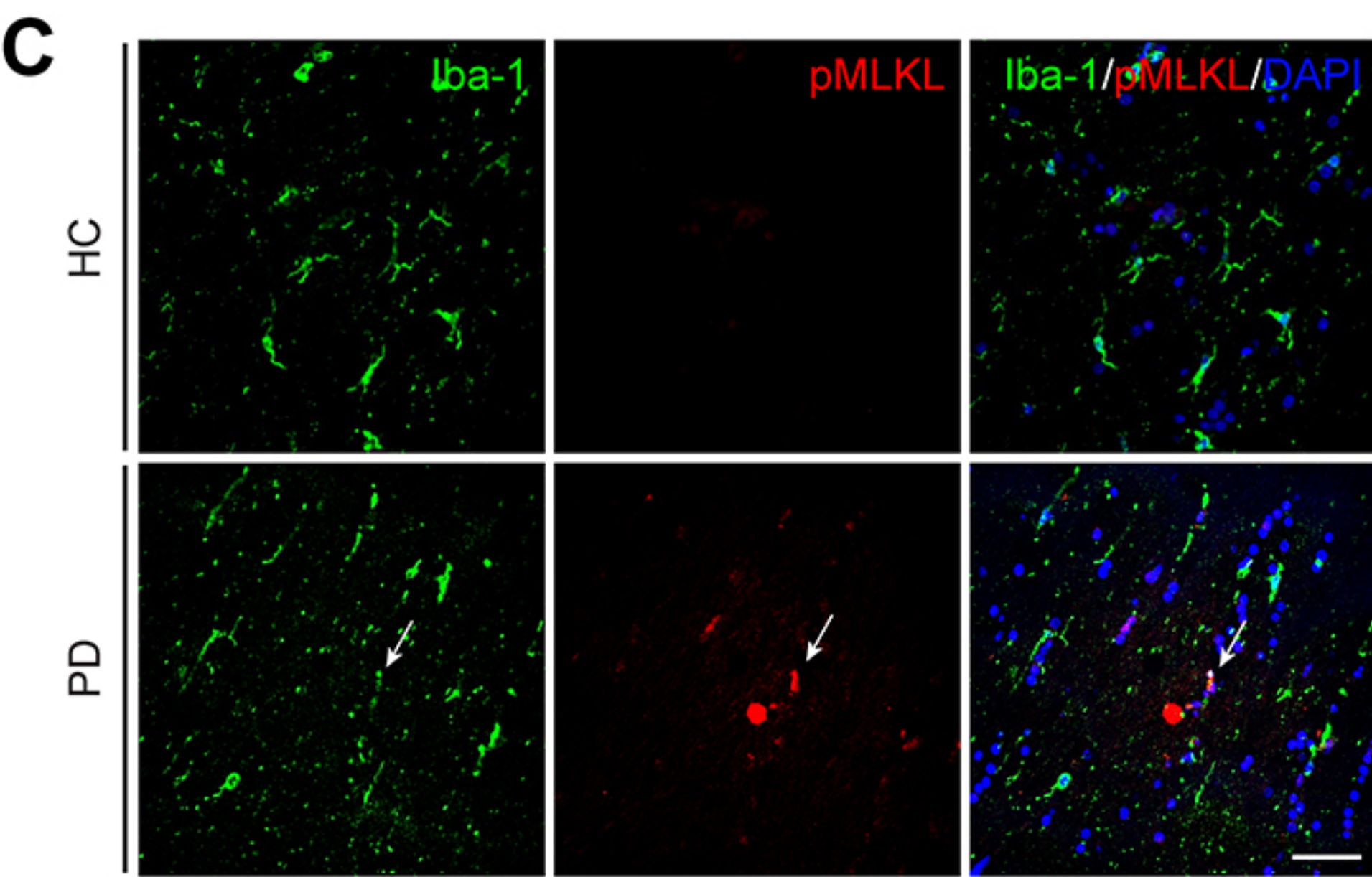
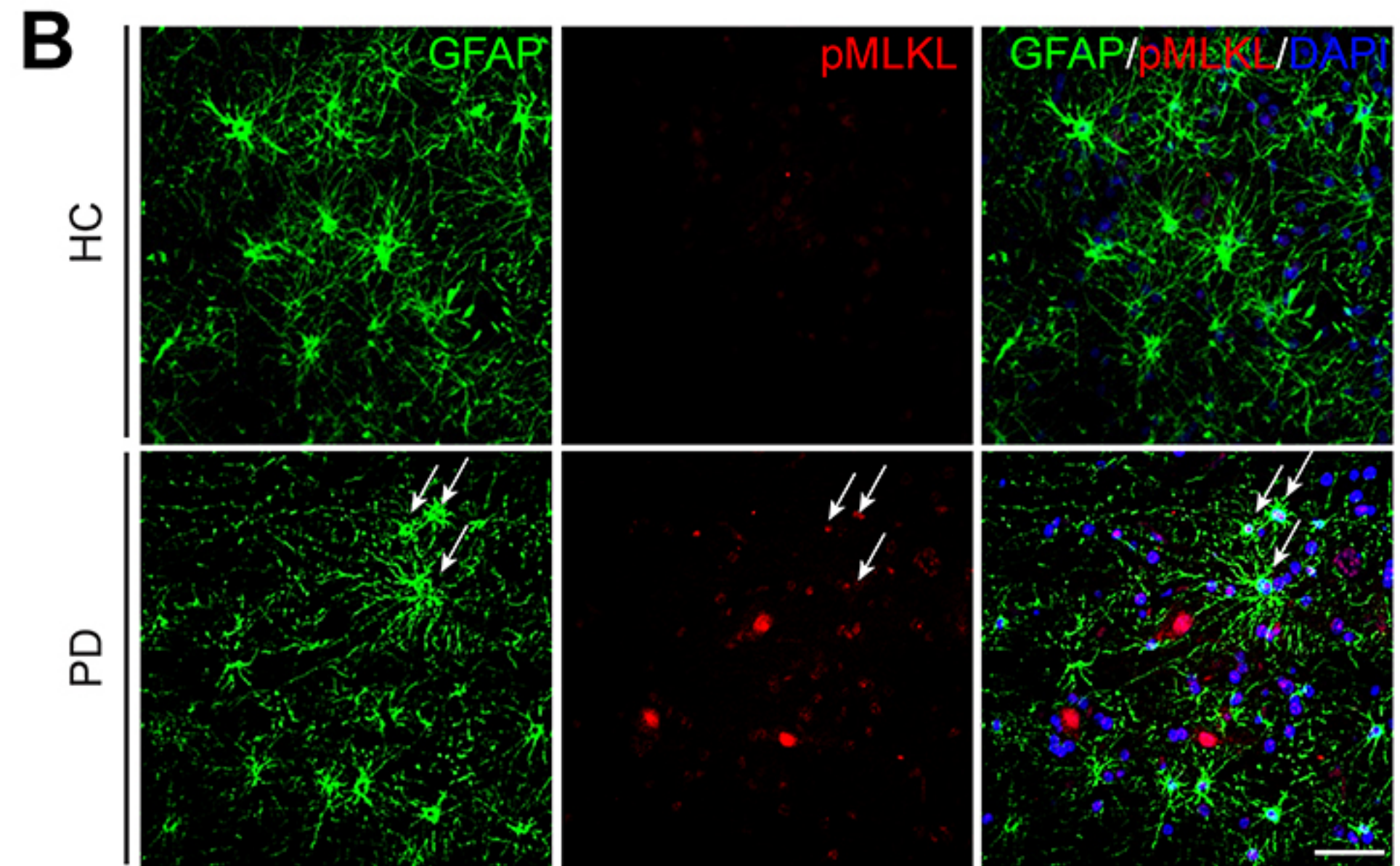
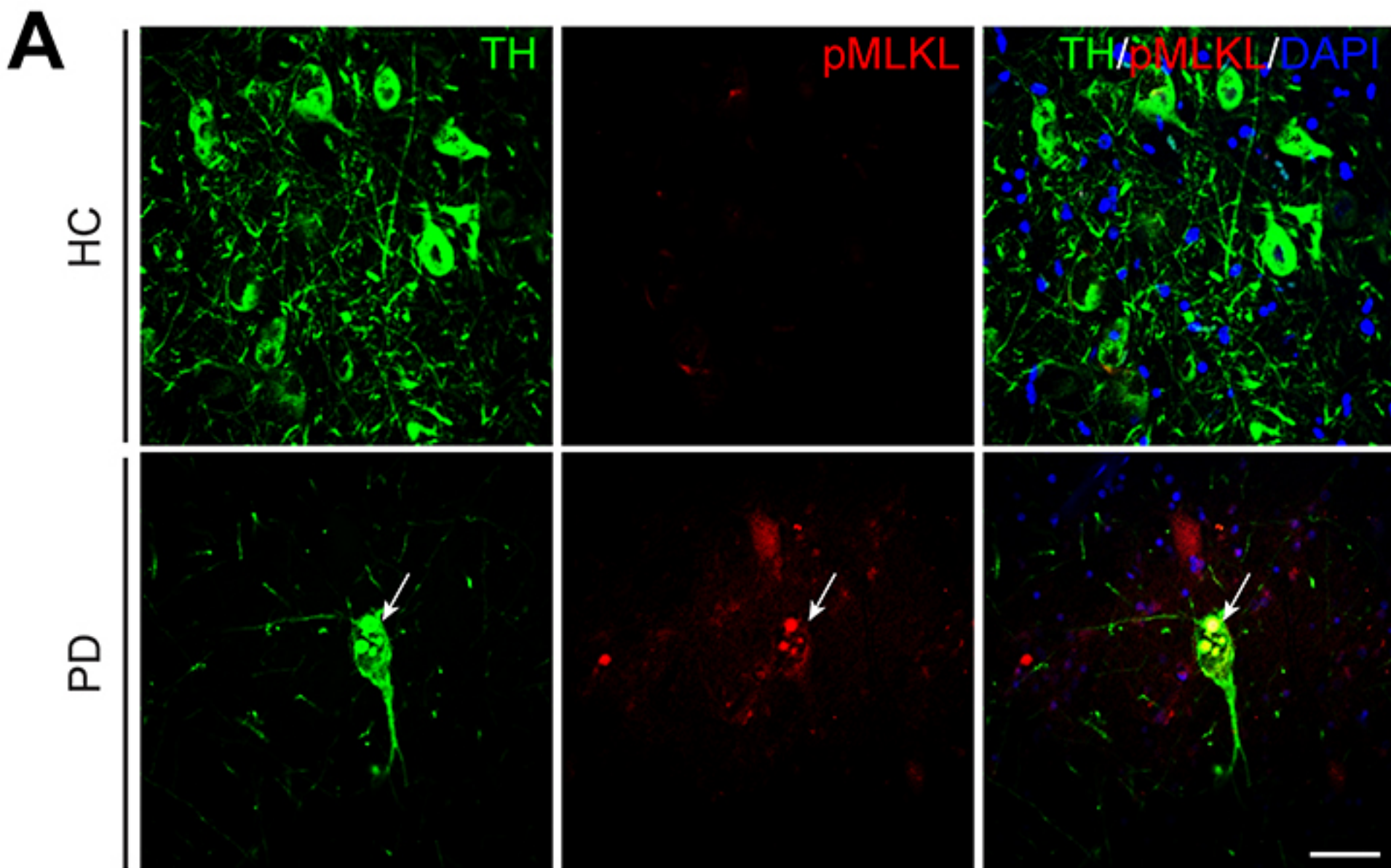
869 **Figure 8. Pharmacological inhibition of RIPK1 contributes to neurodegeneration and**  
870 **motor impairment after 6-OHDA injection.** WT mice were pre-treated with nec-1s for 3

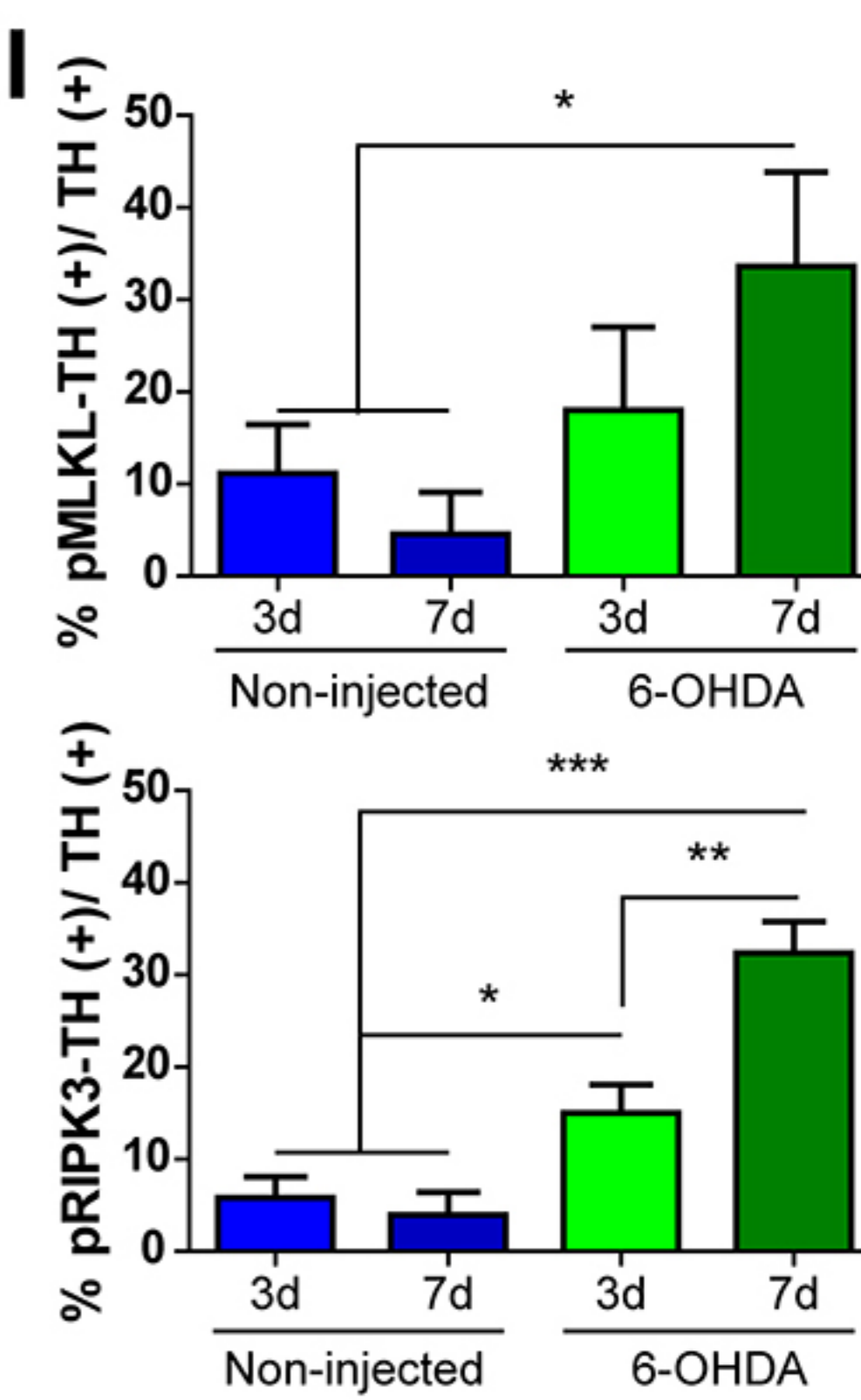
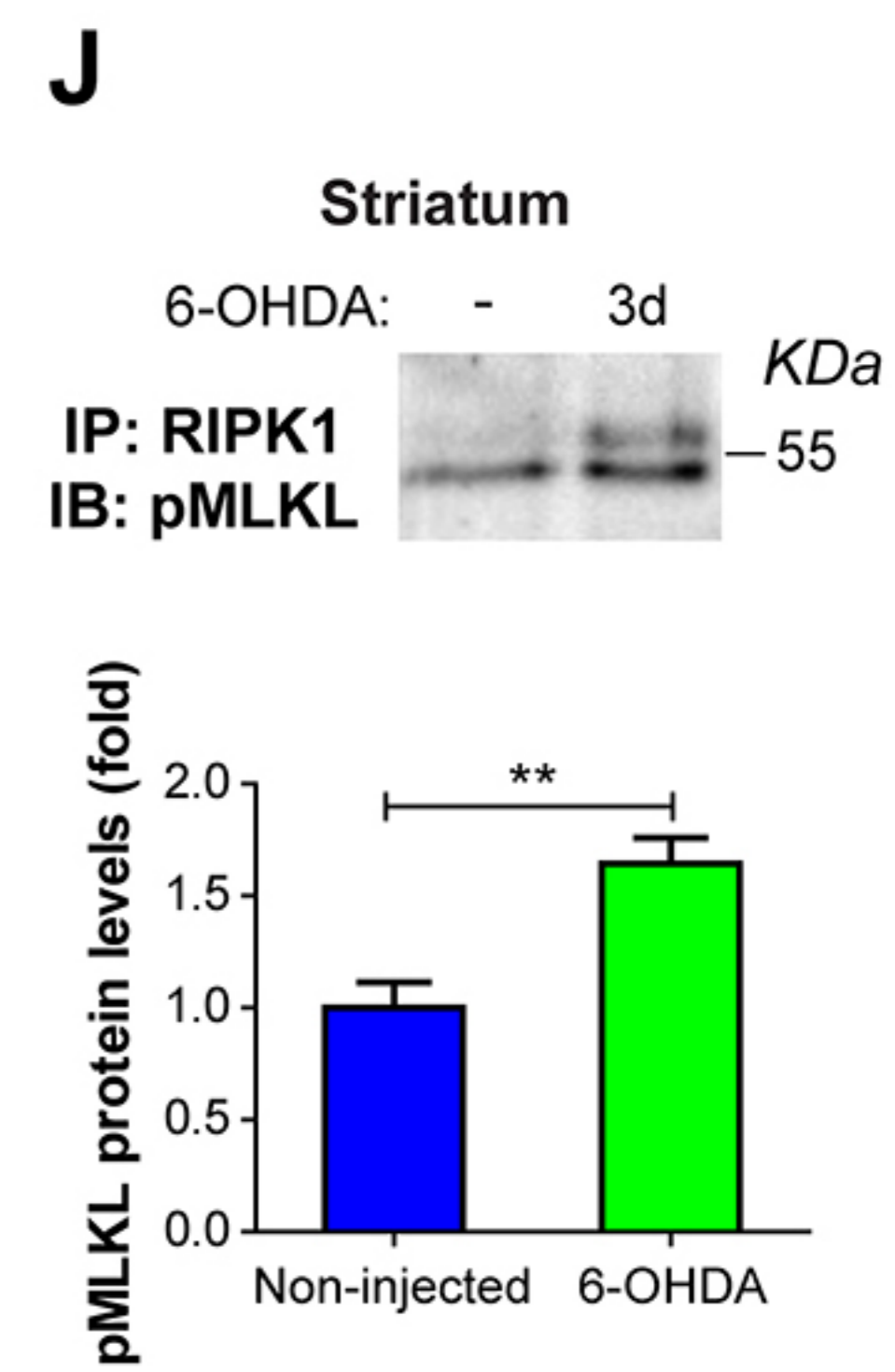
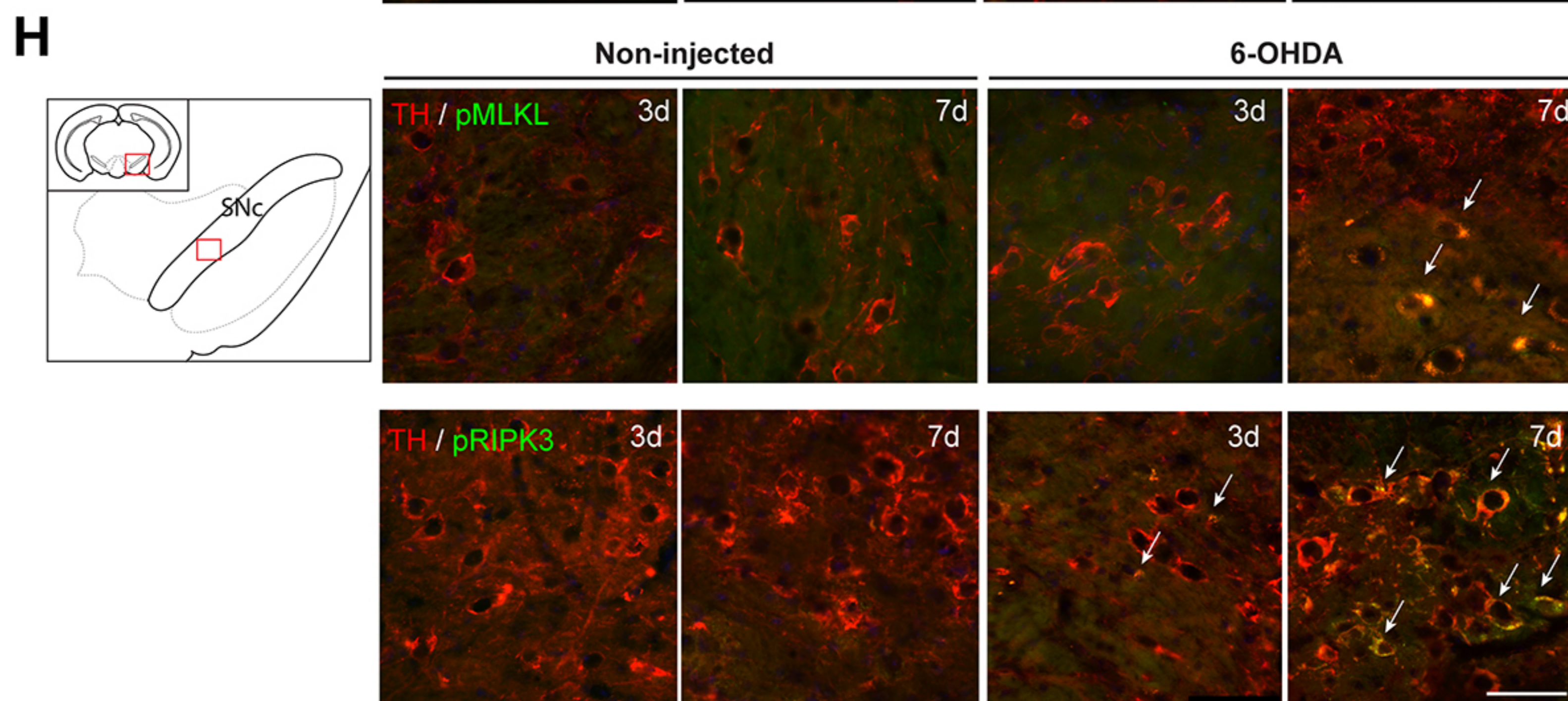
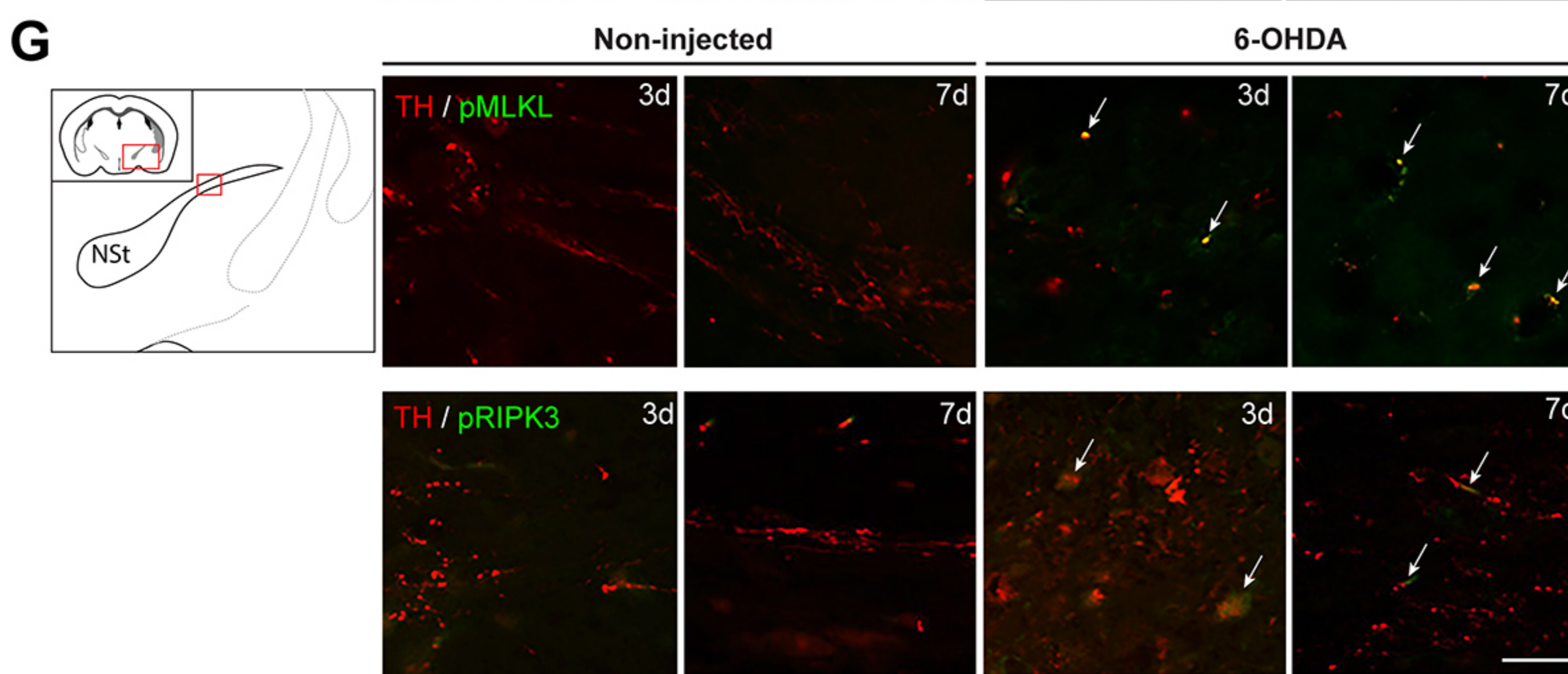
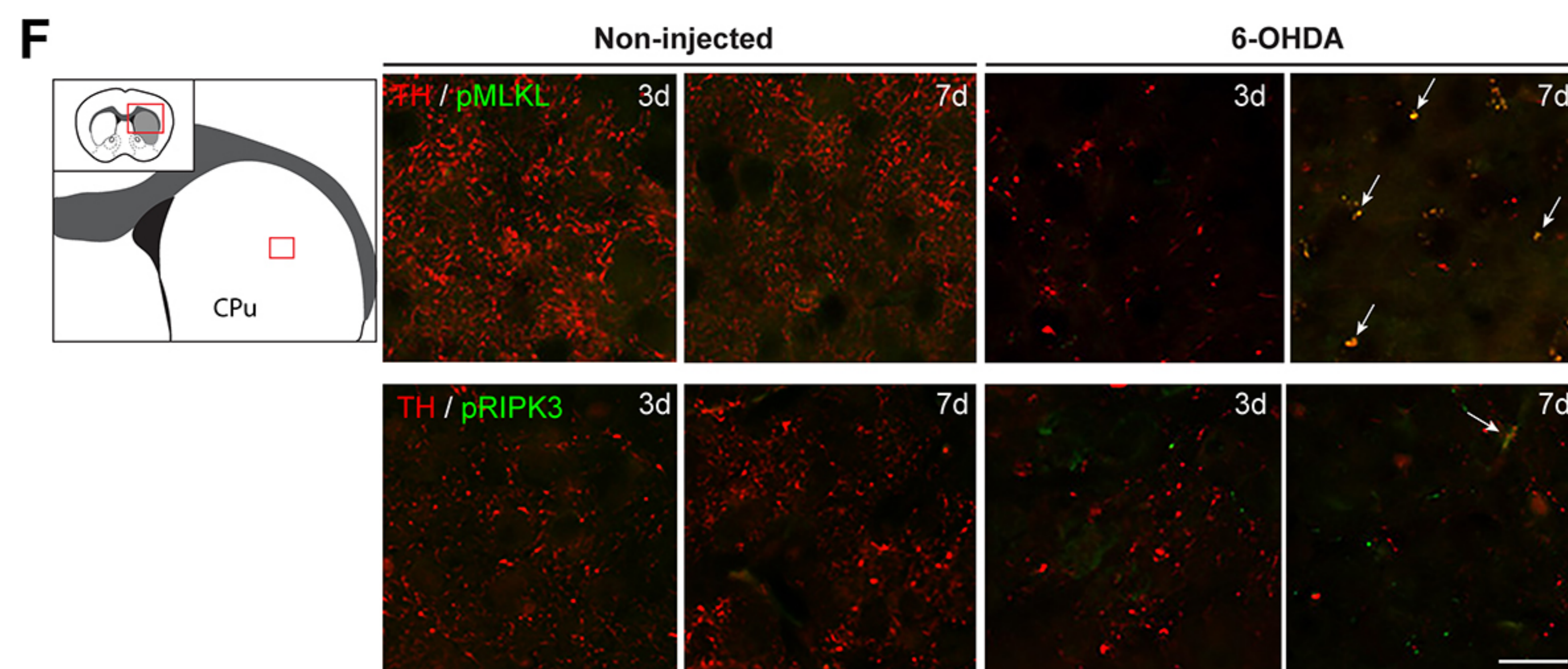
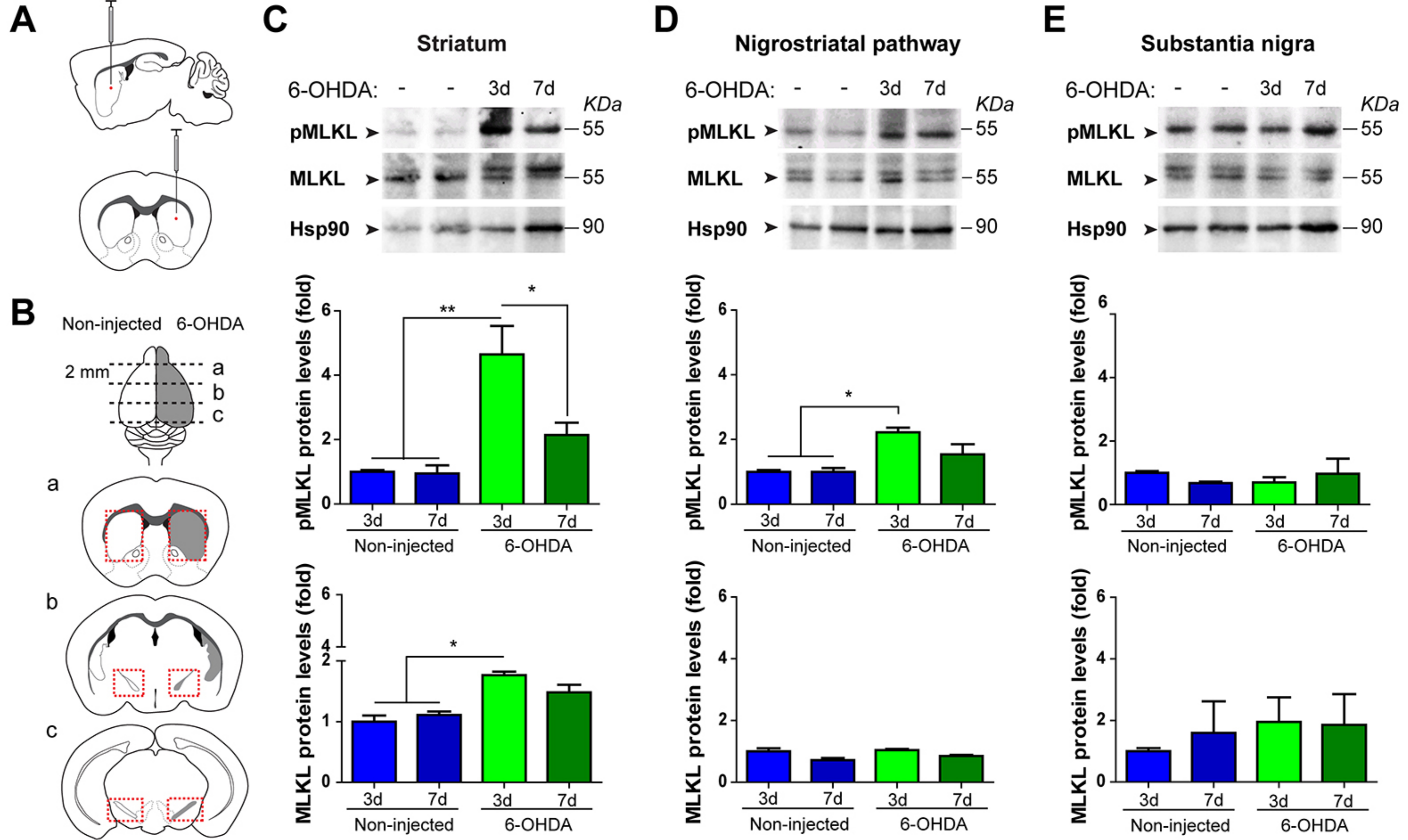
871 days and then unilaterally injected with 6-OHDA in the right striatum. Mice were followed by  
872 7 days daily injections of nec-1s. Vehicle (Veh) treatment was used as a control for the nec-  
873 1s injections. Left striatum was kept non-injected as control. Serial coronal sections of the  
874 entire nigrostriatal circuit were obtained 7 days after 6-OHDA injection and immunostained  
875 for TH. **(A, left)** Scheme of striatal region analyzed. **(A, left)** Representative striatal (CPu)  
876 coronal sections from Veh or nec-1s treated mice unilaterally injected with 6-OHDA. Scale  
877 bar, 1 mm. **(B, left)** Striatal denervation 7 days after 6-OHDA injection was calculated as  
878 the total integrated optical density in non-injected and injected hemispheres from Veh or  
879 nec-1s treated mice. **(B, right)** The percentage of TH loss staining was estimated from  
880 integrated density. **(C, left)** Scheme of nigrostriatal pathway region analyzed. **(C, right)**

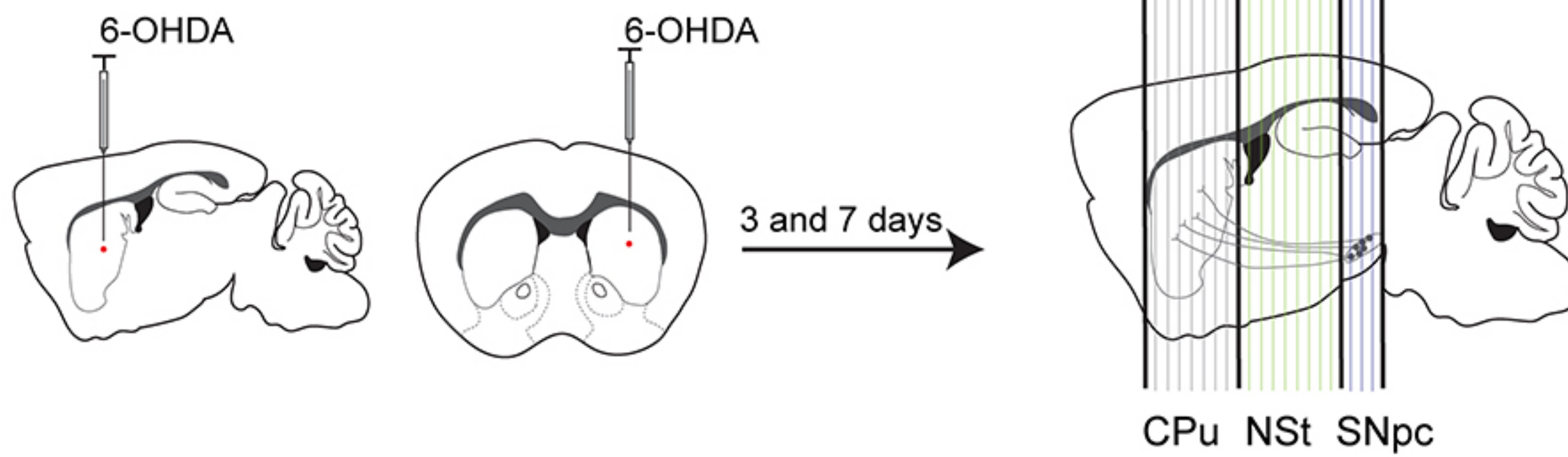
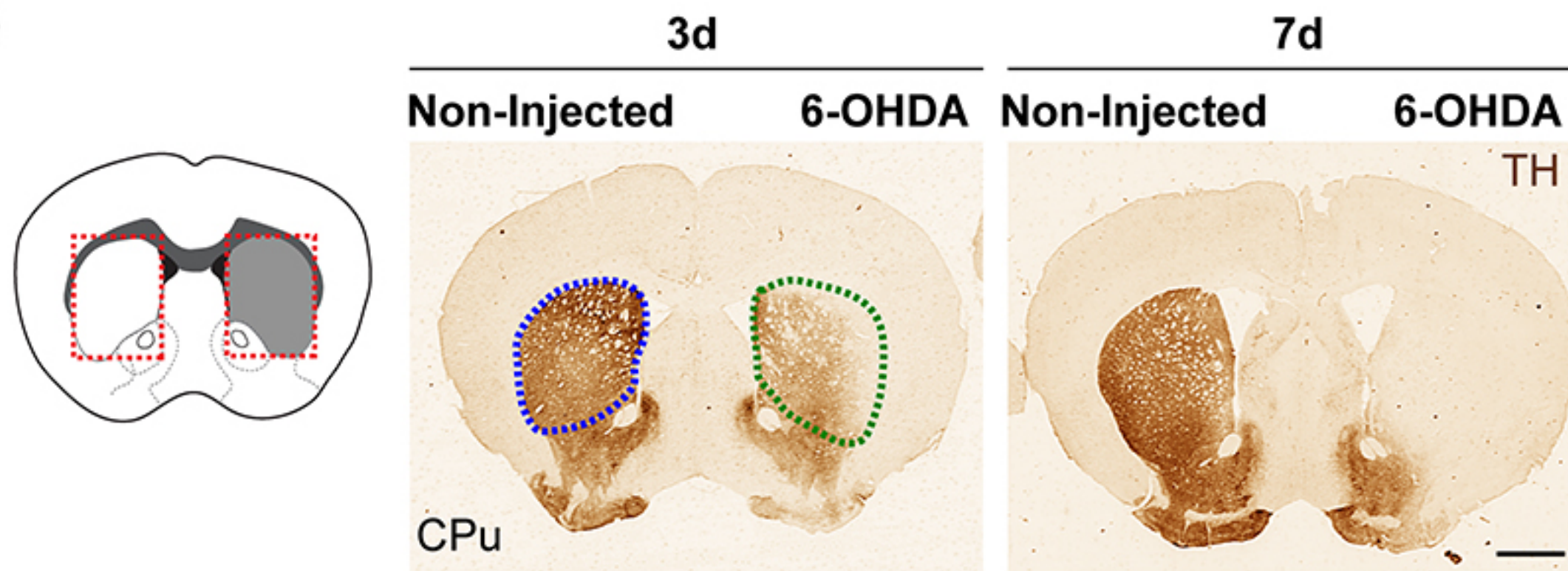
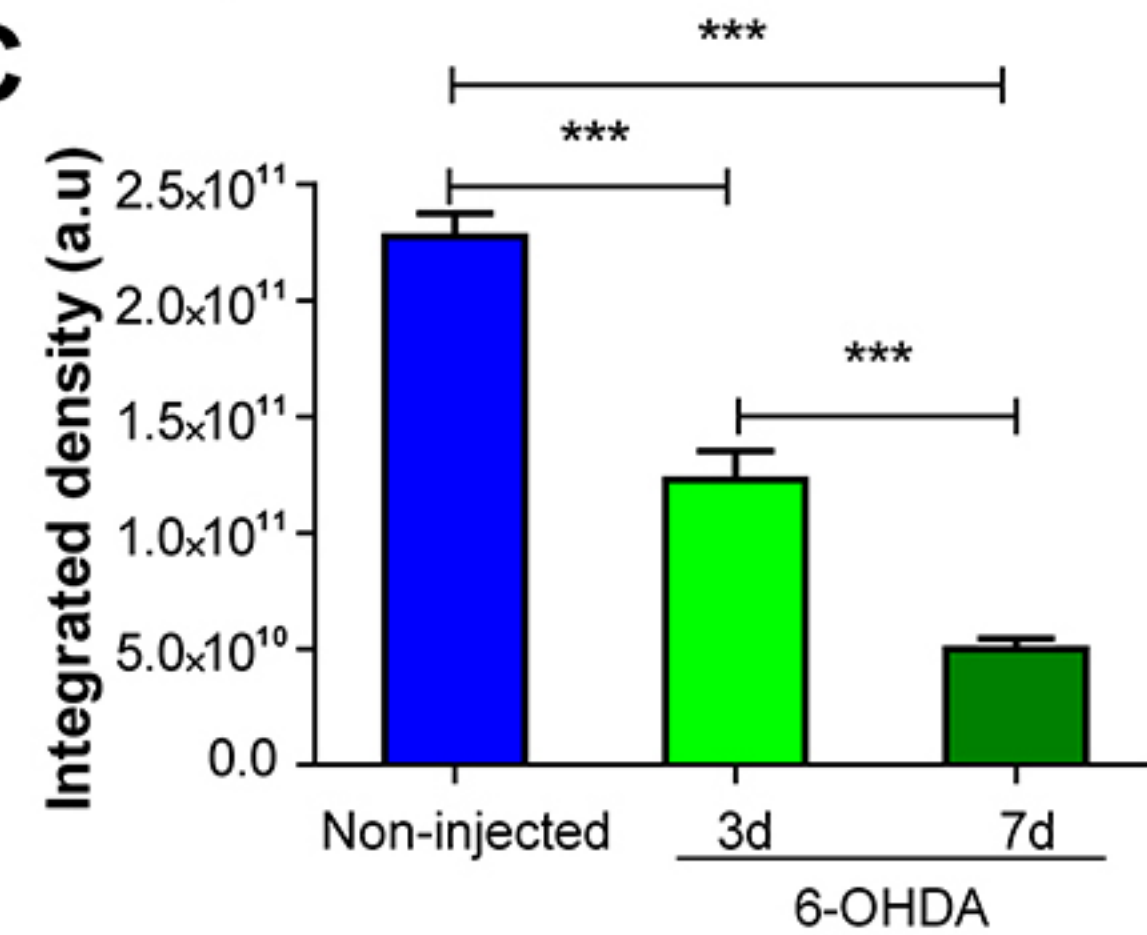
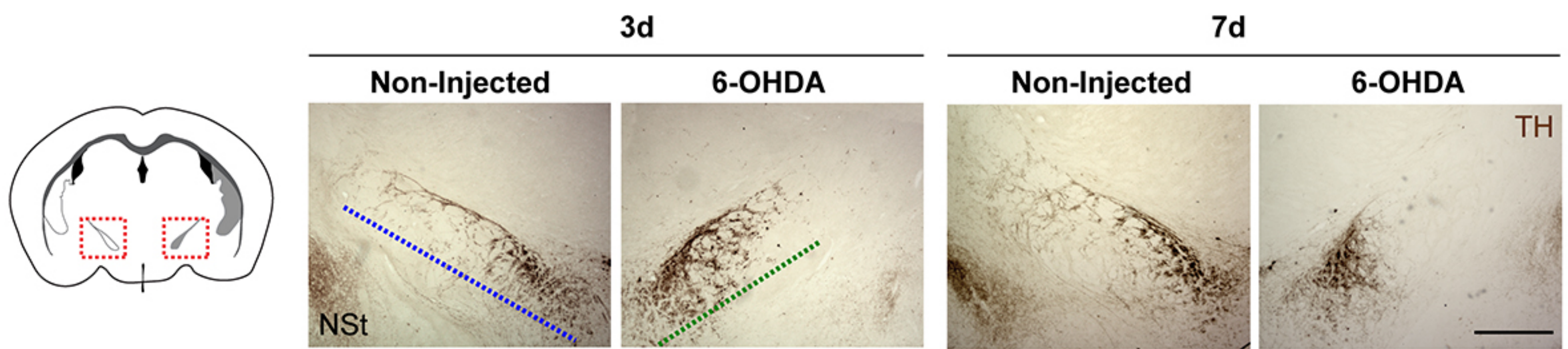
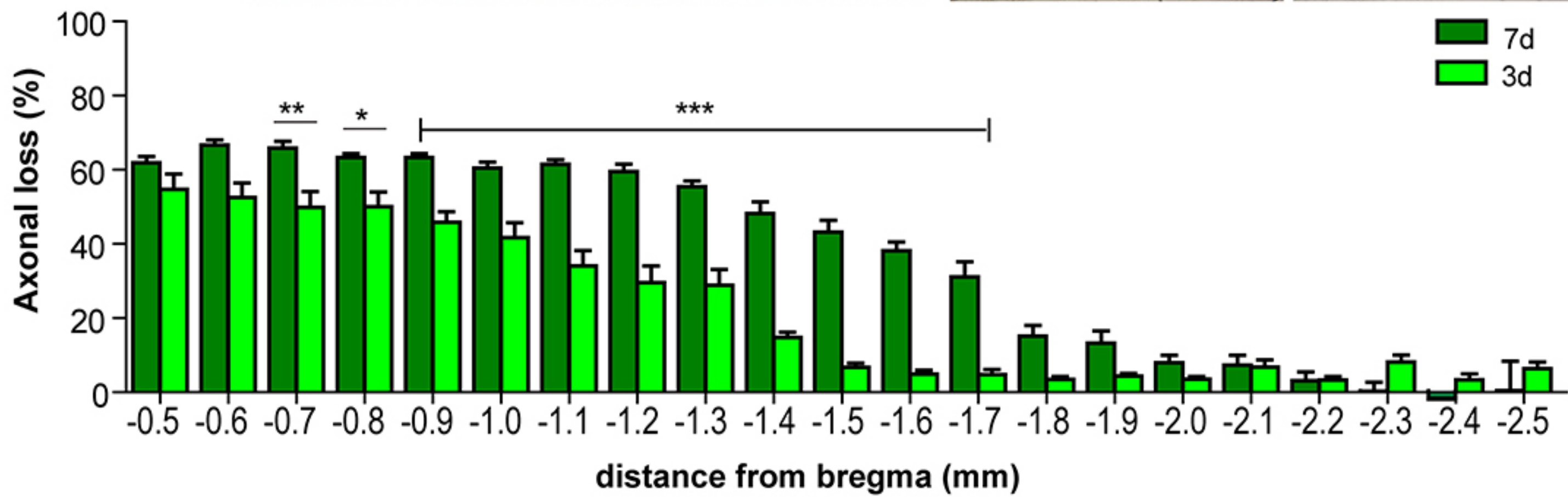
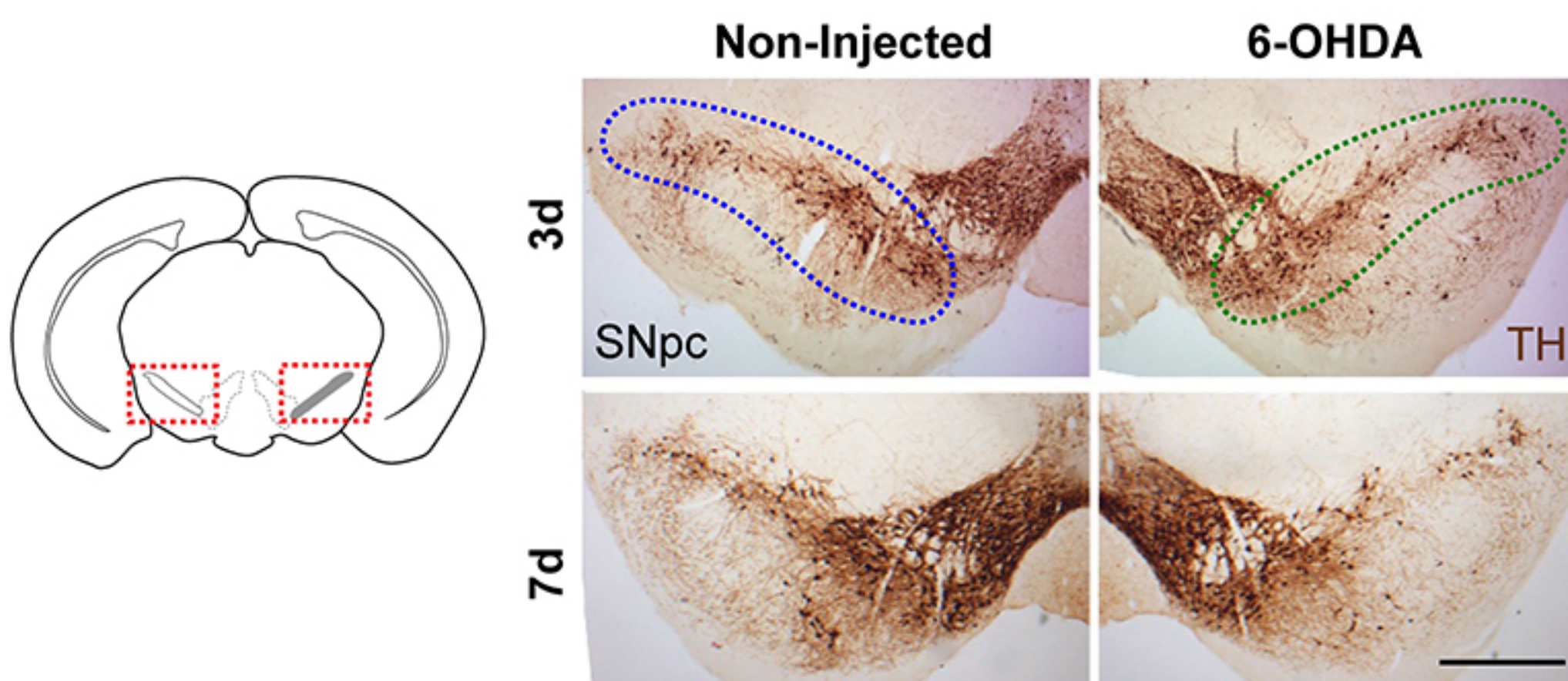
881 Representative images from nigrostriatal axons (NSt) from WT mice unilaterally injected  
882 with 6-OHDA and treated with Veh or nec-1s. Scale bar, 500  $\mu\text{m}$ . **(D)** Spatial distribution of  
883 axonal loss 7 days after 6-OHDA and treated with Veh or nec-1s. **(E)** Forepaw akinesia was  
884 evaluated using the cylinder test. Percentage of touches from the paw contralateral to the  
885 injection side was measured. **(F)** Motor performance was tested using the rotarod test by  
886 measuring the latency to fall in an accelerated protocol. Data are shown as mean  $\pm$  SEM.  
887 Statistical differences were analyzed using two-way ANOVA followed by Bonferroni's *post*  
888 *hoc* test in (B, left), by student's t-test in (B, right), and by two-way repeated measures  
889 ANOVA followed by Bonferroni's *post hoc* test in (B, C and D). \*  $p < 0.05$ ; \*\*  $p < 0.01$ ; \*\*\*  $p$   
890  $< 0.001$ .  $n = 9$  animals per group.



**A****D****G****I****B****E****H****J****C****F**





**A****B****C****D****E****F****G**