**Title**

Dopamine regulates astrocytic IL-6 expression and process formation via dopamine receptors and adrenoceptors

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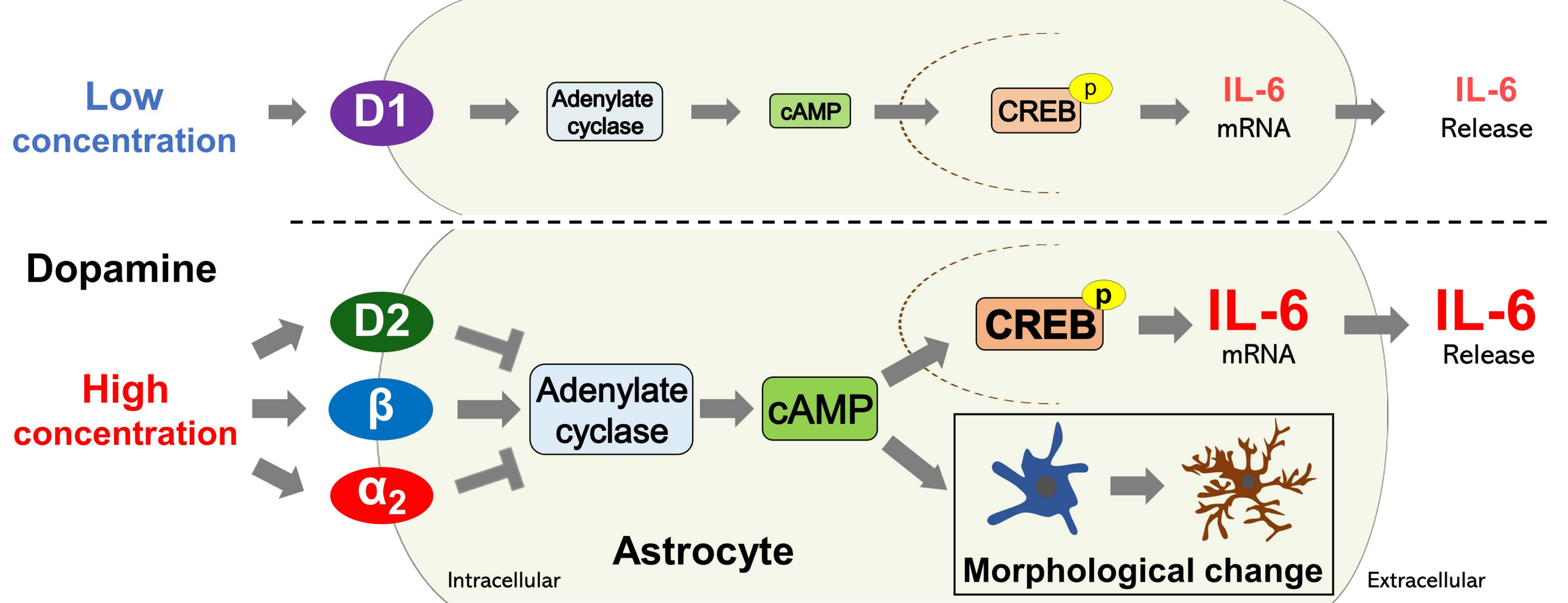
19 **Abstract**

20 Dopamine levels in the central nervous system change under pathological
21 conditions such as Parkinson's disease, Huntington's disease, and addiction. Under those
22 pathological conditions, astrocytes become reactive astrocytes characterized by
23 morphological changes and the release of inflammatory cytokines involved in pathogenesis.
24 However, it remains unclear whether dopamine regulates astrocytic morphology and
25 functions. Elucidating these issues will help us to understand the pathogenesis of
26 neurodegenerative diseases caused by abnormal dopamine signaling. In this study, we
27 investigated the effects of dopamine on IL-6 expression and process formation in rat
28 primary cultured astrocytes and acute hippocampal slices. Dopamine increased IL-6
29 expression in a concentration-dependent manner, and this was accompanied by CREB
30 phosphorylation. The effects of a low dopamine concentration (1 μM) were inhibited by a
31 D1-like receptor antagonist, whereas the effects of a high dopamine concentration (100
32 μM) were inhibited by a β-antagonist and enhanced by a D2-like receptor antagonist.
33 Furthermore, dopamine (100 μM) promoted process formation, which was inhibited by a
34 β-antagonist and enhanced by both an α-antagonist and a D2-like receptor antagonist. In
35 acute hippocampal slices, both a D1-like receptor agonist and β-agonist changed astrocytic
36 morphology. Together, these results indicate that dopamine promotes IL-6 expression and
37 process formation via D1-like receptors and β-adrenoceptors. Furthermore, bidirectional

regulation exists; namely, the effects of D1-like receptors and β-adrenoceptors were negatively regulated by D2-like receptors and $\alpha_2$-adrenoceptors.

**Keywords**

astrocyte, dopamine, adrenoceptor, IL-6, CREB, morphology


Low
concentration
D1
Adenylate
cyclase
cAMP
CREB
p
IL-6
mRNA
IL-6
Release
Dopamine
High
concentration
D2
β
α2
Adenylate
cyclase
cAMP
CREB
p
IL-6
mRNA
IL-6
Release
Morphological change
Astrocyte
Intracellular
Extracellular

44 **1. Introduction**

45 Dopamine is a key neurotransmitter in the central nervous system (CNS) and
46 regulates many brain functions (Klein et al., 2019). Dopaminergic neurons project to most
47 regions of the CNS, including the cerebral cortex, hippocampus, and spinal cord
48 (Descarries et al., 1987; Edelmann and Lessmann, 2018; Ridet et al., 1992). Dopamine is
49 released not only from synapses but also from varicosities (Fuxe et al., 2015), and thus
50 acts on astrocytes, a type of glial cell that surrounds neurons. Astrocytic cytokine
51 production and the morphology of astrocytic processes are closely related to the
52 physiological functions of the CNS. In the CNS, astrocytes represent the major source of
53 interleukin-6 (IL-6) (Gruol and Nelson, 1997), which suppresses neuronal cell death (Day
54 et al., 2014) and acts as a neurotrophic factor (Wagner, 1996). Astrocytic processes contact
55 neurons, forming the "tripartite synapse" that regulates synaptic function (Allen and
56 Eroglu, 2017).

57 Dopamine levels in the CNS change under pathological conditions such as
58 Parkinson's disease, Huntington's disease, and addiction (Klein et al., 2019). Furthermore,
59 astrocytes transform into reactive astrocytes in response to a wide range of
60 neurodegenerative diseases (Hart and Karimi-Abdolrezaee, 2021; Pekny and Nilsson,
61 2005). Reactive astrocytes are characterized by upregulated IL-6 and glial fibrillary acidic
62 protein (GFAP) expression and distinct morphological changes (Escartin et al., 2021; John

63 et al., 2003; Sofroniew, 2009), and regarded as reflections of a detrimental astrocyte
64 phenotype, which contribute to various pathogeneses (Escartin et al., 2021). Chronic IL-6
65 overexpression in astrocytes induces an inflammatory response (Penkowa et al., 2003).
66 Impaired astrocytic morphogenesis links to diminished function of excitatory synapses
67 (Stogsdill et al., 2017), and astrocytic morphology regulates scar formation, facilitating
68 recovery from traumatic brain injury (Schiweck et al., 2021). Furthermore, the activation
69 of astrocytic dopamine receptors regulates neuroinflammation (Montoya et al., 2019; Zhu
70 et al., 2018) and depresses excitatory synaptic transmission (Corkrum et al., 2020). These
71 reports suggest that dopamine is involved in the pathogenesis of CNS diseases via its
72 action on astrocytes.

73 Astrocytes express D1-like receptors (D1, D5), D2-like receptors (D2-D4)
74 (Miyazaki et al., 2004), $\alpha_1$-, $\alpha_2$-, and β-adrenoceptors (Hertz et al., 2010). Dopamine has a
75 low affinity for adrenoceptors (Zhang et al., 2004). High dopamine concentrations act on
76 β-adrenoceptors in astrocytes, activating brain-derived neurotrophic factor (BDNF)
77 transcription and changing cell morphology (Koppel et al., 2018). Our previous reports
78 have demonstrated that noradrenaline acts on astrocytic β-adrenoceptors to enhance IL-6
79 transcription and change cell morphology (Kitano et al., 2021; Morimoto et al., 2021).
80 However, it remains unknown whether dopamine affects astrocytic IL-6 production or
81 morphology and, if so, which receptors are involved. Such knowledge could enhance our

82 understanding of the role that dopamine plays in astrocytic functions and the
83 pathogenesis of neurodegenerative diseases caused by abnormal dopamine signaling. We
84 aimed to comprehensively identify IL-6 expression and morphological changes as a
85 phenotype of reactive astrocytes.

86 In this study, we investigated dopamine-induced IL-6 expression and
87 morphological changes in rat cultured astrocytes and acute brain slices. The results
88 revealed the concentration-dependent effects of dopamine acting via dopamine and
89 adrenergic receptors and its intracellular mechanisms.

90

91 **2. Materials and Methods**

92

93 **2.1. Materials**

94 Antibodies against the following were used: ERK 1/2 (extracellular signal-
95 regulated kinase 1/2) (#4695S, 1:2500), phospho-ERK 1/2 (#9101S, 1:2500), p38 (#9212S,
96 1:2000), phospho-p38 (#9211S, 1:1000), SAPK/JNK (c-jun N-terminal kinase) (#9252S,
97 1:2500), phospho-SAPK/JNK (#9251S, 1:1500), STAT3 (Signal transducer and activator of
98 transcription 3) (#4904S, 1:4000), and phospho-STAT3 (#9145S, 1:2000) (all from Cell
99 Signaling Technology, Danvers, MA, USA); CREB (cAMP response element-binding
100 protein) (#sc-377154, 1:500) and phospho-CREB (#sc-81486, 1:250) (both from Santa Cruz
101 Biotechnology, Santa Cruz, CA, USA); and GFAP (#11051, 1:200, Immuno-Biological
102 Laboratories, Gunma, Japan).

103 The following reagents were used: atenolol, atipamezole hydrochloride, 2-bromo-
104 α-ergocryptine methanesulfonate salt (bromocriptine), and isoproterenol hydrochloride
105 (all from Sigma-Aldrich, St. Louis, MO, USA); propranolol hydrochloride, forskolin,
106 histamine dihydrochloride, and L(+)-ascorbic acid (all from FUJIFILM Wako Pure
107 Chemical, Osaka, Japan); 1-phenyl-2,3,4,5-tetrahydro-1H-3-benzazepine hydrochloride
108 (SCH23390), (±)-6-chloro-2,3,4,5-tetrahydro-1-phenyl-1H-3-benzazepine hydrobromide
109 (SKF81297), 6-chloro-2,3,4,5-tetrahydro-1-(3-methylphenyl)-3-(2-propenyl)-1H-3-

110 benzazepine-7,8-diol hydrobromide (SKF83822), and 6-chloro-2,3,4,5-tetrahydro-3-
111 methyl-1-(3-methylphenyl)-1H-3-benzazepine-7,8-diol (SKF83959) (all from Tocris
112 Bioscience, Bristol, UK); ICI118551 hydrochloride and SR59230A (both from
113 MedChemExpress, Monmouth Junction, NJ, USA); 3-hydroxytyramine hydrochloride
114 (dopamine; Tokyo Chemical Industry, Tokyo, Japan); haloperidol (Pfizer, New York, NY,
115 USA); and 5-hydroxytryptamine hydrochloride (5-HT; Nacalai Tesque, Kyoto, Japan).
116
117 **2.2. Animals**
118 All animal care and experimental protocols were approved by the Committee on
119 Animal Experimentation, Graduate School of Veterinary Medicine, Hokkaido University
120 (No. 19-0009), which was awarded the Accreditation Status by the Association for
121 Assessment and Accreditation of Laboratory Animal Care International. Animal studies
122 were performed in compliance with ARRIVE guidelines (Percie du Sert et al., 2020). Wistar
123 rats were obtained from CLEA Japan (Tokyo, Japan) and were bred to obtain pups. The
124 rats were fed *ad libitum* and kept on a 12 h light-dark cycle at 22 ± 4 °C. Male and female
125 pups (3-5 days old) were used for primary astrocyte cultures, and male pups (14-16 days
126 old) were used for acute brain slice experiments.
127

128 **2.3. Primary cultured astrocytes**

129 Primary cultured astrocytes were obtained as previously described (Morimoto et
130 al., 2020). In brief, the cerebral cortex, hippocampus, and spinal cords were isolated from
131 rat pups (3-5 days old), minced, and incubated with papain (10 U/ml) and DNase (0.1
132 mg/ml). Dissociated cells were suspended in Dulbecco's modified Eagle's medium/Ham's
133 F-12 (#048-29785, FUJIFILM Wako Pure Chemical) containing 10% fetal bovine serum,
134 100 U/ml penicillin, and 0.1 mg/ml streptomycin. The cell suspension was seeded onto a
135 poly-l-lysine-coated T75 flask. After 7-8 days, the flask was shaken at 250 rpm at 37 °C
136 for at least 12 h to remove all cells except astrocytes. Adherent cells were detached with
137 trypsin and re-seeded onto poly-l-lysine-coated 12-well plates or coverslips at a density of
138 $8.0 \times 10^3$ cells/cm$^2$. After 3 days, the cell culture reached confluence, and the medium was
139 changed to serum-free medium. Cell cultures were first treated with antagonists
140 immediately after the medium exchange and were then treated with dopamine or other
141 agonists 1 h after the medium exchange. After a certain amount of time (detailed in the
142 figure legends and results section), the cell culture was used for experiments. The
143 concentrations of dopamine used in this study were determined based on the previous
144 reports (Cragg and Rice, 2004; Koppel et al., 2018). Agonists and antagonists were used at
145 concentrations specific to the target receptors, based on the database ("IUPHAR / BPS
146 Guide to PHARMACOLOGY" https://www.guidetopharmac). The purity of astrocyte

147 cultures was evaluated by immunostaining for the astrocytic marker GFAP. At least 300
148 cells in 12 randomly selected images from three cultures (cerebral cortex, hippocampus,
149 and spinal cord) were evaluated, and all cells we evaluated were positive for GFAP (Fig.
150 S1 and Kitano et al., 2021).
151
152 **2.4. RNA extraction and real-time PCR analyses**
153 Total RNA was extracted from cultured astrocytes using RNAiso Plus (Takara Bio,
154 Tokyo, Japan). To remove genomic DNA and synthesize cDNA, the RNA sample was then
155 incubated with qPCR RT Master Mix with gDNA Remover (TOYOBO, Osaka, Japan).
156 Real-time PCR was performed using Thunderbird SYBR qPCR Mix (TOYOBO), each
157 primer, and the cDNA reaction solution. The primer sequences are provided in Table S1.
158 Thermal cycles were performed using the Eco Real-Time PCR System (Illumina, San Diego,
159 CA, USA). Cycling conditions were 95 °C for 1 min (for initial denaturation), followed by
160 40 cycles of denaturation (95 °C, 15 s), annealing, and extension (temperature: Table S1,
161 45 s). RNAs without reverse transcription were used as a negative control to examine DNA
162 contamination and were not amplified by real-time PCR. Melt curve analysis confirmed
163 that the obtained amplicon was only the one expected in each reaction. The expression
164 levels of IL-6 relative to GAPDH were calculated using the ΔΔCq method and were
165 normalized to the control, which was arbitrarily set to a value of "1.0".

166

167 **2.5. Non-quantitative PCR**

168 Non-quantitative PCR was performed using KOD FX Neo (TOYOBO), each primer,
169 and the cDNA reaction solution obtained by the above method. The primer sequences and
170 product sizes are provided in Table S2. Thermal cycles were performed using a PC320
171 system (ASTEC, Fukuoka, Japan). Cycling conditions were 94 °C for 1 min (for initial
172 denaturation), followed by 40 cycles of denaturation (98 °C, 10 s), annealing (temperature:
173 Table S2, 10 s), and extension (68 °C, 30 s). RNAs without reverse transcription were used
174 as a negative control to examine DNA contamination. PCR products and a 100 bp DNA
175 ladder (Takara Bio) were separated on a 3% agarose gel and visualized with ethidium
176 bromide under UV illumination (Mupid-Scope WD, Mupid, Tokyo, Japan).

177

178 **2.6. Western blotting**

179 Astrocytes were lysed in RIPA buffer containing a protease inhibitor cocktail
180 (Nacalai Tesque). The samples were separated by 10% SDS-PAGE and transferred to
181 polyvinylidene difluoride membranes (Millipore, CA, USA). The membranes were blocked
182 with 5% skimmed milk and then incubated with primary antibodies at 4 °C for at least 12
183 h. Thereafter, the membranes were incubated for 1 h at room temperature (RT) with a
184 horseradish peroxidase-conjugated secondary antibody (#NA931 or #NA934, 1:3000, GE

185 Healthcare, Little Chalfont, UK). Antibody binding was visualized by ECL Prime (GE
186 Healthcare). Band intensities were measured using Fiji-ImageJ software (National
187 Institutes of Health) and normalized to the control, which was arbitrarily set to a value of
188 "1.0".
189
190 **2.7. Enzyme-linked immunosorbent assay (ELISA)**
191 We measured the IL-6 protein levels using IL-6 ELISA Kit (#437107, Biolegend,
192 San Diego, CA, USA). The medium of cultured astrocyte treated with each drug for 6 h
193 was collected. Thereafter, the medium was centrifuged at 1,000 × g for 10 min to remove
194 dead cells and debris, and the supernatant was used for ELISA. The experiment was
195 performed according to the manufacturer's instructions and the ELISA plates were read
196 with an SH-1000 lab fluorescent microplate reader (Corona Electric Co., Ibaraki, Japan)
197 and analyzed using SF6 software (Corona Electric Co.). Quantification of astrocyte total
198 protein was performed for cells in each cultured well using the DC™ Protein Assay reagent
199 (Bio-Rad, Hercules, CA, USA). The IL-6 level was normalized by calculating the IL-6
200 protein content per astrocyte total protein content (pg/mg).
201
202 **2.8. Phalloidin staining and evaluation of astrocytic morphology *in vitro***
203 Phalloidin staining and evaluation of astrocytic morphology were conducted as

204 previously described (Kitano et al., 2021). Astrocytes cultured on coverslips were fixed
205 with 4% paraformaldehyde for 20 min at RT and then permeabilized with phosphate-
206 buffered saline containing 0.1% Triton X-100 at RT for 5 min. To stain filamentous actin
207 (F-actin), cells were incubated with Phalloidin-iFluor 488 reagent (#ab176753, 1:1000,
208 Abcam, Cambridge, UK) in phosphate-buffered saline containing 1% bovine serum
209 albumin at RT for 1 h. Coverslips were mounted onto glass slides with DAPI-Fluoromount
210 G (SouthernBiotech, Birmingham, AL, USA). Fluorescence images were obtained with a
211 fluorescence microscope (BZ-9000, KEYENCE, Osaka, Japan) using a 20× lens objective.
212 Astrocytes with process formation were defined as cells that had one or more processes
213 longer than the width of their cell bodies. The number of astrocytes with process formation
214 was visually counted using Fiji-ImageJ software. The mean percentage from more than
215 200 cells from three random images was used as one independent measurement.
216

217 **2.9. Preparation of acute hippocampal slices**

218 Male pups (14-16 days old) were anesthetized with isoflurane (Pfizer) inhalation
219 and rapidly decapitated. The brains were then quickly detached and transferred into ice-
220 cold artificial cerebrospinal fluid (ACSF) and constantly oxygenated with 95% $O_2$ and 5%
221 $CO_2$. The composition of ACSF was as follows (mM): 125 NaCl, 2.5 KCl, 2.0 $CaCl_2$, 1.0
222 $MgCl_2$, 26 $NaHCO_3$, 1.25 $NaH_2PO_4$, and 25 glucose (pH 7.3-7.4). The brain was glued to a

223 slicer stage (LinearSlicer Pro7, Dosaka EM, Kyoto, Japan), flooded in oxygenated cold
224 ACSF, and cut into 300 μm-thick coronal slices. The slices were incubated for 30 min at
225 22-24 °C in continuously oxygenated ACSF. Afterward, the slices, in continuously
226 oxygenated ACSF containing L(+)-ascorbic acid (200 μM), were incubated for 30 min and
227 treated with isoproterenol, dopamine, or SKF81297 for 90 min at 34 °C.
228
229 **2.10. Immunohistochemistry**
230 The slices obtained by the above method were fixed with 4% paraformaldehyde
231 for 12 h at 4 °C and then blocked for 6 h with a blocking buffer composed of 10% goat
232 serum, 0.5% Triton X-100, and 0.05% sodium azide in phosphate-buffered saline. The
233 slices were then incubated with an anti-GFAP primary antibody at 4 °C for at least 12 h
234 and incubated with an Alexa Fluor 555-conjugated goat anti-mouse antibody (#A21422,
235 1:500, Thermo Fisher Scientific, MA, USA) for 2 h at RT. The slices were mounted onto
236 glass slides with DAPI-Fluoromount G, and images were observed with a laser scanning
237 confocal microscope (LSM 700, Carl Zeiss, Oberkochen, Germany) using a 40× lens
238 objective. The CA1 areas of the hippocampus (shown in Fig. S2) were used to measure the
239 fluorescence intensity. The images (shown in Fig. 6) were used for the fluorescence
240 intensity measurements and were composed of 15 μm Z-stacks consisting of 16 optical
241 slices of 1 μm thickness by maximum intensity projection. The mean grey intensity in the

242 area excluding the neuronal layer was measured using Fiji-ImageJ. The results were
243 expressed as arbitrary units.
244
245 **2.11. Morphological analysis of astrocytes in hippocampal slices**
246 Morphological features of astrocytes in the CA1 area of the hippocampus were
247 assessed using the confocal Z-stack images obtained by immunohistochemistry. For
248 analysis, we applied Simple Neurite Tracer, a free software plugin distributed by Fiji-
249 ImageJ and available at (https://imagej.net/plugins/snt/), as previously described (Tavares
250 et al., 2017). The morphological parameters assessed by Simple Neurite Tracer were the
251 total branch length, number of branches, and average branch length. Additionally, we
252 performed Sholl analysis, which measures the number of intersections at concentric
253 spheres (at 4 μm intervals) originating from the soma. The mean value of 10 cells in one
254 immunohistochemistry image was used as one independent measurement.
255
256 **2.12. Data and statistical analysis**
257 All the studies were designed to generate groups of equal size, using
258 randomization and blinded analysis. Data are expressed as means ± S.E.M (n = number
259 of independent measurements) of at least five independent experiments (biological
260 replicates). After confirming that the data were normally distributed, the following tests

were performed. Statistical comparisons between the two groups were made using the unpaired Student's t-test. For all multiple comparisons, the Dunnett's test or Tukey's test was used. The Dunnett's test or Tukey's test was performed only if F achieved $p < 0.05$ and there was no significant inhomogeneity of variance by one-way ANOVA. A value of $p < 0.05$ was considered statistically significant. All statistical analysis was performed using the statistical analysis software JMP® 14 (SAS Institute, Inc., Cary, NC, USA).

268 **3. Results**

269

270 **3.1. Dopamine increases IL-6 mRNA levels in astrocytes and changes astrocyte cell**

271 **morphology**

272 We first examined the effects of monoamines (dopamine, serotonin, histamine) on

273 the mRNA levels of cytokines (IL-6, IL-1β, tumor necrosis factor-α) and growth factors

274 (fibroblast growth factor 2, BDNF, nerve growth factor). Serotonin and histamine (10 μM)

275 did not affect the mRNA levels of any of the factors (Fig. 1A-F) and dopamine (10 μM) did

276 not affect the mRNA levels of any of the factors except IL-6 (Fig. 1A-E). Conversely,

277 treatment with dopamine for 1 h (but not 3 h) increased IL-6 mRNA levels in cerebral

278 cortical astrocytes (Fig. 1F). In addition, dopamine for 1 h also increased IL-6 mRNA levels

279 in hippocampal and spinal cord astrocytes (Fig. 1G).

280 Next, we investigated the effects of monoamines on astrocytic process formation.

281 We previously demonstrated that noradrenaline induces processes in cultured astrocytes,

282 which peaked after 3 h of treatment (Kitano et al., 2021). According to these results,

283 astrocytic process formation *in vitro* was evaluated after 3 h of treatment in this study.

284 Dopamine (10 μM for 3 h) induced process formation in hippocampal astrocytes, whereas

285 serotonin and histamine had no effect on cell morphology (Fig. 2A and B). Furthermore,

286 dopamine induced process formation in cerebral cortical and spinal cord astrocytes (Fig.

287 2C and D). Next, we confirmed the mRNA expression of dopamine receptor subtypes (D1-
288 D5) and β-adrenoceptor subtypes ($β_1$-$β_3$) in astrocytes. Bands of all receptor subtypes were
289 detected in cerebral cortical, hippocampal, and spinal cord astrocytes (Fig. 2E). Several
290 molecularly distinct types of astrocytes with a region-specific distribution have been
291 reported (Zeisel et al., 2018). However, the effects of dopamine on IL-6 mRNA levels and
292 process formation were almost the same across different brain regions. Furthermore, there
293 were no differences in gene expression of each receptor depending on the site of derivation.
294 Although the cerebral cortical astrocytes are mainly used to examine the function or
295 production mechanism of IL-6, GFAP-staining in the cerebral cortex of slice experiments
296 was very weak (Fig. S3). Therefore, in the following experiments, cerebral cortical
297 astrocytes were used to evaluate IL-6 expression and protein phosphorylation, and
298 hippocampal astrocytes were used to evaluate astrocytic morphology.
299

300 **3.2. Dopamine at low and high concentrations increases IL-6 mRNA levels and release via**
301 **D1-like receptors and β-adrenoceptors, respectively**

302 We investigated which receptors are involved in the dopamine-induced increase
303 in IL-6 mRNA levels. High dopamine concentrations also act on β-adrenoceptors (Koppel
304 et al., 2018). Therefore, we investigated the concentration-response relationships between
305 dopamine and IL-6 mRNA levels in the presence of the β-adrenoceptor antagonist

306 propranolol. Treatment of cerebral cortical astrocytes with dopamine (1 nM to 100 μM)
307 increased IL-6 mRNA levels in a concentration-dependent manner (Fig. 3A). Propranolol
308 (10 μM) inhibited the increase in IL-6 mRNA levels induced by a high concentration of
309 dopamine (100 μM) but not that induced by a low concentration of dopamine (1 μM).
310 Dopamine (1 μM) significantly increased the IL-6 mRNA levels (Fig. 3B). In the following
311 experiments, 1 μM dopamine was used as the lowest concentration that significantly
312 increased IL-6 mRNA levels, while 100 μM dopamine was used as the high concentration
313 of dopamine that was significantly inhibited by propranolol. The D1-like receptor
314 antagonist SCH23390 (10 μM) but not the D2-like receptor antagonist haloperidol (10 μM)
315 inhibited the increase in IL-6 mRNA levels at 1 μM dopamine (Fig. 3B). The increase in
316 IL-6 mRNA levels at 100 μM dopamine was not inhibited by SCH23390, was enhanced by
317 haloperidol (Fig. 3C), and was partially inhibited by the $\beta_1$-adrenoceptor antagonist
318 atenolol (10 μM), the $\beta_2$-adrenoceptor antagonist ICI118551 (1 μM), and the $\beta_3$-
319 adrenoceptor antagonist SR59230A (1 μM) (Fig. 3D). In the presence of a mixture of
320 atenolol, ICI1118551, and SR59230A, dopamine (100 μM) failed to increase IL-6 mRNA
321 levels. Activation of D1-like receptors stimulates adenylate cyclase and phospholipase C
322 (Lee et al., 2004). IL-6 mRNA levels were increased by the D1-like receptor full agonist
323 SKF81297 (10 μM) and the D1-like receptor adenylyl cyclase agonist SKF83822 (10 μM),
324 but not by the D1-like receptor phospholipase C agonist SKF83959 (10 μM) or the D2-like

325 receptor agonist bromocriptine (10 μM) (Fig. 3E). The β-agonist isoproterenol (1 μM) and
326 the adenylate cyclase activator forskolin (10 μM) increased IL-6 mRNA levels (Fig. 3F).
327 None of the antagonists alone exerted any effect on IL-6 mRNA levels (Fig. S4A and B).
328 Next, the protein levels of IL-6 released into the culture medium were measured by ELISA.
329 Similar to the effects of dopamine on the IL-6 mRNA levels, the low (1 μM) and high (100
330 μM) concentrations of dopamine increased the release of IL-6, which were inhibited by
331 SCH23390 and propranolol, respectively (Fig. 3G and H). In addition, the release of IL-6
332 by dopamine (100 μM) was enhanced by haloperidol.
333
334 **3.3. Dopamine promotes CREB phosphorylation**
335 We have previously reported that noradrenaline increases IL-6 mRNA levels via
336 the CREB and ERK phosphorylation (Morimoto et al., 2021). Here, we investigated
337 whether low and high dopamine concentrations regulate the phosphorylation of proteins
338 involved in transcription in cerebral cortical astrocytes. Since the increase in IL-6 mRNA
339 was detected at 1 hour, we assumed that the phosphorylation of these factors occurred
340 before that, and thus we measured the phosphorylation at 30 minutes. Dopamine (1 μM
341 for 30 min) promoted CREB phosphorylation, which was inhibited by SCH23390 but not
342 by haloperidol or propranolol (Fig. 4A). Dopamine (100 μM for 30 min) also promoted
343 CREB phosphorylation, which was inhibited by SCH23390, haloperidol, and propranolol

344 (Fig. 4B). SKF81297, isoproterenol, and forskolin promoted CREB phosphorylation (Fig.
345 4C). Dopamine exerted no effect on STAT3 or mitogen-activated protein kinases (MAPKs);
346 namely ERK, JNK, and p38 phosphorylation at low or high concentrations (Fig. 4D-G).
347

348 **3.4. High dopamine concentrations regulate process formation via D2-like receptors and**
349 **β- and $\alpha_2$-adrenoceptors**

350 Next, we investigated which receptors are involved in dopamine-induced process
351 formation in hippocampal astrocytes. We used hippocampal astrocytes, but not cerebral
352 cortical astrocytes, to investigate the evaluation of morphological changes, because GFAP-
353 fluorescence was hardly detectable in the cerebral cortical slice. The expression level of
354 GFAP in the cerebral cortical astrocytes is much lower than that in the hippocampus
355 astrocytes (Zhang et al., 2019). As shown in Figure 2, there were no regional differences
356 in the effect of dopamine and the receptor expression between the cerebral cortical and
357 hippocampal astrocytes. Dopamine at 1 μM had no effect on process formation (Fig. 5A
358 and B), whereas dopamine at 100 μM induced process formation (Fig. 5C and D). This
359 effect was inhibited by propranolol but not by SCH23390. Haloperidol and the $\alpha_2$-
360 adrenoceptor antagonist atipamezole (10 μM) enhanced dopamine-induced process
361 formation. The effect of dopamine (100 μM) was partially inhibited by atenolol, ICI118551,
362 and SR59230A (Fig. 5E and F). SKF81297, SKF83822, isoproterenol, and forskolin, but

363 not SKF83959 and bromocriptine, induced process formation (Fig. 5G-J). None of the
364 antagonists alone exerted any effect on process formation (Fig. S5).
365
366 **3.5. D1-like receptor and β-adrenoceptor agonists increase GFAP expression and change**
367 **astrocytic morphology in acute hippocampal slices**
368 We investigated whether activating dopamine receptors affects astrocytic
369 morphology in acute hippocampal slices in addition to cultured astrocytes. Dopamine,
370 SKF81297, and isoproterenol (10 μM for 90 min) increased the mean intensity of GFAP
371 expression (Fig. 6A and B). Astrocytes treated with these drugs displayed increases in the
372 total branch length and the number of branches, but not the average branch length (Fig.
373 6C-E). In addition, the Sholl analysis showed increases in intersections and shifts in the
374 curve to the right, which indicates an enhanced complexity of astrocytic processes (Fig.
375 6F).
376

377 **4. Discussion**

378 In this study, we found that dopamine regulates IL-6 expression and process
379 formation in astrocytes. High dopamine concentrations regulated these effects via α- and
380 β-adrenoceptors in addition to dopamine receptors. Furthermore, we observed
381 bidirectional regulation, i.e., the effects of D1-like receptors and β-adrenoceptors were
382 negatively regulated by D2-like receptors and $\alpha_2$-adrenoceptors.

383 Our previous study demonstrated that the noradrenaline-induced increase in IL-
384 6 mRNA levels reached a peak 1 h after treatment (Morimoto et al., 2021). The effect of
385 dopamine in this study was similar. After 1 h, dopamine transiently increased IL-6 mRNA
386 levels in a concentration-dependent manner. Dopamine-induced increases in IL-6 mRNA
387 levels were accompanied by CREB phosphorylation, which was abolished by the D1-like
388 receptor antagonist (when 1 μM dopamine was used) and the β-antagonist (when 100 μM
389 dopamine was used). Furthermore, the D1-like receptor adenylyl cyclase agonist
390 SKF83822, but not the D1-like receptor phospholipase C agonist SKF83959, increased IL-
391 6 mRNA levels. The β-agonist and the adenylyl cyclase activator also increased IL-6 mRNA
392 levels. These results suggest that low dopamine concentrations act via the D1-like
393 receptor/cAMP/CREB pathway, whereas high dopamine concentrations act via the β-
394 adrenoceptor/cAMP/CREB pathway to activate IL-6 transcription. These results agree
395 with our previous study showing that the activation of β-adrenoceptors promotes IL-6

396 transcription via the CREB pathway (Morimoto et al., 2021).

397 ERK and STAT3 phosphorylation promote IL-6 transcription in cerebral cortical
398 astrocytes (Du et al., 2020; Sun et al., 2017). However, dopamine did not affect MAPKs or
399 STAT3 phosphorylation. Therefore, these factors are unlikely to be involved in dopamine-
400 induced increases in IL-6 mRNA levels. Although the D1-like receptor antagonist partially
401 decreased dopamine (100 μM)-induced CREB phosphorylation, it had no effect on IL-6
402 mRNA levels. These effects are likely due to the potent effect of dopamine via β-
403 adrenoceptors, and CREB phosphorylation above a certain level may not contribute to the
404 increase in IL-6 mRNA levels. Furthermore, the D2-like receptor antagonists further
405 enhanced dopamine (100 μM)-induced IL-6 mRNA increases, suggesting that D2-like
406 receptors exert a suppressive effect on IL-6 transcription in the presence of high dopamine
407 concentrations. Contrary to this result, D2-like receptor antagonists suppressed dopamine
408 (100 μM)-induced CREB phosphorylation. Therefore, other pathways are likely to be
409 involved in this suppressive effect.

410 Dopamine (100 μM) induced astrocytic process formation, which was abolished by
411 a β-antagonist. Conversely, an $\alpha_2$-antagonist enhanced dopamine-induced process
412 formation. As we previously reported, the activation of β-adrenoceptors induces astrocytic
413 process formation via cAMP signaling, whereas the activation of $\alpha_2$-adrenoceptors inhibits
414 both cAMP-dependent and -independent astrocytic process formation (Kitano et al., 2021).

415 In this study, we showed that the effects of dopamine on process formation were also
416 regulated by β- and $\alpha_2$-adrenoceptors. In addition, the D1-like receptor agonist induced
417 process formation, and the D2-like receptor antagonist enhanced dopamine-induced
418 process formation. Therefore, D1- and D2-like receptors are likely to play a role in the
419 bidirectional regulation of process formation by dopamine.

420 It has been reported that the dephosphorylation of myosin light chains by down-
421 regulation of the Rho pathway is involved in the mechanism of intracellular cAMP-
422 induced process formation (Rodnight and Gottfried, 2013). Therefore, the dopamine-
423 induced increase in IL-6 mRNA and process formation is likely to occur by different
424 intracellular pathways, and these could be a reason to explain the difference between the
425 duration of dopamine effect on IL-6 mRNA (1 h, but not 3 h) and that on process formation
426 (3 h).

427 The effect of β-adrenoceptor agonists on astrocytic processes have been observed
428 in brain slices *in situ* (Sherpa et al., 2016), *in vivo* (Hodges-Savola et al., 1996; Sutin and
429 Griffith, 1993), and in cultured astrocytes *in vitro* (Kitano et al., 2021). The adenylate
430 cyclase activator forskolin increases the overall thickness of the primary processes in the
431 hippocampal slice (Ujita et al., 2017). In this study, we found that dopamine or the D1-like
432 receptor agonist changed astrocytic morphology and upregulated GFAP expression in
433 acute hippocampal slices, suggesting that dopamine receptors are involved in modulating

434 astrocytic morphology *in vivo*. IL-6 upregulation may contribute to dopamine-induced
435 morphological changes in astrocytes. However, it has been reported that the morphology
436 of astrocytes in GFAP-IL6 transgenic mice does not differ from that in normal mice, even
437 though GFAP-IL6 transgenic mice exhibit high IL-6 expression in astrocytes (Penkowa et
438 al., 2003). Therefore, IL-6 is not likely to exert effects on astrocytic morphology.

439 Dopamine is a direct precursor in the synthesis of noradrenaline, and thus
440 dopamine and noradrenaline are structurally similar. Whereas dopamine and
441 noradrenaline normally only interact with their respective receptors, they can also
442 interact with each other's receptors (Lei, 2014). Dopamine has been shown to activate all
443 adrenoceptor subtypes expressed in Chinese hamster ovary cells (Zhang et al., 2004).
444 Intravenously administered dopamine at low doses activates dopamine receptors in blood
445 vessels, whereas dopamine at higher doses activates mainly adrenoceptors (Frishman and
446 Hotchkiss, 1996). These findings support our results, namely, that dopamine at low
447 concentrations acted on dopamine receptors, whereas dopamine at high concentrations
448 acted mainly on adrenoceptors in astrocytes.

449 Adrenergic and dopamine receptors can form homodimers and heterodimers
450 (Franco et al., 2000). The dimerization may lead to different properties from the monomers.
451 For example, D2-like and $\beta_2$ adrenergic receptors form homodimers, which transduce
452 enhanced signals compared to monomers (Hebert et al., 1996; Wouters et al., 2019).

453 Furthermore, D2-like receptors form heterodimers with $\beta_2$ adrenergic receptors and
454 enhance adenylate cyclase activity when stimulated by dopamine (Rebois et al., 2012;
455 Watts and Neve, 1997). A dopamine D2 receptor antagonist decreases the level of D2-like
456 receptors dimer formation (Wouters et al., 2019). In addition, an adenosine A1 receptor
457 antagonist enhances the activation of the dopamine D1 receptor coupled with the A1
458 receptor (Franco et al., 2000). Therefore, the antagonists used in this study may affect not
459 only monometric receptors but also receptor complexes. Further studies are needed to
460 address this issue.

461 In this study, the D2-like, but not the D1-like, receptor antagonist had no effects
462 on the dopamine (1 μM)-induced increase in IL-6 mRNA levels. D2-like receptors have a
463 higher affinity for dopamine (Seeman and Grigoriadis, 1987); however, D2-like receptor
464 expression is lower than D1-like receptor expression in at least 21 brain regions (Richfield
465 et al., 1989). In a simulation with model parameters for dopamine receptors in striatal
466 neurons, the amount of dopamine binding to D2-like receptors was approximately 10 times
467 lower than that to D1-like receptors in the presence of 1 μM dopamine (Hunger et al.,
468 2020). The D2-like receptor antagonist may not have exerted any effect because of the low
469 numbers of D2-like receptors available for dopamine binding.

470 The concentration of dopamine in human cerebrospinal fluid *in vivo* has been
471 reported to be 39.5 ± 19.8 nM (Strittmatter et al., 1997), while the concentration of

472 dopamine in the synaptic gap reaches 10-100 μM (Cragg and Rice, 2004; Koppel et al.,
473 2018), and astrocytic processes contact neurons and synaptic gaps (i.e., the "tripartite
474 synapse"), thus regulating synaptic function (Allen and Eroglu, 2017). Furthermore,
475 dopamine concentrations increase under acute stress and ischemic conditions (Baker et
476 al., 1991; Chang et al., 1993; Pascucci et al., 2007). Rat models of drug abuse and
477 pathological gamblers have higher dopamine levels than healthy groups (Egenrieder et al.,
478 2020; van Holst et al., 2018). It is likely that dopamine concentrations (1-100 μM) used in
479 this study could be reached, at least transiently, *in vivo* under physiological and
480 pathological conditions. Further investigations are needed to evaluate the effects of brief
481 exposure of astrocytes to dopamine.

482 In this study, dopamine (1 μM) increased IL-6 mRNA levels and release via D1-
483 like receptors. IL-6 has been widely reported to play a beneficial role in brain function,
484 e.g., by acting as a neurotrophic factor (Wagner, 1996), suppressing neuronal cell death
485 (Day et al., 2014), and improving learning and memory impairment after traumatic brain
486 injury (Willis et al., 2020). Furthermore, the D1-like agonist SKF83959 is suggested to
487 protect nigral neurons from MPTP neurotoxicity via astrocytic D1-like receptors (Zhang
488 et al., 2009). In addition, the activation of astrocytic D1-like receptors enhances the
489 recovery of brain function after experimental stroke (Kuric et al., 2013). As IL-6 and D1-
490 like receptors appear to contribute to recovery from various diseases, the effects of low

491 dopamine concentrations on IL-6 transcription via astrocytic D1-like receptors may be
492 beneficial. Conversely, high dopamine concentrations are likely to cause CNS
493 inflammation and induce reactive astrocytes. In this study, dopamine (100 μM) induced
494 morphological changes in astrocytic cultures and acute brain slices, and increased GFAP
495 expression in astrocytes in acute brain slices. These characteristics are consistent with
496 those of reactive astrocytes in CNS inflammation, including elongated and complex
497 processes and increased GFAP expression (Pekny and Pekna, 2014). Dopamine and IL-6
498 levels increase under ischemic conditions (Baker et al., 1991; Chang et al., 1993; Clark et
499 al., 1999). In addition, astrocytes upregulate ß-adrenoceptors and downregulate $\alpha_2$-
500 adrenoceptors in neurodegenerative diseases (Mantyh et al., 1995; Shao and Sutin, 1992).
501 Taken together, adrenoceptors in astrocytes may be involved in the pathogenesis of
502 neuroinflammatory diseases associated with extremely elevated dopamine levels. Thus,
503 the inhibitory role of dopamine via D2-like receptors and $\alpha_2$-adrenoceptors may improve
504 such pathological conditions.
505

## CRediT authorship contribution statement

**Kohei Morimoto**: Conceptualization, Formal analysis, Investigation, Writing - original draft, Visualization, Funding acquisition. **Mai Ouchi**: Formal analysis, Investigation, Writing - original draft. **Taisuke Kitano**: Conceptualization, Writing - review & editing. **Ryota Eguchi**: Conceptualization, Writing - review & editing, Funding acquisition. **Ken-ichi Otsuguro**: Conceptualization, Writing - review & editing, Supervision, Funding acquisition.

## Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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757

758

759 **Figure legends**

760

761 **Figure 1. The effects of monoamines on mRNA levels of multiple factors and cell**

762 **morphology in cultured astrocytes.**

763 (A-F) The mRNA levels of tumor necrosis factor-α (A), IL-1β (B), nerve growth factor (C),

764 brain-derived neurotrophic factor (D), fibroblast growth factor 2 (E), and IL-6 (F) in

765 cerebral cortical astrocytes treated with serotonin (5-HT, 10 μM), histamine (HA, 10 μM),

766 and dopamine (DA, 10 μM) for 1 and 3 h. The mRNA levels of each factor were normalized

767 to the control level, which was arbitrarily set to a value of "1.0". $^{**}p < 0.01$ vs. control

768 (Dunnett's test), n = 6. (G) IL-6 mRNA levels in hippocampal and spinal cord astrocytes

769 treated with DA (10 μM) for 1 h. $^{**}p < 0.01$ (unpaired Student's t-test), n = 6. All data are

770 presented as means ± S.E.M.

771

772 **Figure 2. The effects of dopamine on IL-6 mRNA levels, process formation, and receptor**

773 **expression in astrocytes from different brain regions.**

774 (A, B) Representative images of F-actin (green) and DAPI (blue) in hippocampal astrocytes

775 treated with serotonin (5-HT, 10 μM), histamine (HA, 10 μM), and dopamine (DA, 10 μM)

776 for 3 h (A). Scale bars = 100 μm. The percentage of cells with process formation (B). More

777 than 200 cells in three random fields were counted. $^{**}p < 0.01$ vs. control (Dunnett's test),

778 n = 6. (C, D) Representative images of F-actin (green) and DAPI (blue) in cerebral cortical

779 and spinal cord astrocytes treated with DA (10 μM) for 3 h (C). Scale bars = 100 μm. The

780 percentage of cells with process formation (D). More than 200 cells in three random fields

781 were counted. $^{**}p < 0.01$ (unpaired Student's t-test), n = 6. All data are presented as means

782 ± S.E.M. (E) Bands for all dopamine receptor and adrenoceptor subtypes were detected in

783 cerebral cortical (upper), hippocampal (middle), and spinal cord (lower) astrocytes. RT (+)

784 and (–) indicates samples reverse-transcribed (+) or not (–), respectively.

785

786 **Figure 3. The effects of dopamine receptor and adrenoceptor agonists or antagonists on**

787 **IL-6 mRNA levels and release.**

788 (A) IL-6 mRNA levels in cerebral cortical astrocytes treated with dopamine (DA, 1 nM to

789 100 μM) in the presence or absence of the β-antagonist propranolol (PROP, 10 μM) for 1 h.

790 $^{**}p < 0.01$ (unpaired Student's t-test), n = 6. (B-D) IL-6 mRNA levels in astrocytes treated

791 with DA (B: 1 μM, C and D: 100 μM) in the presence or absence of the D1-like receptor

792 antagonist SCH23390 (SCH, 10 μM), D2-like receptor antagonist haloperidol (HAL, 10

793 μM), PROP (10 μM), $\beta_1$-adrenoceptor antagonist atenolol (ATE, 10 μM), $\beta_2$-adrenoceptor

794 antagonist ICI118551 (ICI, 1 μM), and $\beta_3$-adrenoceptor antagonist SR59230A (SR, 1 μM)

795 for 1 h. n.s.: not significant, $^{*}p < 0.05$, $^{**}p < 0.01$ (vs. DA alone, B and C, Dunnett's test),

796 (D, Tukey's t-test), n = 6. (E, F) IL-6 mRNA levels in astrocytes treated with the D1-like

797 receptor full agonist SKF81297 (10 μM), D1-like receptor adenylyl cyclase agonist
798 SKF83822 (10 μM), D1-like receptor phospholipase C agonist SKF83959 (10 μM), D2-like
799 receptor agonist bromocriptine (BRO, 10 μM), β-agonist isoproterenol (ISO, 1 μM), or
800 adenylate cyclase activator forskolin (FSK, 10 μM) for 1 h. *$p < 0.05$, **$p < 0.01$ vs. control
801 (Dunnett's test), n = 6. (G, H) IL-6 protein levels of the medium were measured by ELISA.
802 Astrocyte was treated with each drug for 6 h. IL-6 levels were normalized by astrocyte
803 total protein. *$p < 0.05$, **$p < 0.01$ vs. DA alone (Dunnett's test), n = 5. All data are
804 presented as means ± S.E.M.
805
806 **Figure 4. The effects of dopamine receptor and adrenoceptor agonists or antagonists on**
807 **CREB, MAPKs, and STAT3 phosphorylation.**
808 (A-C) The protein expression levels of phosphorylated and total CREB were quantified,
809 and representative blots are shown. Cerebral cortical astrocytes were treated with
810 dopamine (A: 1 μM, B: 100 μM), D1-like receptor full agonist SKF81297 (10 μM), β-agonist
811 isoproterenol (ISO, 1 μM), and adenylate cyclase activator forskolin (FSK, 10 μM) in the
812 presence or absence of the D1-like receptor antagonist SCH23390 (SCH, 10 μM), D2-like
813 receptor antagonist haloperidol (HAL, 10 μM), and β-antagonist propranolol (PROP, 10
814 μM) for 30 min. *$p < 0.05$, **$p < 0.01$ (vs. DA alone, A and B, Dunnett's test), (vs. control,
815 C, Dunnett's test), n = 6. (D-G) The protein expression levels of phosphorylated and total

816 ERK (D), JNK (E), p38 (F), and STAT3 (G) were quantified, and representative blots are

817 shown. Astrocytes were treated with dopamine (1 or 100 μM), n = 6. All data are presented

818 as means ± S.E.M.

819

820 **Figure 5. The effects of dopamine receptor and adrenoceptor agonists or antagonists on**

821 **astrocytic process formation.**

822 (A, C, E, G, I) Representative images of F-actin (green) and DAPI (blue) in hippocampal

823 astrocytes treated with dopamine (DA, A: 1 μM, C and E: 100 μM), D1-like receptor full

824 agonist SKF81297 (10 μM), D1-like receptor adenylyl cyclase agonist SKF83822 (10 μM),

825 D1-like receptor phospholipase C agonist SKF83959 (10 μM), D2-like receptor agonist

826 bromocriptine (BRO, 10 μM), β-agonist isoproterenol (ISO, 1 μM), and adenylate cyclase

827 activator forskolin (FSK, 10 μM) in the presence or absence of the D1-like receptor

828 antagonist SCH23390 (SCH, 10 μM), D2-like receptor antagonist haloperidol (HAL, 10

829 μM), β-adrenoceptor antagonist (PROP, 10 μM), $\alpha_2$-adrenoceptor antagonist atipamezole

830 (ATIP, 10 μM), $\beta_1$-adrenoceptor antagonist atenolol (ATE, 10 μM), $\beta_2$-adrenoceptor

831 antagonist ICI118551 (ICI, 1 μM), and $\beta_3$-adrenoceptor antagonist SR59230A (SR, 1 μM)

832 for 3 h. Scale bars = 100 μm. (B, D, F, H, J) The percentage of cells with process formation.

833 More than 200 cells in three random fields were counted. n.s.: not significant (B, unpaired

834 Student's t-test), $^{**}p < 0.01$ (vs. DA alone, D, Dunnett's test), (F, Tukey's t-test), (vs. control,

835 H and J, Dunnett's test), n = 6. All data are presented as means ± S.E.M.

836

837 **Figure 6. The effects of dopamine receptor and adrenoceptor agonists on GFAP expression**

838 **in acute hippocampal slices.**

839 (A) Representative GFAP-stained images of the CA1 areas in acute hippocampal slices

840 treated with dopamine (DA, 10 μM), D1-like receptor full agonist SKF81297 (SKF, 10 μM),

841 and β-agonist isoproterenol (ISO, 10 μM) for 90 min (Upper left panel: high magnification

842 of the representative astrocyte). Yellow scale bars = 100 μm, green scale bars = 20 μm. (B)

843 The mean grey intensity of GFAP was qualified. The results are expressed as arbitrary

844 units (A.U.). (C-F) The morphology of GFAP-stained astrocytes in the CA1 areas was

845 analyzed using the Fiji-ImageJ Simple Neurite Tracer plugin. The morphological

846 parameters assessed were the total branch length (C), number of branches (D), and

847 average branch length (E); Sholl analysis was also performed (F), which measures the

848 number of intersections at concentric spheres (at 4 μm intervals) originating from the

849 soma (upper right panel). Scale bars = 20 μm. *$p < 0.05$, **$p < 0.01$ vs. control (Dunnett's

850 test), n = 5. All data are presented as means ± S.E.M.

Fig.1

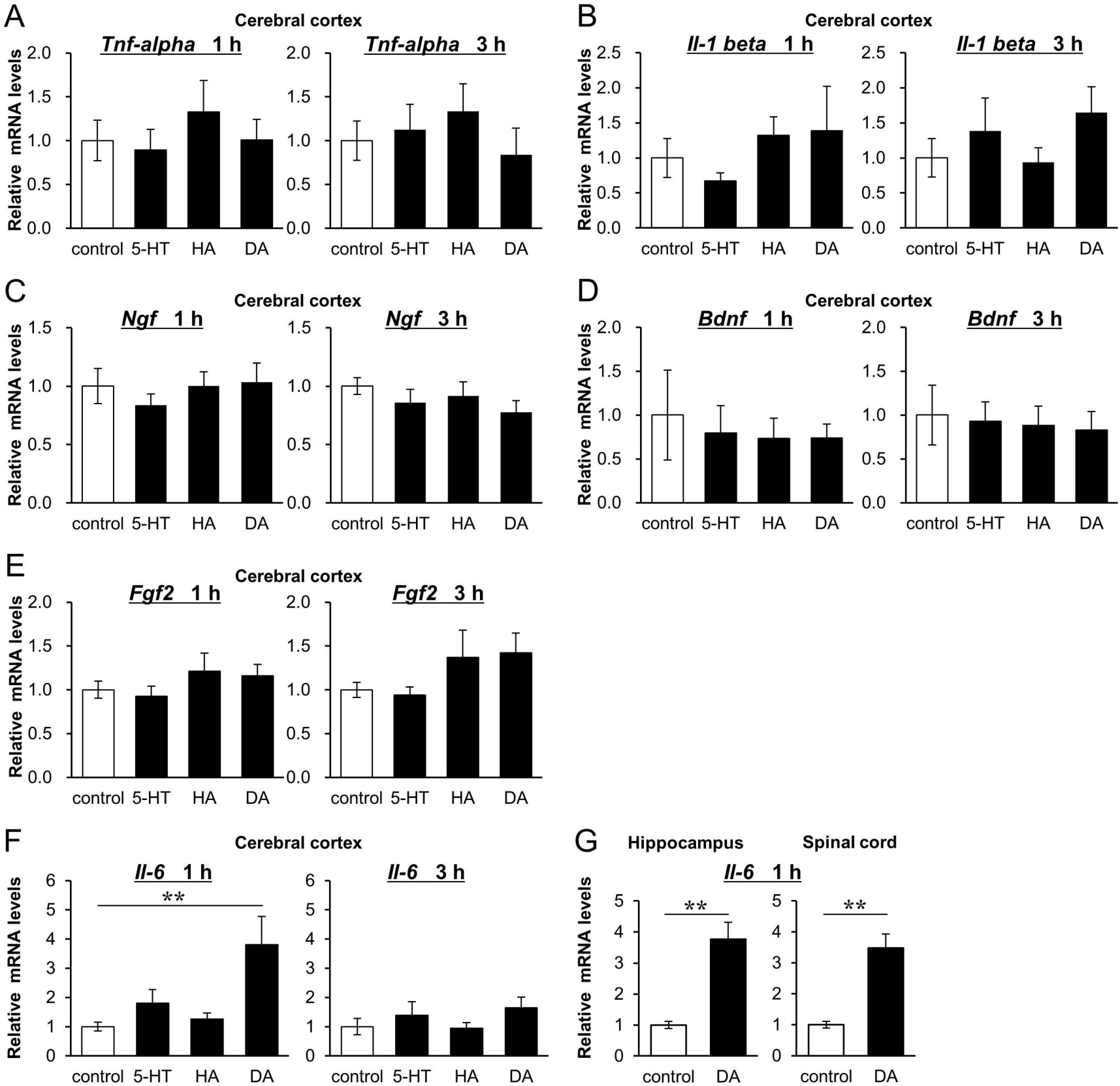

A
Cerebral cortex
Tnf-alpha 1 h
Tnf-alpha 3 h
Relative mRNA levels
control 5-HT HA DA
B
Cerebral cortex
Il-1 beta 1 h
Il-1 beta 3 h
C
Cerebral cortex
Ngf 1 h
Ngf 3 h
D
Cerebral cortex
Bdnf 1 h
Bdnf 3 h
E
Cerebral cortex
Fgf2 1 h
Fgf2 3 h
F
Cerebral cortex
Il-6 1 h
Il-6 3 h
**
G
Hippocampus
Spinal cord
Il-6 1 h
control DA

Fig.2

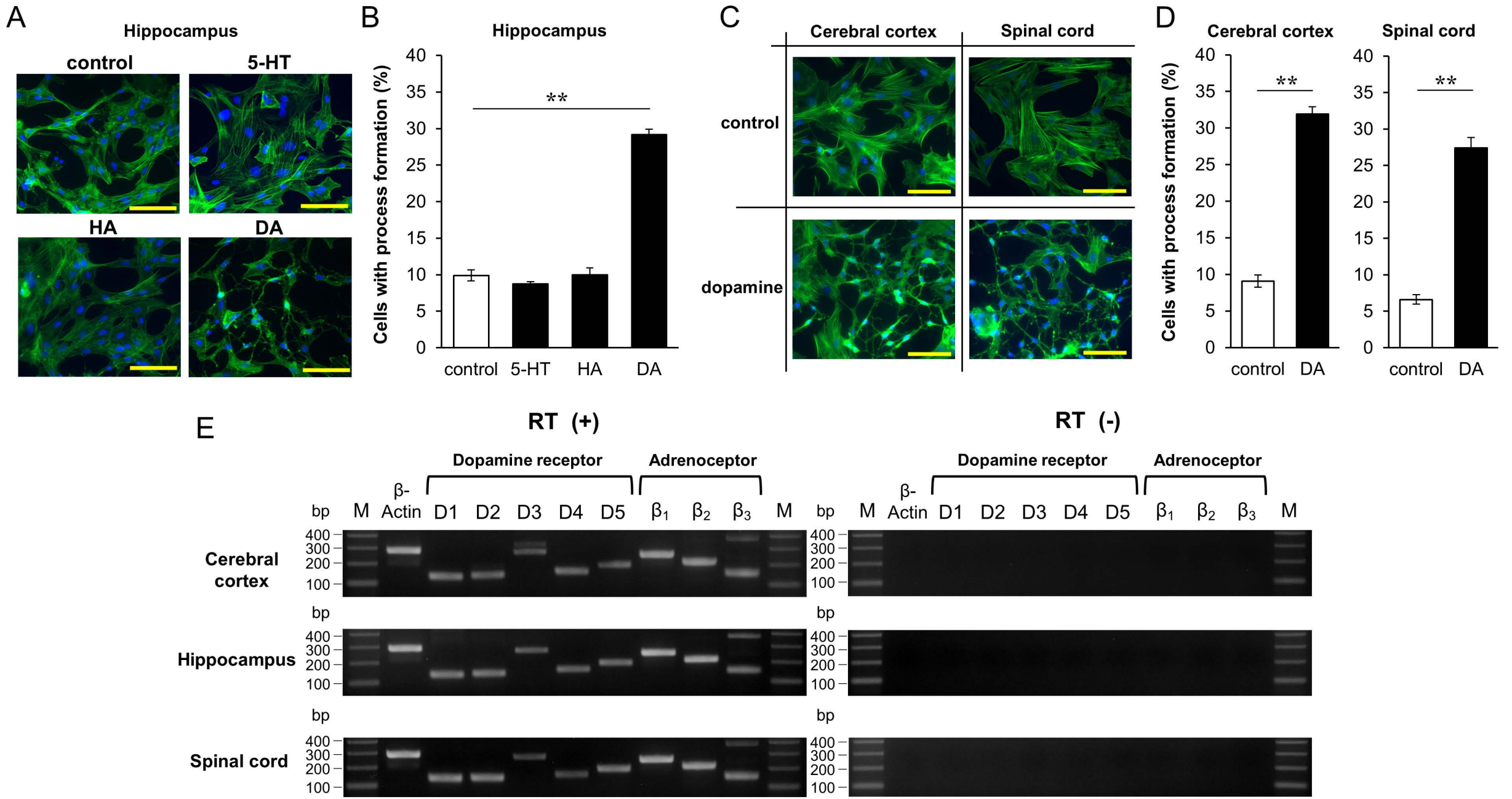

A
Hippocampus
control
5-HT
HA
DA
B
Hippocampus
Cells with process formation (%)
40
35
30
25
20
15
10
5
0
**
control
5-HT
HA
DA
C
Cerebral cortex
Spinal cord
control
dopamine
D
Cerebral cortex
Spinal cord
Cells with process formation (%)
**
control
DA
E
RT (+)
RT (-)
Dopamine receptor
Adrenoceptor
β-Actin
D1
D2
D3
D4
D5
β1
β2
β3
M
bp
400
300
200
100
Cerebral cortex
Hippocampus
Spinal cord

Fig.3

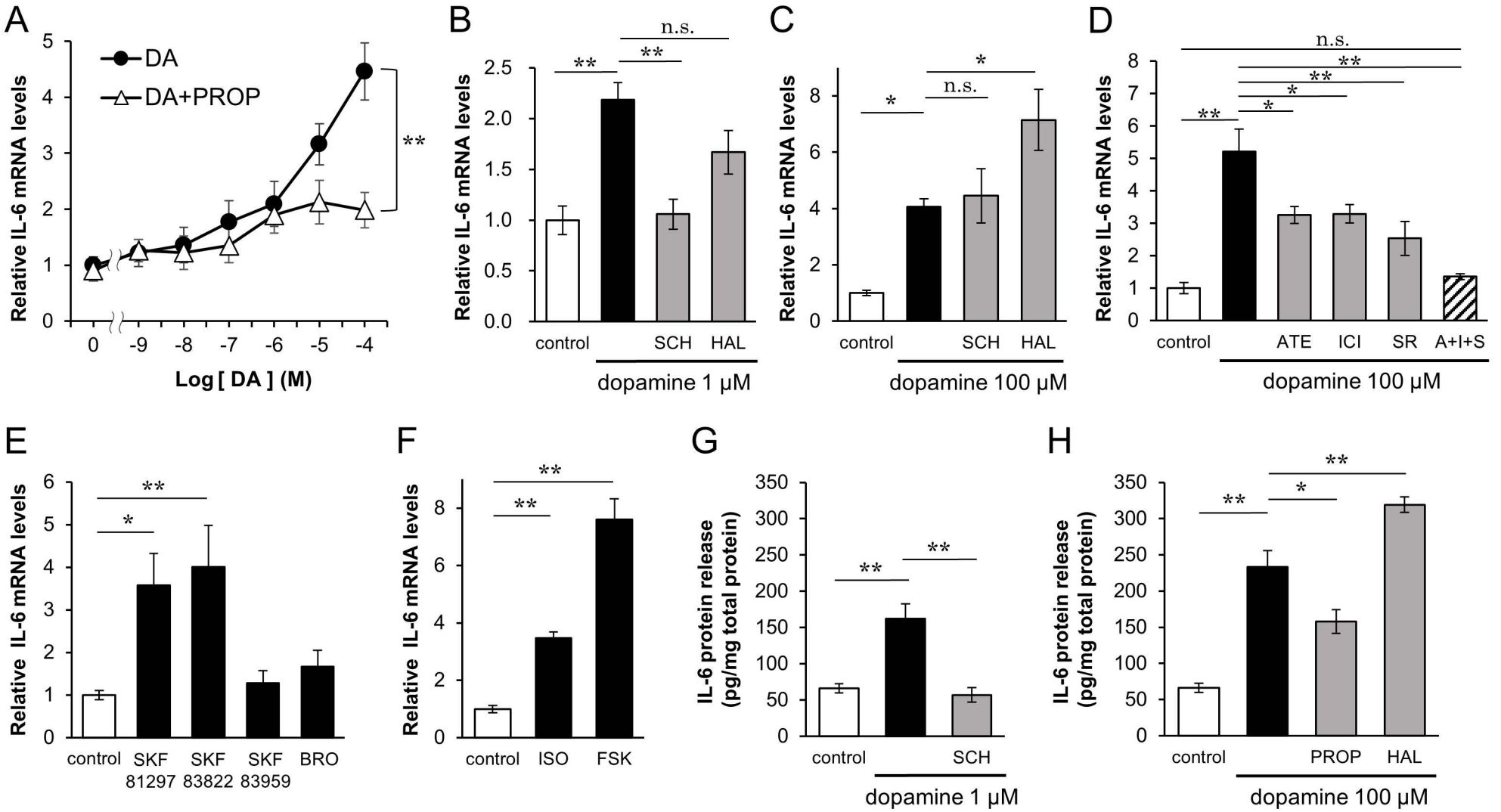

A
DA
DA+PROP
Relative IL-6 mRNA levels
Log [ DA ] (M)
0
-9
-8
-7
-6
-5
-4
**
B
Relative IL-6 mRNA levels
control
SCH
HAL
dopamine 1 μM
n.s.
**
**
C
Relative IL-6 mRNA levels
control
SCH
HAL
dopamine 100 μM
*
n.s.
*
D
Relative IL-6 mRNA levels
control
ATE
ICI
SR
A+I+S
dopamine 100 μM
n.s.
**
**
*
*
**
E
Relative IL-6 mRNA levels
control
SKF 81297
SKF 83822
SKF 83959
BRO
**
*
F
Relative IL-6 mRNA levels
control
ISO
FSK
**
**
G
IL-6 protein release (pg/mg total protein)
control
SCH
dopamine 1 μM
**
**
H
IL-6 protein release (pg/mg total protein)
control
PROP
HAL
dopamine 100 μM
**
*
**

Fig. 4

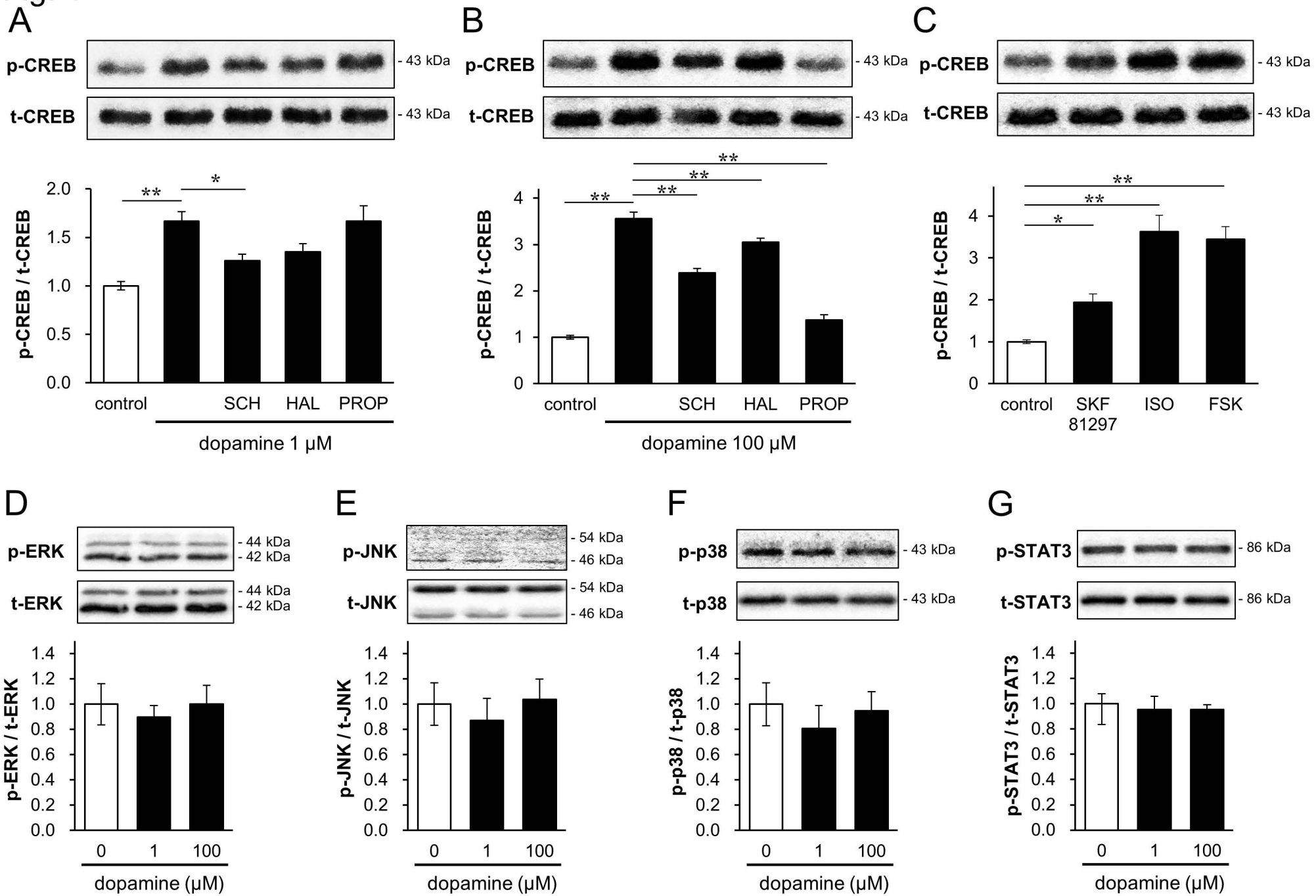

A
p-CREB
t-CREB
- 43 kDa
- 43 kDa
p-CREB / t-CREB
control
SCH
HAL
PROP
dopamine 1 μM
B
p-CREB
t-CREB
- 43 kDa
- 43 kDa
p-CREB / t-CREB
control
SCH
HAL
PROP
dopamine 100 μM
C
p-CREB
t-CREB
- 43 kDa
- 43 kDa
p-CREB / t-CREB
control
SKF 81297
ISO
FSK
D
p-ERK
t-ERK
- 44 kDa
- 42 kDa
p-ERK / t-ERK
dopamine (μM)
E
p-JNK
t-JNK
- 54 kDa
- 46 kDa
p-JNK / t-JNK
dopamine (μM)
F
p-p38
t-p38
- 43 kDa
p-p38 / t-p38
dopamine (μM)
G
p-STAT3
t-STAT3
- 86 kDa
p-STAT3 / t-STAT3
dopamine (μM)

Fig.5

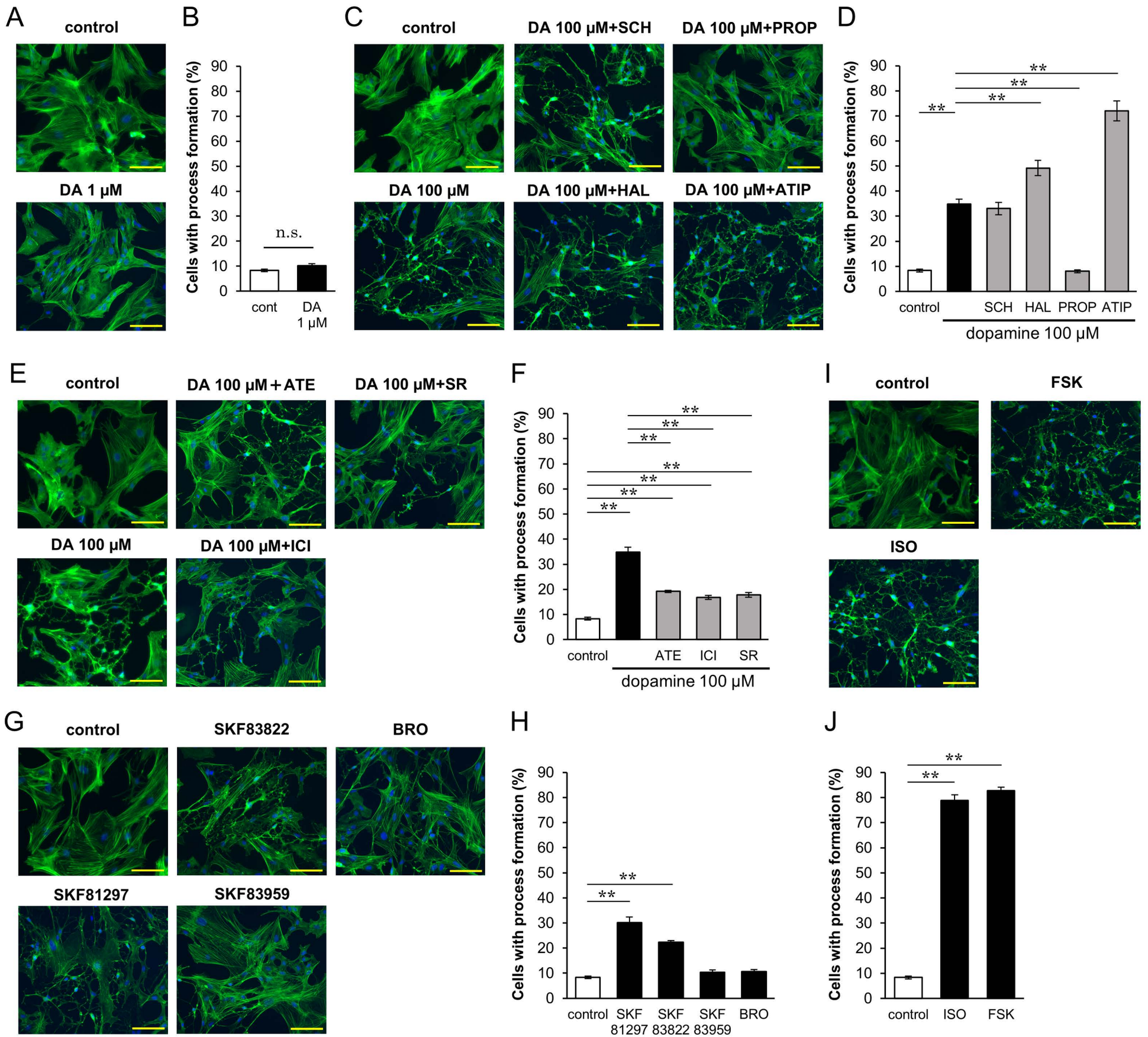
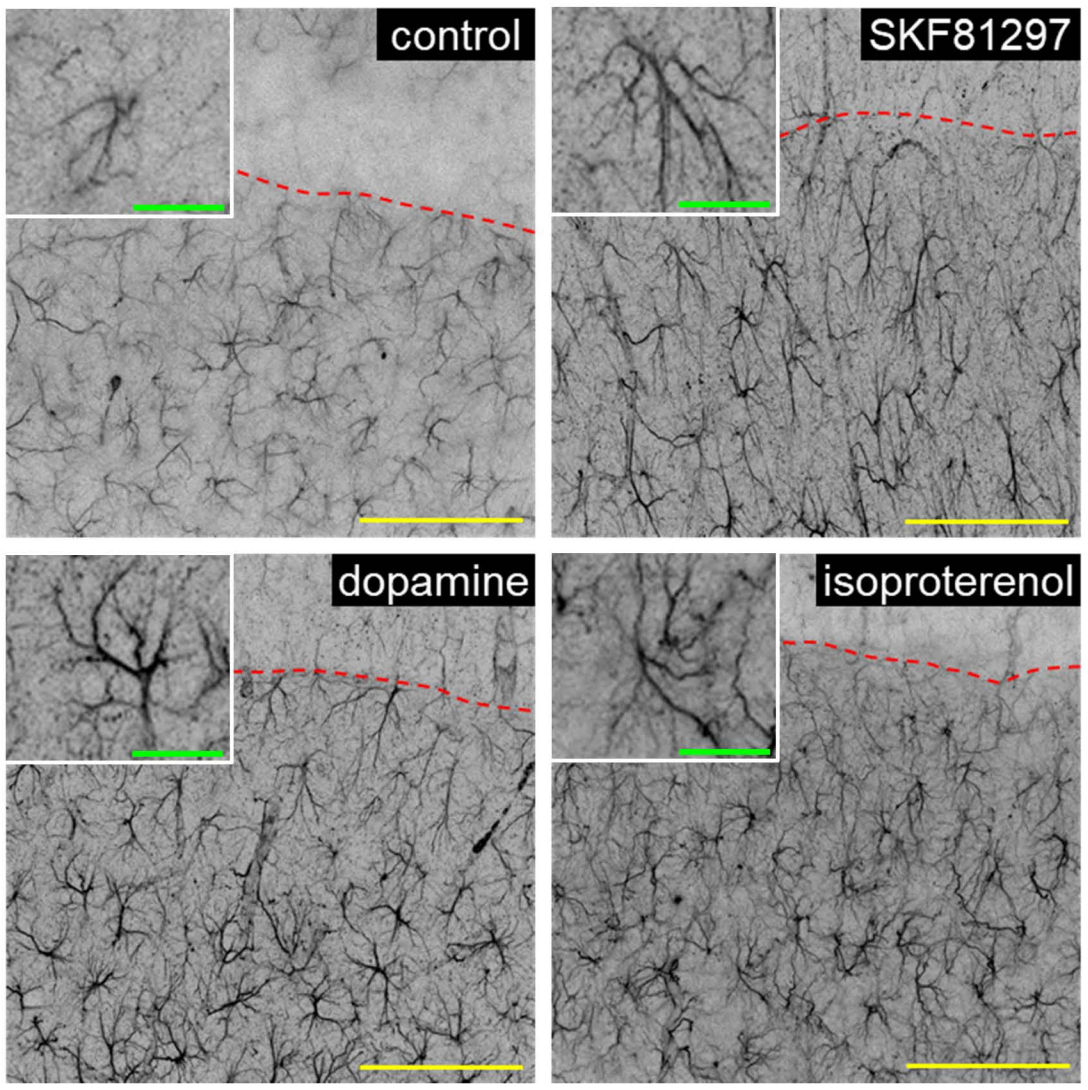
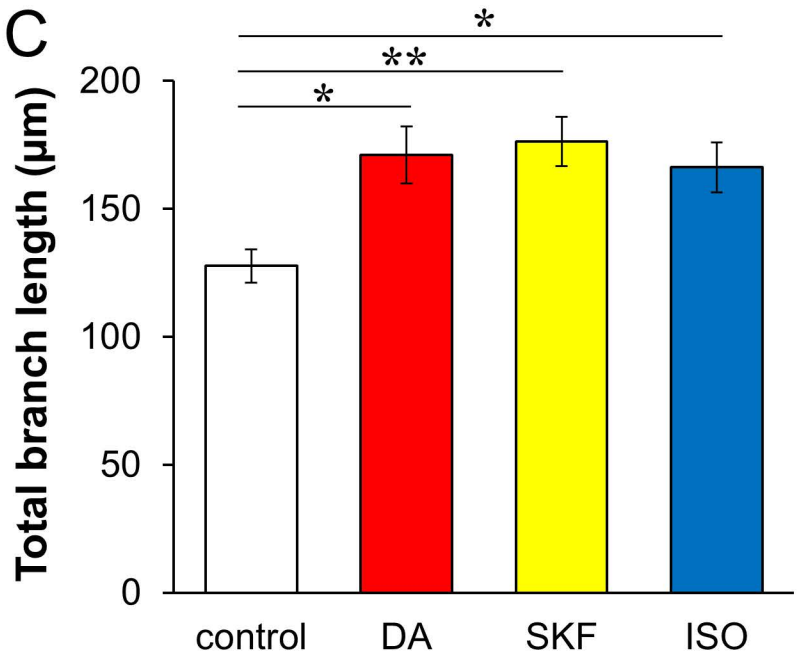

A
control
DA 1 μM
B
Cells with process formation (%)
n.s.
cont
DA 1 μM
C
control
DA 100 μM+SCH
DA 100 μM+PROP
DA 100 μM
DA 100 μM+HAL
DA 100 μM+ATIP
D
Cells with process formation (%)
control
SCH
HAL
PROP
ATIP
dopamine 100 μM
E
control
DA 100 μM＋ATE
DA 100 μM+SR
DA 100 μM
DA 100 μM+ICI
F
Cells with process formation (%)
control
ATE
ICI
SR
dopamine 100 μM
I
control
FSK
ISO
G
control
SKF83822
BRO
SKF81297
SKF83959
H
Cells with process formation (%)
control
SKF 81297
SKF 83822
SKF 83959
BRO
J
Cells with process formation (%)
control
ISO
FSK

Fig.6

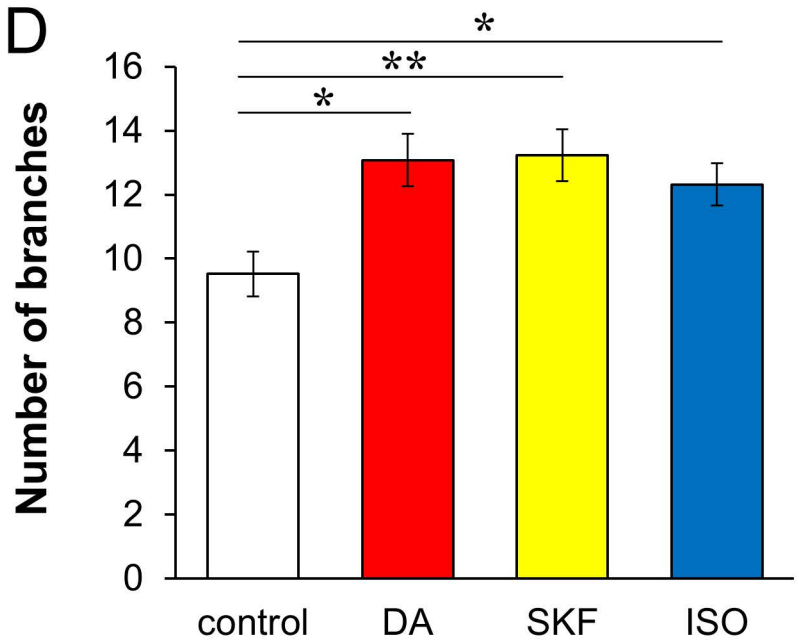
A


control
SKF81297
dopamine
isoproterenol

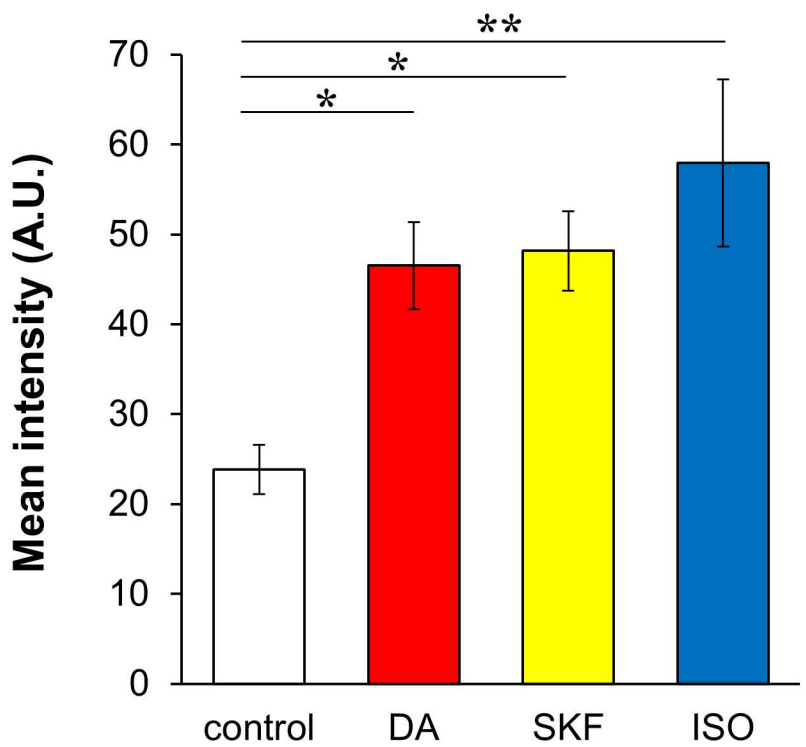
C


Total branch length (μm)
200
150
100
50
0
control
DA
SKF
ISO
*
**
*

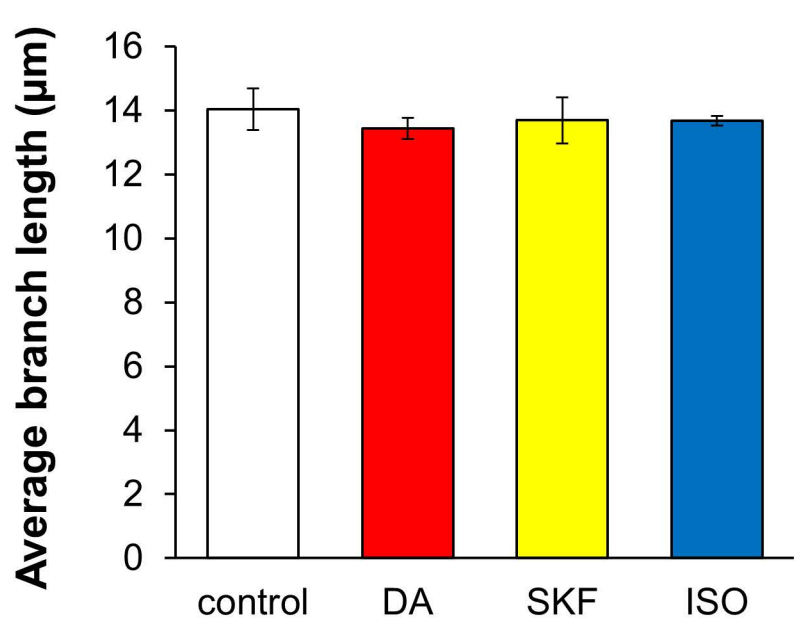
D


Number of branches
16
14
12
10
8
6
4
2
0
control
DA
SKF
ISO
*
**
*

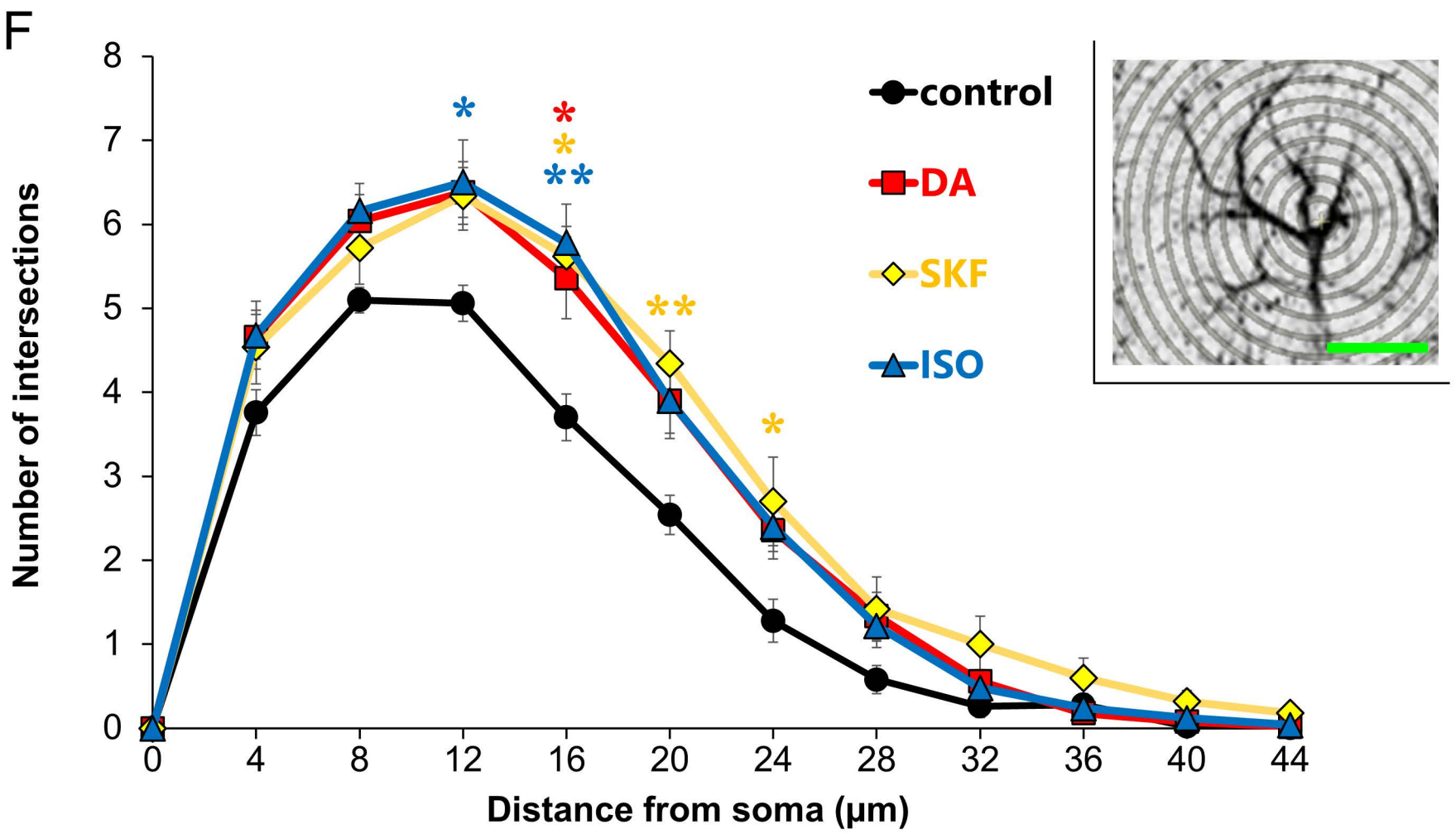
B


Mean intensity (A.U.)
70
60
50
40
30
20
10
0
control
DA
SKF
ISO
**
*
*

E


Average branch length (μm)
16
14
12
10
8
6
4
2
0
control
DA
SKF
ISO

F


Number of intersections
8
7
6
5
4
3
2
1
0
0
4
8
12
16
20
24
28
32
36
40
44
Distance from soma (μm)
control
DA
SKF
ISO
*
*
*
**
**
*