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## **Distinct populations of neurons activated by heroin and cocaine in the striatum as assessed by catFISH**

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1 **Distinct populations of neurons activated by heroin and cocaine in the**  
2 **striatum as assessed by catFISH**

3

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## 26 **Abstract**

27 Despite the still prevailing notion of a shared substrate of action for all addictive drugs, there is  
28 evidence suggesting that opioid and psychostimulant drugs differ substantially in terms of their  
29 neurobiological and behavioural effects. These differences may reflect separate neural circuits  
30 engaged by the two drugs. Here we used the catFISH technique to investigate the degree of  
31 overlap between neurons engaged by heroin versus cocaine in adult male Sprague-Dawley rats.  
32 The catFISH technique is a within-subject procedure that takes advantage of the different  
33 transcriptional time-course of the immediate-early genes *homer 1a* and *arc* to determine to  
34 what extent two stimuli separated by an interval of 25 min engage the same neuronal  
35 population. We found that throughout the striatal complex the neuronal populations activated  
36 by non-contingent intravenous injections of cocaine (800 µg/kg) and heroin (100 and 200  
37 µg/kg), administered at an interval of 25 min from each other, overlapped to a much lesser  
38 extent than in the case of two injections of cocaine (800 µg/kg), also 25 min apart. The greatest  
39 reduction in overlap between populations activated by cocaine and heroin was in the  
40 dorsomedial and dorsolateral striatum (~30% and ~22%, respectively, of the overlap observed  
41 for the sequence cocaine-cocaine). Our results point toward a significant separation between  
42 neuronal populations activated by heroin and cocaine in the striatal complex. We propose that  
43 our findings are a proof of concept that these two drugs are encoded differently in a brain area  
44 believed to be a common neurobiological substrate to drug abuse.

## 45 **Significance statement**

46 Despite significant advances in the substance use disorders field, effective prevention and  
47 treatment strategies are scarce and still under active development. Here we add to growing  
48 evidence indicating major differences in the neurobiological effects of opioid versus  
49 psychostimulant drugs, which is at odds with the still prevailing notion of a shared substrate of  
50 action for all addictive drugs. This suggests that, to be effective, development of prevention and  
51 treatment strategies should not look for a “silver bullet” solution to all drug addictions.  
52 Instead, they should be tailored to the specific drug preference of pathological users.

## 53 Introduction

54 Virtually all current theories of drug abuse posit that the addictive properties of drugs depend  
55 on common neurobiological processes, including hyper-reactivity of motivational systems (e.g.,  
56 Wolf, 2010; Berridge and Robinson, 2016), impaired impulse control (e.g., Jentsch and Taylor,  
57 1999), and aberrant learning (e.g., Everitt and Robbins, 2005). Regardless of the core process  
58 on which each theory focuses, the biological substrate of said processes involves the  
59 mesotelencephalic dopamine system projecting from ventral tegmental area (VTA) and  
60 substantia nigra (SN) to the striatal complex, including caudate and nucleus accumbens (NAcc),  
61 and to the prefrontal cortex (PFCx). Indeed, it is commonly assumed that all substances of  
62 abuse increase dopamine levels in the terminal regions of the dopaminergic system (Di Chiara  
63 and Imperato, 1988; Robinson and Berridge, 1993; Wise, 1996; Nestler, 2001, 2004; Hyman et  
64 al., 2006; Koob and Volkow, 2010; Berridge, 2012; Covey et al., 2014; Keiflin and Janak, 2015;  
65 Volkow and Morales, 2015; Berridge and Robinson, 2016; Keramati et al., 2017; Volkow et al.,  
66 2017) albeit via different mechanisms of action. Psychostimulant drugs, such as cocaine and  
67 amphetamines, produce dopamine overflow by binding the dopamine transporter (for reviews,  
68 see Kuczenski et al., 1982; Johanson and Fischman, 1989). Opioid agonists, such as heroin and  
69 morphine, are thought to increase dopamine concentrations indirectly by binding mu-opioid  
70 receptors located on inhibitory interneurons in the VTA, hence disinhibiting dopaminergic  
71 neurons (Gysling and Wang, 1983; Matthews and German, 1984; Johnson and North, 1992).  
72 Yet, the magnitude of drug-induced dopamine overflow differs enormously from one drug to  
73 another, even within the same pharmacological class. For example, some opioids produce  
74 dramatic increases in dopamine whereas others have very little effect (e.g., Gattas et al., 2014;  
75 Vander Weele et al., 2014). Furthermore, electrophysiological experiments have shown that  
76 neurons in the striatum respond in a very different manner to heroin versus cocaine self-  
77 administration (e.g., Chang et al., 1998; Wei et al., 2018), suggesting that the effects of the two  
78 drugs are encoded differently in this brain area.

79 The aim of the two experiments reported here was to further explore this hypothesis  
80 using the catFISH technique, which is a within-subject technique that takes advantage of the  
81 different transcriptional time-course of the immediate-early genes (IEGs) *homer 1a* and *arc* to

82 detect the activation of partly distinct neuronal populations in response to two temporally  
83 distinct stimuli (see Fig.1, Guzowski et al., 1999; Vazdarjanova et al., 2002; Vazdarjanova and  
84 Guzowski, 2004). To date, a few studies have looked at the effects of cocaine on *arc* (Caffino et  
85 al., 2011) or *homer 1a* expression (Unal et al., 2009), whereas there is no information on the  
86 effects of heroin administration on the expression of these two IEGs. As in the case of the IEG *c-*  
87 *fos*, which is known to be transcribed across the striatum in response to heroin and cocaine  
88 administration (Harlan and Garcia, 1998; Paolone et al., 2007; Celentano et al., 2009), both *arc*  
89 and *homer 1a* are activated by the transcription factor CREB; that is, they are transcribed  
90 following activation of the ERK/MAPK pathway, elevated cAMP activity, or calcium influx to the  
91 cell (Impey et al., 1998; Sato et al., 2001; Kawashima et al., 2014). Considering these shared  
92 mechanisms of expression, we expected that *arc* and *homer 1a* would be suitable markers of  
93 neuronal activity produced by drug administration. We predicted that intravenous (i.v.)  
94 injections of heroin and cocaine will produce a rapid and transient IEG transcription in the  
95 striatum. Indeed, we found that intravenous administration of low doses (i.e., those typically  
96 used in self-administration experiments) of heroin and cocaine produce temporally distinct  
97 increases in the expression of *homer 1a (h1a)* and/or *arc* suggesting that both drugs induce  
98 neuronal activity across the striatum. In a second experiment, we used the catFISH technique  
99 to establish to what extent this activity occurs in overlapping vs. drug-specific neuronal  
100 populations. Based on electrophysiological evidence suggesting distinct neuronal activity  
101 produced by heroin vs cocaine (Chang et al., 1998), we predicted that administration of heroin  
102 following cocaine would activate non-overlapping neuronal populations across the striatum.

103

## 104 **Methods**

### 105 **Subjects**

106 A total of 66 male Sprague-Dawley rats ( $n = 37$  in Exp. 1 and  $n = 29$  in Exp. 2) from ENVIGO  
107 (Netherlands) were tested at a weight of 300-375 g. The rats were housed and tested in a  
108 temperature- and humidity-controlled room ( $21\pm 1^\circ\text{C}$ ; 50%) with a reversed 12 h light/dark cycle  
109 (lights on at 19:00 hours). The rats were housed in groups of 3 or 4 until surgery and  
110 individually thereafter. Food and water were provided *ad libitum* except during testing sessions.

111 All regulated procedures were carried out in accordance with the UK 1986 Animal Scientific  
112 Procedures Act (ASPA) and received approval from the relevant Animal Welfare and Ethics  
113 Review Board. After their arrival in the animal facilities, the rats were given a period of at least  
114 7 days before undergoing experimental procedures.

#### 115 ***Drugs***

116 Anesthesia was induced with 110 mg/kg of ketamine (Anesketin, Dechra) and 2 mg/kg of  
117 xylazine (Rompun, Bayer HealthCare). Cocaine and heroin hydrochloride (Johnson Matthey-  
118 MacFarlan Smith, Edinburgh, UK) were dissolved in sterile saline and infused i.v. at the doses  
119 specified in the next paragraphs. Each infusion consisted of a volume of 40  $\mu$ l of the  
120 appropriate drug solution delivered over 4 s. Saline-treated rats received equivalent volumes  
121 of saline.

#### 122 ***Intravenous catheter surgery***

123 The surgical procedures were similar to those recently described by Avvisati et al. (2019).  
124 Briefly, after anesthesia, an 11 cm silicone catheter (0.37-mm inner diameter and 0.94-mm  
125 outer diameter), sheathed at 3.4 cm from its proximal end by a silicone bead, was implanted in  
126 the right jugular vein, externalised at the nape of the neck, and attached to a cannula secured  
127 to the top of the skull with dental cement. Following surgery, rats were allowed to recover for  
128 at least 7 days. Catheter patency was maintained by flushing the catheters daily with 0.1 ml  
129 saline.

#### 130 ***Catheter patency test***

131 At the appropriate time (see next sections) the rats were killed via an i.v. infusion of  
132 pentobarbital (120 mg/kg in 200  $\mu$ l of saline) through the catheter. This also served as a  
133 catheter patency test: the rats that did not become ataxic and die within 5 s would be excluded  
134 from the data analysis. All catheters were found to be patent.

#### 135 ***Drug administration procedures***

136 **Experiment 1.** After recovery, the rats received, while briefly restrained, an i.v. infusion of  
137 either 400  $\mu$ g/kg cocaine ( $n = 18$ ) or 50  $\mu$ g/kg heroin ( $n = 19$ ) in their home cage. These doses  
138 were selected based on the findings of previous self-administration experiments (Caprioli et al.,

139 2007b; Caprioli et al., 2008). The rats received the lethal pentobarbital injection and were then  
140 decapitated at different time points after the cocaine or heroin infusion: 0 min ( $n = 3$  for both  
141 the cocaine and heroin groups), 8 min ( $n = 3$  for both the cocaine and heroin groups), 16 min ( $n$   
142 = 4 for both the cocaine and heroin groups), 25 min ( $n = 4$  for both the cocaine and heroin  
143 groups), and 35 min ( $n = 4$  and  $n = 5$  for the cocaine and heroin groups, respectively).

144 **Experiment 2.** After recovery, the rats were moved to testing chambers used for self-  
145 administration experiments (PRS Italia; see Avvisati et al., 2019). In order to reduce the  
146 potentially confounding effects of environmental novelty on drug-induced IEG expression  
147 (Uslaner et al., 2001; Paolone et al., 2007) we let the rats habituate to these chambers for 18 h  
148 before tethering them to the infusion lines. Food and water were available *ad libitum* during  
149 this habituation period and were removed immediately prior to tethering. The use of self-  
150 administration chambers allowed us to deliver drug infusions remotely via a computer-  
151 controlled infusion pump. The infusion pumps were programmed to start automatically, in the  
152 absence of the experimenter, 1 h after tethering. This way we avoided the confounding effects  
153 usually associated to signalled drug administration (Crombag et al., 1996) and/or handling. All  
154 rats received two i.v. infusions, 25 min apart, of: saline–saline ( $n = 4$ ), cocaine 800  $\mu\text{g}/\text{kg}$ –saline  
155 ( $n = 6$ ), cocaine 800  $\mu\text{g}/\text{kg}$ –cocaine 800  $\mu\text{g}/\text{kg}$  ( $n = 6$ ), cocaine 800  $\mu\text{g}/\text{kg}$ –heroin 100  $\mu\text{g}/\text{kg}$  ( $n =$   
156 6), or cocaine 800  $\mu\text{g}/\text{kg}$ –heroin 200  $\mu\text{g}/\text{kg}$  ( $n = 7$ ). To administer two separate injections  
157 through the same catheter, the infusion lines were back-filled with the appropriate drug  
158 solutions, separated by a tiny air bubble, just before tethering of the rats. The rationale for  
159 using higher doses of cocaine and heroin in Exp. 2 was to boost the magnitude of IEG  
160 expression. These doses were still within the range of those used in self-administration  
161 experiments (e.g. Zito et al., 1985; Dai et al., 1989; Roberts et al., 1989; Pettit and Justice, 1991;  
162 Shaham and Stewart, 1994; Wise et al., 1995; Mantsch et al., 2001; Wee et al., 2007; Mandt et  
163 al., 2012).

164 Five minutes after the second infusion, the rats were given 120 mg/kg pentobarbital,  
165 i.v., and, after decapitation, their brains were snap-frozen in isopentane at  $-50^{\circ}\text{C}$ .

166 **Brain slicing**

167 The brains were excised and snap-frozen in 400 ml of isopentane cooled to -50°C and later  
168 sectioned on a cryostat at 16- or 20- $\mu$ m thickness. In Exp. 1, sectioning started from the tip of  
169 the olfactory bulbs and brain sections were removed until the Sylvian fissure no longer reached  
170 the midline (+3.70 mm from bregma). At this point, either 100 or 80 sections were removed  
171 (when sectioning at 16  $\mu$ m and 20  $\mu$ m, respectively) to reach +2.00 mm from bregma at which  
172 point the sections contained anterior dorsal striatum and NAcc core (Fig. 2A). Two coronal  
173 sections per rat (16 or 20  $\mu$ m-thick) were obtained at this point. An identical procedure was  
174 used in Exp. 2 in order to collect two coronal sections containing NAcc core and shell, DMS, and  
175 DLS at +1.70 mm from bregma (Fig. 3A).

#### 176 *In situ hybridization*

177 Immediately after cutting, the brain tissue sections were mounted on Superfrost Plus slides. On  
178 the first day of staining, the slides were incubated in 10% neutral buffered formalin (Sigma, cat.  
179 No. HT501128-4L) for 20 min at 4°C, followed by 2x1 min washes in 1xPBS, and then serial  
180 dehydration in ascending concentrations of ethanol (5-min incubation in 50%, 70%, and 2x  
181 100%). Following this, the tissue was stored in 100% ethanol overnight. On day 2, the tissue was  
182 air dried, and then incubated with protease for 20 minutes, followed by 2x1 min washes in  
183 dH<sub>2</sub>O. Protease, probe and amplifier solutions were supplied by ACDBio as part of a  
184 commercially available RNAscope® kit (Advanced Cell Diagnostics, ACDBio). *Arc* and *h1a*  
185 hybridization probes (ACDBio, cat. No. 317071-C2 & 433261, respectively) were hybridized to  
186 fresh frozen brain coronal sections sliced on a Leica CM1900 cryostat. The signal was amplified  
187 with an RNAscope® Multiplex Fluorescent Reagent Kit (ACDBio, cat. No. 320850). The *arc* probe  
188 targeted the region spanning 1519-2621 base pairs of the *arc* gene mRNA, accession No.  
189 NM\_019361.1. The *h1a* probe targeted the 3' untranslated region of the *h1a* gene mRNA,  
190 spanning 5001-5625 base pairs, accession No. U92079.1.

191 The *arc* and *h1a* probes were applied (50  $\mu$ l per section), and the sections were  
192 incubated for 2h at 40°C in a humidity-controlled oven. After incubation with the probes, the  
193 signal was amplified at 4 separate stages with 15 min, 30 min, 15 min and 30 min of incubation  
194 in between (respectively) at 40°C in the hybridization oven. The probe and amplifier solutions  
195 were applied to the sections with the help of a hydrophobic pen barrier. There were 2x2 min



196 washes in wash buffer after each incubation (including after probe hybridization). Finally,  
197 sections were coverslipped and counterstained with DAPI mounting medium (Vector  
198 Laboratories, cat. No. H-1500) and left at 4°C overnight.

199

#### 200 ***Image acquisition and analysis***

201 Fluorescent signal was detected using a Zeiss Axioskop 2 plus epifluorescent microscope, and  
202 images were acquired using an Axiovision software (Zeiss).

203 Greyscale images were taken from both hemispheres of 2 adjacent sections for each rat  
204 at 20x magnification. This yielded four images per brain area for each rat. Final counts of DAPI-,  
205 *arc*-, and *h1a*-positive nuclei were averaged from these four images. The resulting images  
206 represented a region of interest (ROI) of 700 x 550 µm. These images were analysed using the  
207 RIO Montpellier extension of the ImageJ software (Baecker and Travo, 2006). Greyscale images  
208 were analysed separately for each channel – DAPI, Alexa 488 (*h1a*) and Cy3 (*arc*) – as follows.  
209 First, each DAPI image was analysed by applying a Gaussian blur filter (sigma = 2), then a  
210 “rolling ball” background subtraction algorithm (ball radius = 20), followed by the application of  
211 the default automatic global thresholding algorithm. This yielded a binary image which was  
212 then used to count objects selected on the basis of their size and circularity using the “analyse  
213 particles” function of ImageJ. The size criterion was set to 0.0045-0.045 square inches, and the  
214 circularity - to 0.7-1.00. This analysis resulted in a binary mask image containing only objects  
215 fulfilling the aforementioned criteria.

216 The images from the Alexa 488 and Cy3 channels were first adjusted for brightness so  
217 that the most visible signal was that coming from nuclear staining for *arc* and *h1a*. This was  
218 defined as any signal representing one or two bright dots close to each other, as opposed to  
219 cytoplasmic signal which is less bright and more diffused (Guzowski et al., 1999). A global  
220 threshold was then applied to the images (default algorithm), and the “analyse particles”  
221 function was used again to select only objects of 4-90 square pixels, and to create a binary  
222 image mask showing only the defined particles.

223 A Windows 10 Dell OptiPlex 7060 desktop computer ran a MATLAB script to overlay the  
224 three binary mask images and count instances where objects defined as DAPI nuclei coincided

225 with objects defined as either *arc* mRNA, *h1a* mRNA or both. The MATLAB code will be made  
226 available on request. Thus, IEG expression was measured by counting DAPI-positive (DAPI+) cell  
227 nuclei also positive for *h1a*, *arc*, or both.

228

### 229 **Statistical analyses**

230 The data from Exp. 1 were analysed by two-way mixed ANOVAs with time and IEG as fixed  
231 factors. Amount of IEG-positive cell nuclei (as percentage of all DAPI-stained nuclei) was the  
232 dependent variable. The data from Exp. 2 were analysed using a two-way ANOVA with brain  
233 area and treatment group as fixed factors. The outcome variable was overlap (expressed as  
234 percent of the cocaine-cocaine group). All analyses were carried out in SPSS 25 (IBM) software.  
235 An alpha value of 0.05 or less was used for determining statistically significant effects.

236

## 237 **Results**

### 238 **Experiment 1 (time course of *arc* and *h1a* expression following i.v. drug administration)**

239 Figure 2B shows the amount of *arc*- and *h1a*-positive nuclei in the NAcc core and DMS  
240 expressed as a percentage of all DAPI-positive nuclei and as a function of time elapsed since i.v.  
241 injections of cocaine and heroin. Table 1 shows the same data before conversion to percentage.

242 **Arc and *h1a* expression in the NAcc core.** Cocaine administration increased both *arc*  
243 and *h1a* mRNA levels in the NAcc core, but at different time points. A two-way mixed ANOVA  
244 showed non-significant main effects of IEG  $F(1,13) = 0.08$ ,  $p = .782$ ,  $\eta^2 = .006$ , and time  $F(4,13) =$   
245  $1.62$ ,  $p = .227$ ,  $\eta^2 = .333$  but a significant time x IEG interaction,  $F(4,13) = 7.93$ ,  $p = .002$ ,  $\eta^2$   
246  $= .977$ .

247 Heroin produced a similar pattern of mRNA expression, but the effect did not reach  
248 significance: a two-way mixed ANOVA showed non-significant main effects of IEG  $F(1,14) =$   
249  $2.32$ ,  $p = .150$ ,  $\eta^2 = .142$ , and time  $F(4,14) = 0.72$ ,  $p = .596$ ,  $\eta^2 = .17$ , and a non-significant time x  
250 IEG interaction,  $F(4,14) = 2.15$ ,  $p = .129$ ,  $\eta^2 = .38$ .

251 **Arc and *h1a* expression in the DMS.** As in the NAcc core, cocaine treatment increased  
252 IEG levels in a time-dependent manner. A two-way mixed ANOVA showed significant main

253 effects of IEG  $F(1,13) = 18.93, p = .001, \eta^2 = .593$ , and time  $F(4,13) = 5.36, p = .009, \eta^2 = .623$ ,  
254 and a significant time x IEG interaction,  $F(4,13) = 44.58, p < .001, \eta^2 = .932$ .

255 Heroin produced a similar effect. A two-way mixed ANOVA showed non-significant main  
256 effects of IEG  $F(1,14) = 3.17, p = .097, \eta^2 = .185$ , and time  $F(4,14) = 0.22, p = .924, \eta^2 = .059$ , but  
257 a significant time x IEG interaction,  $F(4,14) = 3.58, p = .033, \eta^2 = .506$ .

258

### 259 **Experiment 2 (overlap in neuronal populations activated by cocaine and heroin)**

260 Table 2 shows the average number of *arc*-only, *h1a*-only and double-stained cell nuclei as a  
261 function of brain area and drugs administered in experiment 2. Figures 4-7 show representative  
262 images from all brain areas analysed using catFISH.

263 Figure 3B shows the extent of overlap between neuronal populations activated by  
264 cocaine and heroin as a percent change from the cocaine-cocaine group. Overlap was  
265 quantified as the number of nuclei co-expressing *arc* and *h1a* expressed as a percent of all  
266 mRNA-positive nuclei (single and double-labelled). In all four brain areas examined, there was a  
267 substantial reduction in overlap when cocaine and heroin were administered in succession,  
268 relative to the overlap seen when cocaine was administered twice, and regardless of heroin  
269 dose (figure 3). A two-way mixed ANOVA showed a significant main effect of treatment group  
270  $F(3,19) = 20.97, p < .001, \eta^2 = .768$  and brain area  $F(3,57) = 3.40, p = .024, \eta^2 = .152$  but not  
271 treatment x brain area interaction  $F(9,57) = 0.79, p = .619, \eta^2 = .112$ .

272

### 273 **Discussion**

274 We have shown that intravenous injections of heroin and cocaine at doses typically self-  
275 administered by rats produce a quick and transient increase of *homer 1a* and *arc* expression  
276 across the striatum. More importantly, using the catFISH technique, we took advantage of the  
277 difference in timing of expression between the two IEGs to show that heroin and cocaine  
278 activate partly distinct neuronal populations in this brain area.

279 In line with our findings, previous studies have shown that heroin and cocaine increase  
280 *c-fos* levels in the ventral and dorsomedial striatum (Hope et al., 1992; Harlan and Garcia, 1998;

281 Uslaner et al., 2001; Ferguson et al., 2004; Paolone et al., 2007; Celentano et al., 2009). The IEG  
282 *c-fos* is a marker of neuronal activity expressed under similar conditions of *arc* and *homer 1a*  
283 (Guzowski et al., 2001). In addition, our findings indicate that this activity occurs in separate  
284 neuronal populations and may explain why only a small proportion of neurons show similar  
285 electrophysiological responses to heroin and cocaine (Chang et al., 1998).

286 It is likely that drug-induced IEG expression represents glutamatergic activity modulated  
287 by DA, because NMDA and DA D1 receptors play a key role in IEG expression through activation  
288 of CREB (Impey et al., 1998; Mattson et al., 2005; Surmeier et al., 2007; Guez-Barber et al.,  
289 2011; Tritsch and Sabatini, 2012), and both DA and glutamate levels are increased in the  
290 striatum following heroin and cocaine administration. Note, however, that DA release alone  
291 does not produce IEG expression in the absence of glutamatergic activity (Kreuter et al., 2004).  
292 In addition, NMDA receptor activity and DA transmission in the accumbens are necessary for  
293 food and cocaine self-administration, but not heroin self-administration (Ettenberg et al., 1982;  
294 Pettit et al., 1984; Pulvirenti et al., 1992; Kelley et al., 1997). Finally, D1 receptor-expressing  
295 MSNs in the dorsal striatum appear to be sufficient to sustain operant behaviour (Kravitz et al.,  
296 2012) and these neurons express IEGs (i.e. are activated) following cocaine administration.  
297 Thus, loss and gain of function studies have provided evidence that activity of cells in the  
298 striatum plays a key role for cocaine, but not heroin, reinforcement through DA and glutamate  
299 transmission. It remains to be determined what is the functional role of the distinct neuronal  
300 populations engaged by heroin relative to cocaine.

### 301 **A case for drug-specific neural circuitries**

302 Perhaps the most intriguing interpretation of the results presented here is that partly distinct  
303 neuronal populations activated by heroin and cocaine across the striatum are suggestive of  
304 dissociated circuitry processing the acute effects of the two drugs. There is already existing  
305 evidence that the striatum is functionally and structurally organised to accommodate circuits  
306 which operate in parallel but carry out separate functions. First, striatal medium spiny neurons  
307 (MSNs) are characterised by more or less excitable states, i.e. “up” and “down” states (Wolf et  
308 al., 2001; O'Donnell, 2003) and in order for MSNs to be excited (and to express IEGs), they must  
309 receive input from several sources which may include different combinations of amygdala,

310 hippocampus, thalamus, PFCx and VTA/SNc afferent inputs (Pennartz et al., 1994). Each of the  
311 brain areas sending these afferent projections: i) is affected differently by heroin, cocaine, and  
312 natural rewards (Chang et al., 1998; Mukherjee et al., 2018); ii) contains neuronal ensembles  
313 involved in distinct functions (Zelikowsky et al., 2014; Warren et al., 2016), and iii) might be  
314 comprised of genetically distinct projection neurons. Thus, considering the integrative function  
315 of the striatum, the diverse connectivity and specialised functions of its input regions, and the  
316 necessity for synchronised excitatory input to elicit action potentials from MSNs, it is quite  
317 possible that the activation of partly distinct neuronal populations in the striatum reflects the  
318 activation of dissociated circuitries. Here it must be noted that, although the afferent inputs of  
319 the striatum from limbic and cortical areas are topographically organised in a ventromedial-  
320 dorsolateral fashion, they are not constrained to perfectly defined striatal subregions, but are  
321 overlapping, with higher concentrations of certain afferents in, e.g., shell vs core (Voorn et al.,  
322 2004). It should also be considered that MSNs send collateral GABAergic projections to  
323 neighbouring MSNs. This mutual inhibition between MSNs is another functional-anatomical  
324 feature predisposing the accumbens and rest of striatum to accommodate neuronal ensembles  
325 embedded in distinct circuitries – whilst one ensemble is active, it can decrease the activity in  
326 other ensembles so that only one computation is taking place over others (Pennartz et al.,  
327 1994). The experiments presented here are only suggestive of distinct striatal circuitry engaged  
328 by heroin and cocaine. Future studies should address this hypothesis by expanding on our  
329 findings in three ways. First, single-cell quantitative PCR studies can further elucidate  
330 phenotypic differences between neuronal populations activated by heroin and cocaine in terms  
331 of their genetic make-up (Hrvatin et al., 2018). Second, retro- and anterograde labelling studies  
332 in conjunction with immunohistochemistry can reveal whether these neuronal populations  
333 connect to distinct up- and downstream targets. And third, selective loss- and gain-of-function  
334 studies can be used to test whether inactivation of neurons responding to cocaine in the dorsal  
335 striatum and accumbens core would impair heroin reinforcement. The Daun02 technique (Koya  
336 et al., 2009; Koya et al., 2016) would be a useful technique in this regard, as well as other  
337 techniques which manipulate neuronal ensembles such as the TetTag approach using the Fos-

338 tTA mouse line combined with optogenetics (Reijmers and Mayford, 2009; Liu et al., 2012; Du  
339 and Koffman, 2017).

#### 340 **Methodological considerations**

341 Two caveats to the experimental design used here are worthy of mention. There are known  
342 differences between the effects of non-contingent vs contingent exposure to heroin and  
343 cocaine (e.g. Galici et al., 2000; Lecca et al., 2007; Radley et al., 2015). In the present study we  
344 administered heroin and cocaine in a non-contingent but unsignalled manner as we were  
345 interested in comparing the acute pharmacological effects of these two drugs using IEG  
346 expression as a marker of neuronal activation. Contingent administrations (e.g., self-  
347 administration) require repeated exposure to drugs over several test sessions which has been  
348 shown to produce habituation to IEG expression (Hope et al., 1992; Unal et al., 2009). Of  
349 course, we recognise the value of studying the encoding of drug-related information in the  
350 striatum during and after periods of drug self-administration. Future studies could employ *in*  
351 *vivo* imaging techniques such as the UCLA/Inscopix© miniscope to address this question  
352 directly. A second, somewhat related caveat is that our paradigm includes a multi-substance  
353 component. It is possible that circuit activity may differ following polysubstance versus single-  
354 drug use histories. However, electrophysiological evidence from rats self-administering both  
355 substances is congruent with our findings (Chang et al., 1998). Also, we administered only two  
356 injections of cocaine and/or heroin to drug-naïve rats so it is unlikely that any long-term  
357 polysubstance use effects would have influenced our observations.

#### 358 **Conclusion**

359 In summary, we found a significant dissociation in the neuronal populations responding to self-  
360 administration doses of heroin versus cocaine, as indicated by *arc* and *homer 1a* expression.  
361 Our findings provide a proof of concept that heroin and cocaine effects on the brain must be  
362 studied as separate phenomena, adding to the evidence of major differences between the  
363 various drugs of abuse (for a review, see Badiani et al., 2011). Although the functional  
364 significance of these differences remains to be fully explored, they might have implications for  
365 both research and treatment. It is remarkable, for example, that the functional or anatomical  
366 integrity of the dopaminergic system is required for the reinforcing properties of cocaine but

367 not of heroin (e.g., Ettenberg et al., 1982; Pettit et al., 1984; Pisanu et al., 2015), that distinct  
368 projections from the PFCx to the shell of the NAcc are implicated in the relapse to cocaine  
369 versus heroin seeking after abstinence (Peters et al., 2008; Bossert et al., 2012), and that basic  
370 environmental manipulations gate in opposite directions the reinforcing, affective, and  
371 neurobiological response to heroin versus cocaine in rats and humans (Uslaner et al., 2001;  
372 Ferguson et al., 2004; Caprioli et al., 2007a; Paolone et al., 2007; Caprioli et al., 2008; Caprioli et  
373 al., 2009; Celentano et al., 2009; Montanari et al., 2015; Avvisati et al., 2016; De Pirro et al.,  
374 2018; De Luca et al., 2019).

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622

623 **Table 1.** Mean (SE) number of *arc*- and *h1a*-stained cell nuclei as a function of brain area and  
624 drug administered in Exp. 1. The brains were excised at different time points after drug  
625 administration: 0 min ( $n = 3$  for both the cocaine and heroin groups), 8 min ( $n = 3$  for both the  
626 cocaine and heroin groups), 16 min ( $n = 4$  for both the cocaine and heroin groups), 25 min ( $n =$   
627 4 for both the cocaine and heroin groups), and 35 min ( $n = 4$  and  $n = 5$  for the cocaine and  
628 heroin groups, respectively).

629 **Table 2.** Mean (SE) number of *h1a*-only, *arc*-only and double-stained cell nuclei as a function of  
630 brain area and drugs administered, 25 min apart, in Exp. 2: saline-saline ( $n = 4$ ), and cocaine  
631 (800  $\mu\text{g}/\text{kg}$ )-saline ( $n = 6$ ), cocaine (800  $\mu\text{g}/\text{kg}$ )-cocaine (800  $\mu\text{g}/\text{kg}$ ) ( $n = 6$ ), cocaine (800  $\mu\text{g}/\text{kg}$ )-  
632 heroin 100  $\mu\text{g}/\text{kg}$ , ( $n = 6$ ), and cocaine (800  $\mu\text{g}/\text{kg}$ )-heroin 200  $\mu\text{g}/\text{kg}$  ( $n = 7$ ).

633

634 **Figure 1. The catFISH paradigm.** Working hypothesis based on Guzowski et al. 2005: the  
635 expression of mRNA encoding for *h1a* and *arc* should be detectable at different time points  
636 after drug administration. **A)** *arc* mRNA expression in the nucleus should peak at about 5 min  
637 after drug administration, whereas *h1a* mRNA should peak at about 30 min. **B)** Overlap in the  
638 expression of Drug 1-induced *h1a* mRNA and *arc* mRNA should be observed at time 30 min (25  
639 min after Drug 1 and 5 min after Drug 2).

640 **Figure 2. Effect of single drug injections on IEG expression.** Time-course of *arc* mRNA and *h1a*  
641 mRNA expression in experiment 1. **A)** Regions of interest (plate from Paxinos & Watson, 1986).  
642 **B)** Average number of *arc*- or *homer 1a* (*h1a*)-positive cell nuclei as a function of brain area and  
643 administered drug (expressed as a percentage of all DAPI-positive nuclei). The brains were  
644 excised at different time points after drug administration: 0 min ( $n = 3$  for both the cocaine and  
645 heroin groups), 8 min ( $n = 3$  for both the cocaine and heroin groups), 16 min ( $n = 4$  for both the  
646 cocaine and heroin groups), 25 min ( $n = 4$  for both the cocaine and heroin groups), and 35 min  
647 ( $n = 4$  and  $n = 5$  for the cocaine and heroin groups, respectively).

648 **Figure 3. Overlap in the neuronal populations engaged by heroin and cocaine.** Co-expression  
649 of *arc* and *h1a* mRNAs in experiment 2. **A)** Regions of interest. (plate from Paxinos & Watson,  
650 1986). **B)** Overlap expressed as % of overlap in the cocaine-cocaine condition as a function of  
651 brain area and drugs administered, 25 min apart, in Exp. 2: saline-saline ( $n = 4$ ), cocaine (800  
652  $\mu\text{g}/\text{kg}$ )-saline ( $n = 6$ ), cocaine (800  $\mu\text{g}/\text{kg}$ )-cocaine (800  $\mu\text{g}/\text{kg}$ ) ( $n = 6$ ), cocaine (800  $\mu\text{g}/\text{kg}$ )-  
653 heroin 100  $\mu\text{g}/\text{kg}$  ( $n = 6$ ), and cocaine (800  $\mu\text{g}/\text{kg}$ )-heroin 200  $\mu\text{g}/\text{kg}$  ( $n = 7$ ).

654 **Fig. 4. Representative microscope images taken from the NAcc core.** DAPI-stained cell nuclei  
655 (blue) co-express either only *h1a* (green), only *arc* (red), or both. The columns show green and  
656 red channels separately and then merged. Taken from Nacc core. Scale bars correspond to 0.1  
657 mm. Arrows point to mRNA-positive nuclei.

658 **Fig. 5. Representative microscope images taken from the NAcc shell.** DAPI-stained cell nuclei  
659 (blue) co-express either only *h1a* (green), only *arc* (red), or both. The columns show green and  
660 red channels separately and then merged. Scale bars correspond to 0.1 mm. Arrows point to  
661 mRNA-positive nuclei.

662 **Fig. 6. Representative microscope images taken from the DMS.** DAPI-stained cell nuclei (blue)  
663 co-express either only *h1a* (green), only *arc* (red), or both. The columns show green and red  
664 channels separately and then merged. Scale bars correspond to 0.1 mm. Arrows point to mRNA-  
665 positive nuclei.

666 **Fig. 7. Representative microscope images taken from the DLS.** DAPI-stained cell nuclei (blue)  
667 co-express either only *h1a* (green), only *arc* (red), or both. The columns show green and red  
668 channels separately and then merged. Scale bars correspond to 0.1 mm. Arrows point to mRNA-  
669 positive nuclei.

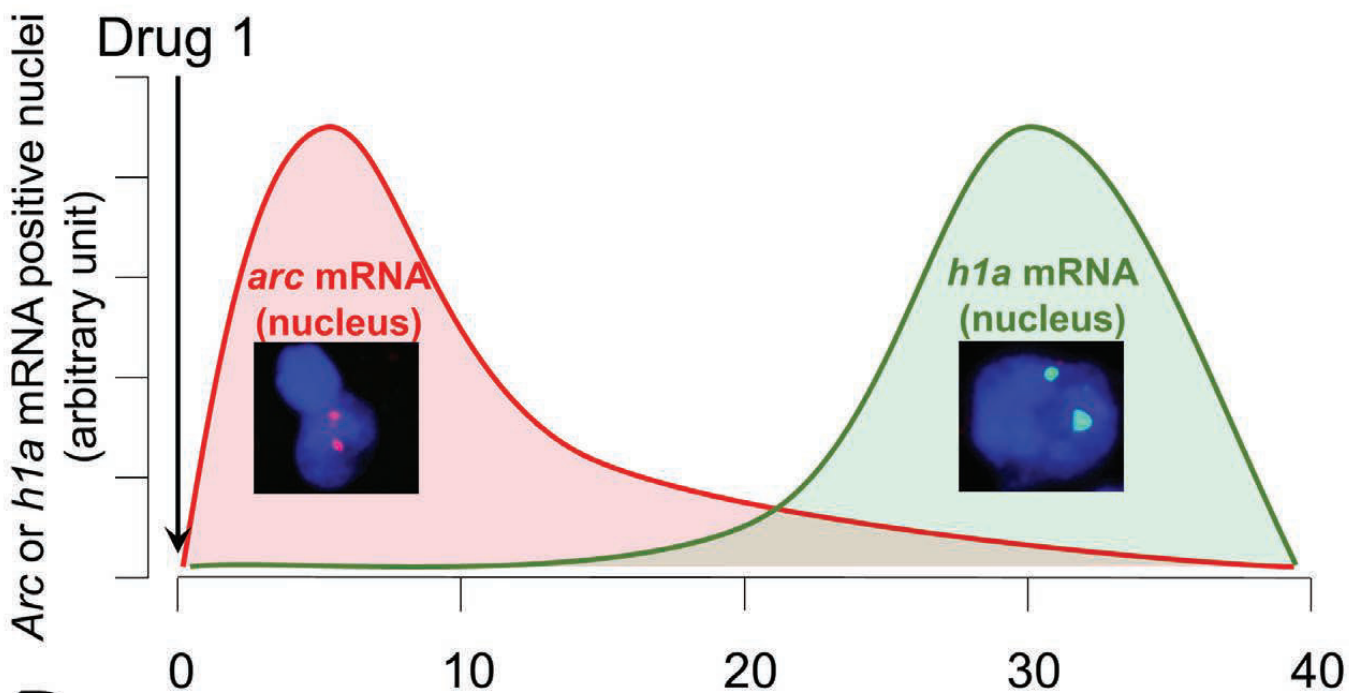
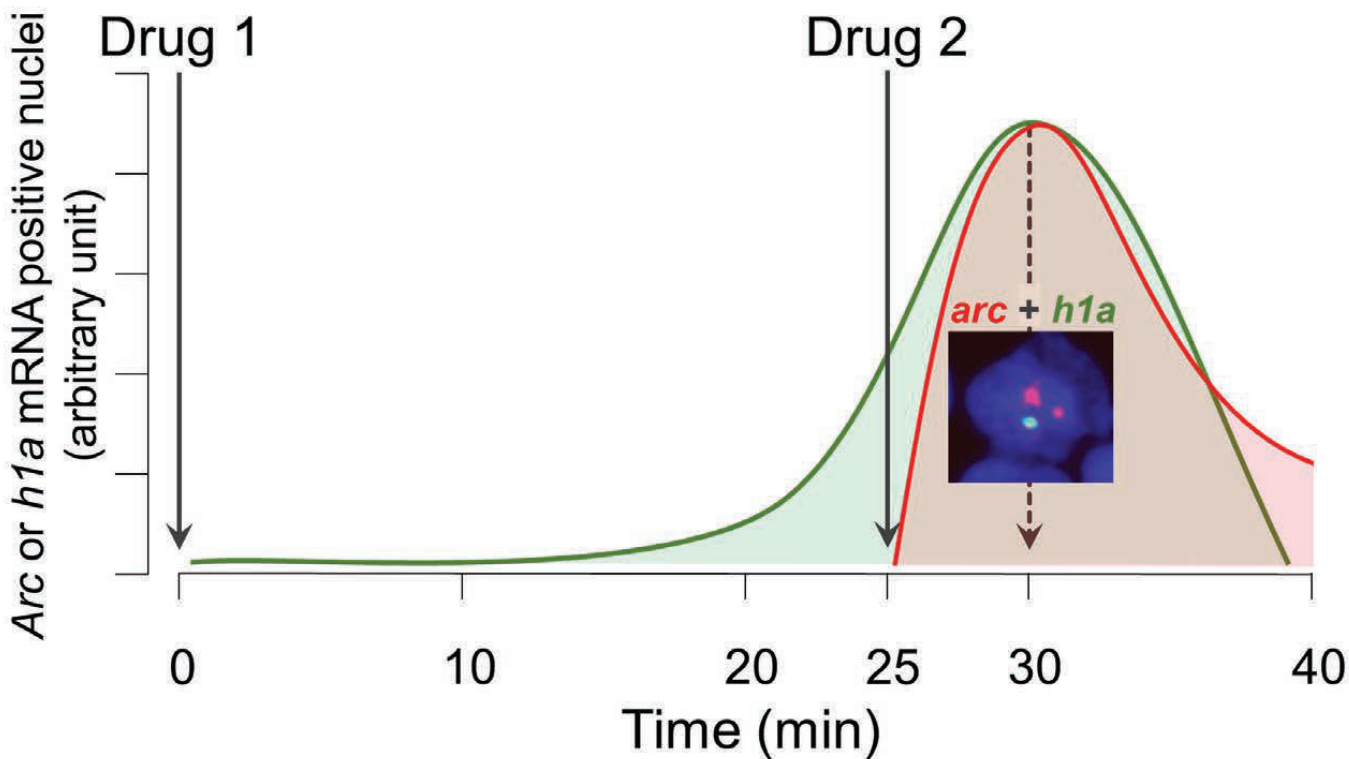
Table 1.

	NAcc				DS			
	Cocaine (400 µg/kg)		Heroin (50 µg/kg)		Cocaine (400 µg/kg)		Heroin (50 µg/kg)	
	Arc	H1a	Arc	H1a	Arc	H1a	Arc	H1a
0 min	19.50 (1.52)	12.50 (4.44)	18.42 (6.86)	11.33 (1.8)	30.10 (1.97)	16.42 (5.85)	21.17 (6.59)	14.83 (1.02)
8 min	25.50 (5.36)	9.75 (2.38)	26.92 (13.66)	8.83 (2.71)	44.58 (4.43)	11.83 (1.91)	26.92 (8.21)	14.58 (5.27)
16 min	16.31 (3.35)	14.81 (1.22)	13.88 (3.63)	8.63 (2.94)	21.81 (3.08)	32.19 (3.73)	12.31 (2.86)	20.88 (5.59)
25 min	9.94 (2.78)	17.81 (0.82)	18.19 (3.95)	18.69 (4.29)	15.00 (3.89)	48.88 (5.99)	16.44 (3.76)	31.75 (7.3)
35 min	11.00 (1.67)	26.25 (5.13)	8.00 (2.22)	14.6 (4.26)	11.25 (1.44)	58.50 (1.52)	7.50 (2.2)	28.35 (4.58)

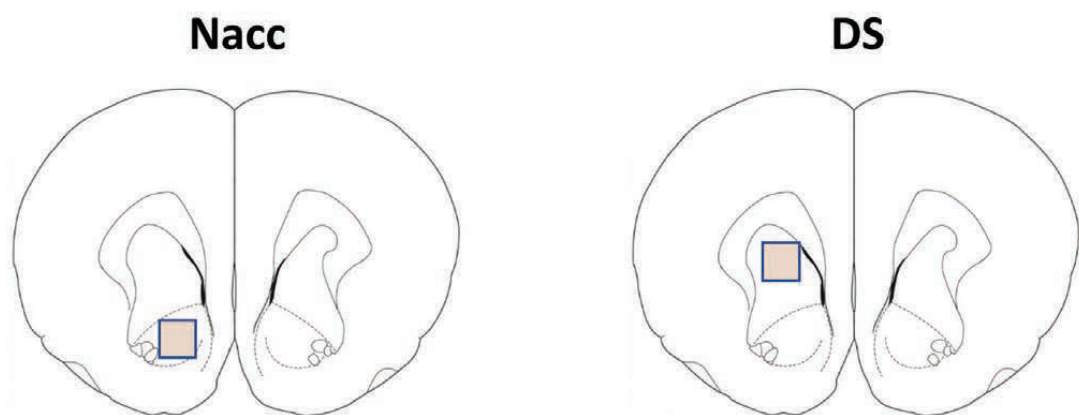


Table 2.

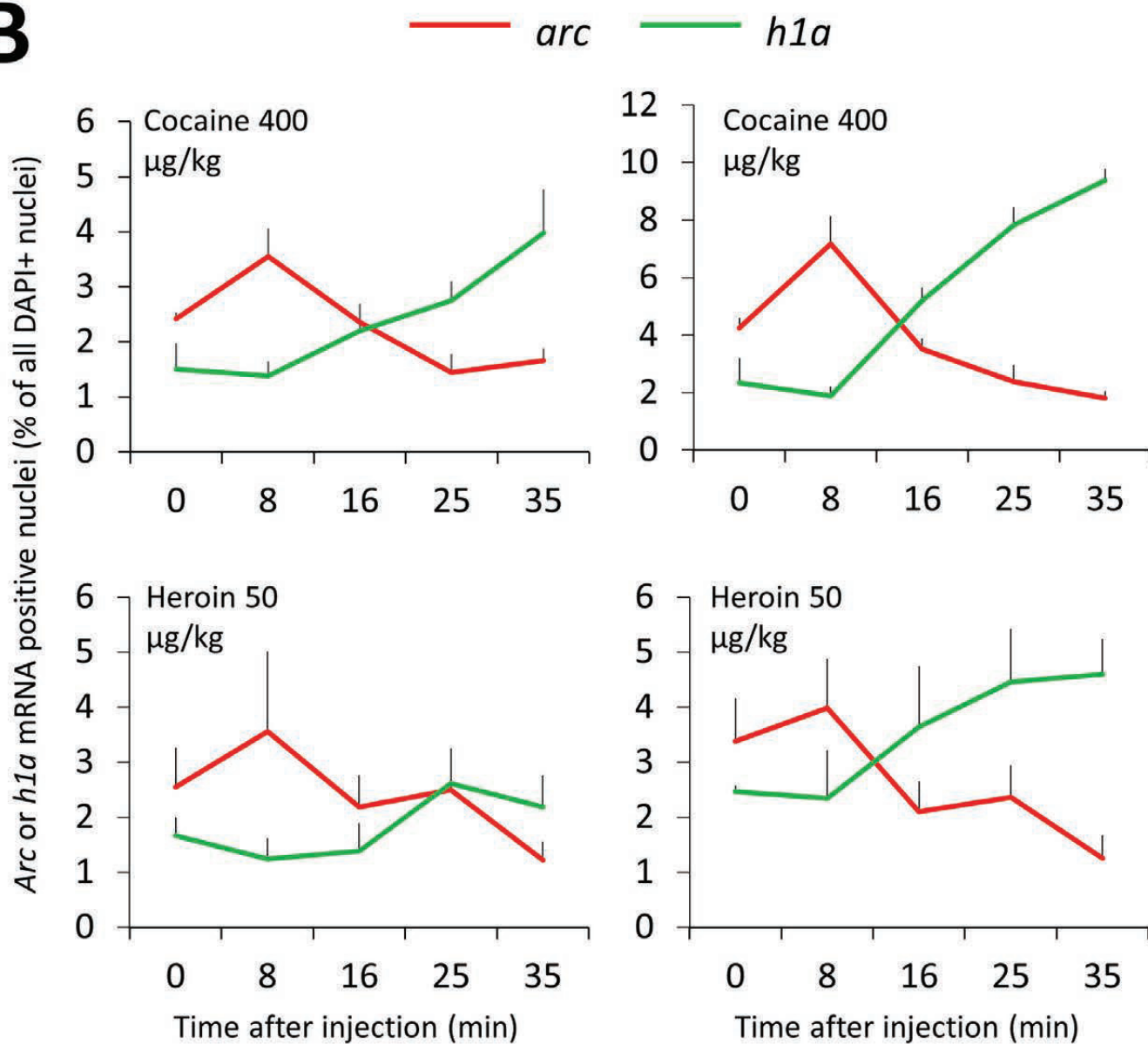
		NAcc core	NAcc shell	DMS	DLS
1 <sup>st</sup> saline 2 <sup>nd</sup> saline	h1a	4.94 (0.66)	4.44 (1.61)	5.98 (2.18)	8.5 (2.54)
	arc	2.38 (0.94)	2.48 (0.75)	1.98 (0.95)	3.75 (1.59)
	double	0.13 (0.07)	0.38 (0.16)	0.13 (0.07)	0.13 (0.13)
1 <sup>st</sup> cocaine (800 µg/kg) 2 <sup>nd</sup> saline	h1a	21.5 (4.44)	8.67 (2.36)	51.1 (7.99)	65.1 (7.83)
	arc	4.25 (0.77)	3.13 (0.68)	5.58 (1.77)	4.33 (0.95)
	double	1.46 (0.25)	1.67 (0.35)	3.63 (0.43)	4.46 (1.49)
1 <sup>st</sup> cocaine (800 µg/kg) 2 <sup>nd</sup> cocaine (800 µg/kg)	h1a	20.54 (5.45)	11.00 (2.87)	42.17 (8.65)	49.71 (7.5)
	arc	8.08 (0.59)	5.13 (0.67)	12.54 (1.99)	15.33 (3.89)
	double	5.46 (1.04)	3.33 (0.77)	14.17 (3.17)	21.33 (5.19)
1 <sup>st</sup> cocaine (800 µg/kg) 2 <sup>nd</sup> heroin (100 µg/kg)	h1a	20.33 (3.72)	8.04 (2.36)	58.67 (16.42)	66.04 (11.36)
	arc	23.29 (9.55)	18.92 (9.13)	3.63 (0.70)	3.04 (1.06)
	double	5.00 (2.07)	2.46 (0.87)	4.17 (1.19)	3.63 (0.96)
1 <sup>st</sup> cocaine (800 µg/kg) 2 <sup>nd</sup> heroin (200 µg/kg)	h1a	18.96 (4.33)	7.57 (1.75)	50.68 (7.34)	56.46 (7.64)
	arc	12.11 (1.53)	12.61 (2.91)	5.17 (0.74)	4.32 (1.16)
	double	3.14 (0.59)	1.75 (0.49)	3.11 (0.46)	3.93 (0.93)

**A****B**

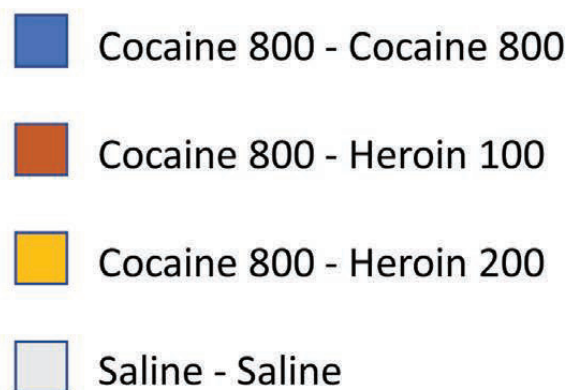
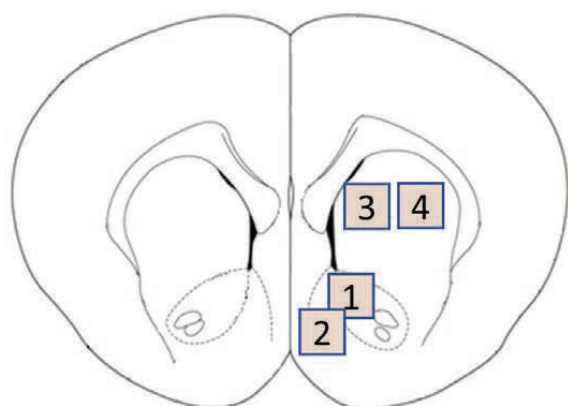
**A**



**B**



**A**



**B**

Co-expression of *arc* and *h1a* mRNAs  
(% of cocaine-cocaine)

