

1 **NFκB1 and NFκB2 gene expression in the prefrontal cortex and hippocampus of early life**
2 **stressed mice exposed to cocaine-induced conditioned place preference during adolescence**

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23

24 **ABSTRACT**

25 **Background:** Neuro-immune pathways contribute to the onset and maintenance of cocaine-
26 seeking behaviors, particularly through activation of NFκB signaling in the brain. However, the
27 molecular mechanisms of this relationship are still not completely understood, especially
28 considering the effects of early life stress, a major risk factor to cocaine addiction. The goal of
29 this study was to investigate NFκB1 and NFκB2 gene expression in the prefrontal cortex (PFC)
30 and hippocampus of mice exposed to early life stress and cocaine-induced conditioned place
31 preference (CPP) within adolescence. Male BALB/c mice were randomly assigned to one of four
32 groups: animal facility reared (AFR) with or without CPP training; maternal separation (MS)
33 with or without CPP training. The MS animals were subjected to daily 3-h maternal separation
34 from postnatal day (PND) 2 to 15. CPP was performed following three sequential phases:
35 habituation (PND 34), conditioning (PND 35 to PND 44) and post-conditioning test (PND 45).
36 Gene expression was determined by qPCR.

37 **Findings:** NFκB1 mRNA levels were decreased in the PFC of animals exposed to CPP
38 compared to drug-naïve animals, while no difference was detected regarding rearing conditions.
39 NFκB2 expression was upregulated in the PFC of animals exposed to CPP when compared to
40 drug-naïve animals, particularly in animals exposed to MS with higher CPP scores. No
41 significant effects were detected in the hippocampus.

42 **Conclusions:** Cortical NFκB2 up-regulation may be involved with the enhanced motivational
43 salience for cocaine-paired cues observed in animals exposed to MS during adolescence.

44

45 **Key words:** Cocaine, Conditioned place preference, Early life stress, Maternal separation, NFκB
46 mRNA.

47 INTRODUCTION

48 Early life stress can promote immune alterations across lifespan, including increased
49 innate immunity/inflammation (1). Following chronic stress, up-regulation of an array of genes
50 involved in pro-inflammatory pathways has been documented in both peripheral immune cell
51 populations and brain-specific cells, such as astrocytes and microglia (2). Among possible
52 pathways, Toll-like Receptors (TLRs) are the first line of defense against invading
53 microorganisms and are responsible for triggering innate immune responses. TLRs expression
54 are also modulated by stress exposure, inducing activation of the transcription factor nuclear
55 factor-kappa B (NFκB) (3).

56 The NFκB regulates the expression of many pro-inflammatory cytokine genes (e.g.,
57 Interleukin-1, Interleukin-6, Tumor Necrosis Factor-Alpha) and it is also involved in the
58 processes of synaptic plasticity and memory formation (4, 5). Therefore, chronic activation of
59 this innate immune pathway can enhance inflammation in brain areas associated with mental
60 disorders, and, consequently, contributing to the development of anxiety, depression and
61 substance abuse disorder (6). In this sense, both clinical and pre-clinical investigations have
62 pointed out a role for NFκB in the pathophysiology of cocaine drug addiction (7, 8). For
63 instance, blocking the expression of NFκB in the nucleus accumbens (NAc) reduces the overall
64 preference for cocaine use in mice, and decreases the number of dendritic spines in the NAc (8).
65 Additionally, NFκB signaling in the frontal cortex was associated with cocaine-induced
66 neurotoxicity and cocaine-induced memory impairments (9).

67 The NFκB family is composed of five proteins: relA (p65), NFκB1 (p50 and its precursor
68 p105), NFκB2 (p52 and its precursor p100), c-Rel and RelB (10). Both p52 and p50 subunits do
69 not have a transcription activation domain, so they need to interact with other factors to regulate

70 transcription (10). For this reason, RelB will interact with p100 and p52, and RelA and c-Rel will
71 interact with p50 (11). Furthermore, both NFκB1 and NFκB2 have an important role in the
72 regulation of immune and inflammatory responses, and the dysregulation in the expression of
73 these genes can provoke the overproduction of pro-inflammatory cytokines, which are associated
74 with several inflammatory disorders (12). Although NFκB1 and NFκB2 present similar structure
75 and both are responsible for immune and inflammatory responses, they also have distinct
76 functions. NFκB1 is mainly responsible for innate immune response, as well as controlling
77 lymphocyte and macrophage function. On the other hand, NFκB2 has a main role in the
78 organogenesis of peripheral lymphoid tissues and B-cell development, with a more pronounced
79 function related to adaptive immunity (12).

80 Clinical and epidemiological evidence demonstrated that early life stress promotes higher
81 vulnerability to substance use disorder development (13, 14). Therefore, using the Maternal
82 Separation (MS) model of early life stress we have investigated the effects of neonatal chronic
83 stress on cocaine-related phenotypes within adolescence, using the well-known Conditioning
84 Place Preference (CPP) behavioral paradigm in mice (15). In this experiment stressed and control
85 animals were exposed to cocaine and saline solution during 10 days of CPP training, and tested
86 for cocaine/saline preference. We observed that adolescent male mice exposed to MS presented
87 higher preference for cocaine paired cues and reduced cortical brain-derived neurotrophic factor
88 (BDNF) gene expression, suggesting that exposure to early life stress can enhance the
89 motivational salience for cocaine-paired cues, and that this effect may be partially mediated by
90 BDNF signaling (Figure 1).

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118 Maternal Separation (MS)

119 MS is a robust early life stress protocol (18), where pups were separated daily from their
120 dams for a period of 180 min (3 p.m. to 6 p.m.) from PND2 to PND14. Dams were transferred to
121 another room and allocated in a new cage to avoid ultrasonic vocalization between dam and
122 pups. The litter was kept together during the MS protocol in a new cage with clean bedding, and
123 the temperature was controlled with a heating pad (33 ± 2 C) to maintain adequate body
124 temperature. After the MS protocol both the litter and the dam were returned to the home cage.
125 AFR group was left undisturbed until weaning, except for routine cage cleaning.

126

127 Cocaine-induced CPP

128 The Cocaine-induced CPP apparatus consisted two (15 x 15 cm) acrylic chambers
129 divided by an acrylic barrier (3.2mm tick). Each chamber had a different visual cue printed on a
130 white floor and background. One chamber had black lines and the other black circles. During the
131 adolescence period three phases of the CPP were executed: Habituation (PND 34), conditioning
132 (PND 35 to PND 44), and post-conditioning test (PND45). Mice spent 30 min exploring the
133 chamber each day during all phases. During the habituation the animals had free access to both
134 chambers, and the time in each of the chambers was recorded to verify pre-conditioning place
135 preference. No animals exceeded the threshold of 70% time spent in only one chamber. During
136 the conditioning procedure mice were assigned to receive cocaine in the chamber that they spent
137 less time during habituation and saline in the other chamber. During the period of 10 days each
138 animal received alternately cocaine and saline solution (5 days each). Half of the animals

139 received cocaine on days 1, 3, 5, 7, 9 and saline solution on days 2, 4, 6, 8, 10. The other half
140 was exposed to the inverse pattern. After the last conditioning day, to test for cocaine-induced
141 CPP the animals were allowed to explore both chambers with no drug/saline administration. To
142 calculate the CPP score the time spent in seconds in the cocaine-paired chamber was subtracted
143 by the time spent during the pre-test habituation phase. For more detailed information review
144 Viola et al (2016). To perform the cocaine administration for the CPP a concentration of 20
145 mg/ml was obtained after dissolving Cocaine Hydrochloride in a sterile 0.9 % saline solution. A
146 cocaine dose of 20 mg/kg (i.p.) was used since previous studies with BALB/c mice revealed that
147 this cocaine concentration optimally induces place preference (19).

148

149 **Brain sample preparation**

150 Two hours after the CPP test at PND 45 euthanasia was performed by cervical dislocation
151 without anesthesia in order to avoid pharmacological interference. After decapitation the brains
152 were removed and stored overnight at 4 C in RNAlater (Ambion) solution (1 ml). All areas of the
153 PFC were dissected by freehand technique without the olfactory bulb. After storing the samples
154 at -80 C the hemispheres were randomly alternated within the groups, so half of the samples
155 were from the left hemisphere and half from the right hemisphere.

156

157 **Transcript mRNA levels**

158 Total RNA was extracted using QIAzol (Qiagen) according to the manufacturer's
159 protocol and suspended in RNase-free water. The RNA concentration was quantified using Qubit
160 RNA Broad Range Assay in a Qubit Fluorometer 2.0 (Life Technologies). 250 ng of RNA from
161 each sample was transformed into cDNA using miScript II RT kit (Qiagen) for mRNA gene

162 expression. The resulting 20 ul of cDNA were diluted with 80 ul of RNase-free water to a total
163 volume of 100ul and stored at -20 C. 1 ul of each cDNA sample was used for RT-qPCR reaction,
164 executed in a StepOne PCR machine (Applied Biosystems), using miScript SYBR Green PCR
165 Kit (Qiagen). Each reaction was run in duplicate, and the $\Delta\Delta C_t$ method was used to calculate the
166 relative fold change expression with the AFR drug-naïve group as a reference. The Primers:
167 NFkB1 (QT00154091), NFkB2 (QT00129864) and GAPDH (QT01658692) were purchased
168 from Qiagen.

169

170 **Statistical analyses**

171 All data are presented as group mean \pm standard error of the mean, and tested for
172 normality using the Shapiro-Wilk test. A two-way analysis of variance (ANOVA) followed by
173 Bonferroni post-tests was used to investigate the difference in mRNA expression between
174 groups. A P-value of < 0.05 was considered statistically significant. All statistical analyses were
175 conducted using SPSS software v.20.0 (SPSS, Chicago, IL, USA) and the graphs were generated
176 using GraphPad Prism 6 (GraphPad Software Inc., La Jolla, CA, USA).

177

178 **RESULTS**

179 **Cocaine-induced CPP effects on NFkB mRNA levels**

180 We observed a decrease in NFkB1 mRNA levels in the PFC in animals exposed to
181 cocaine-CPP compared to drug-naïve animals showing a significant treatment effect ($F(3,24) =$
182 $4.38, p < 0.05$) (Figure 2A). No significant group effect was detected between AFR and MS
183 animals. In the Hippocampus we did not observed significant difference when comparing

184 animals exposed to cocaine-CPP and drug naïve animals or MS and AFR groups ($F(3,24) = 1.21$,
185 n. s.) (Figure 2C).

186 Regarding NF κ B2 mRNA levels in the PFC we identified higher levels in animals
187 exposed to cocaine-CPP when compared to drug-naïve animals demonstrating a significant
188 treatment effect ($F(3,24) = 33.11$, $p < 0.01$) (Figure 2B). When comparing the MS-CPP group to
189 the AFR drug-naïve group we observed statistically significant difference (Bonferroni, $p < 0.01$),
190 but this difference could not be observed when comparing AFR-CPP group to AFR drug-naïve
191 animals (Bonferroni, n. s.). No significant group effect between AFR and MS from the same
192 treatment group was detected. Moreover, there was no significant group or treatment effects on
193 NF κ B2 mRNA levels in the hippocampus ($F(3,24) = 3.21$, n. s.).

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FIGURE 2 HERE

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198 **DISCUSSION**

199 The data presented here suggest a novel role for the NF κ B1 and NF κ B2 genes regarding
200 the acquisition of cocaine-induced CPP, showing dynamically opposite effects in their
201 transcriptional activity following CPP test. Interestingly, we suggest that NF κ B2 up-regulation in
202 the PFC may be involved with the enhanced motivational salience for cocaine-paired cues
203 observed in animals exposed to MS; while NF κ B1 down-regulation was associated with cocaine-
204 induced CPP effects in the cortex, but its expression was not affected by stress-related effects.

205 Our results are in agreement with a previous study that observed a decrease in NF κ B1
206 protein levels in the frontal cortex, an affect associated with memory impairments in animals

207 treated with intraperitoneal cocaine injections of 15mg/kg during 20 days (9). Thus although we
208 did not test memory performance, the downregulation in the expression of cortical NFκB1
209 observed in our experiments could be involved with cocaine-induced neuroplasticity and PFC-
210 dependent cognitive functioning alterations. Such hypothesis is supported by evidence that
211 activation of NFκB1 is necessary for the regulation of synaptic plasticity and memory formation
212 (4), in addition to its role of regulating innate immune functioning.

213 On the other hand, the enhanced motivational salience for cocaine-paired cues observed
214 in the animals exposed to early life stress could possibly be associated with altered NFκB2 gene
215 expression. Evidence suggests that NFκB2 transcription is induced by the B-cell activating factor
216 (BAFF) (20). Although we did not investigate the expression pattern of BAFF, it is possible to
217 hypothesize that an increase in BAFF levels triggers the activation of the IκB kinase complex,
218 which induces the processing of p100 to p52, both subunits of NFκB2. There is evidence
219 indicating that exposure to cocaine can enhance the release of BAFF, which will activate the
220 cascade and increase the levels of NFκB2 (21). Although NFκB2 function in the has been
221 understudied, the cortical BAFF-NFκB2 signaling pathway potentially has an important role in
222 regulating cue-induced drug craving and drug-seeking behavior, and altered functioning of this
223 pathway could be involved with higher cocaine-induced CPP attributed to early maternal
224 separation exposure. There is evidence suggesting that the acquisition of CPP involves histones
225 alterations resulting in increased expression of several genes including NFκB2 (22). Here we
226 presented for the first time the effects of early life stress on cortical NFκB2 gene expression after
227 CPP. In this regard, there is one study that found a pattern of demethylation in the promoter
228 region of the NFκB2 in a genome-wide DNA methylation analysis from umbilical cord blood of
229 neonates exposed to prenatal anxiety or depression of their mothers (23).

230 Moreover, while NFκB1 is constantly expressed in cortical regions (<http://mouse.brain->
231 [map.org/experiment/show/77454688](http://mouse.brain-map.org/experiment/show/77454688)), NFκB2 has a low expression in the brain cortex during
232 basal conditions (<http://mouse.brain-map.org/experiment/show/76098410>). Thus, activation of
233 this “silenced” gene in the PFC may be a potential mechanism for higher vulnerability to cocaine
234 and other drugs of abuse following repeated stress due to increased neuroinflammatory signaling.
235 In this sense, previous evidence have showed that NFκB2 was downregulated in
236 methamphetamine low drinking animals but upregulated in methamphetamine high drinking
237 animals in the NAc (24). This is relevant since the majority of NAc neurons receive significant
238 glutamatergic inputs from the PFC, an effect involved in the transferring of information
239 regarding drug-associated environmental stimuli (25). Thus, synaptic alterations between the
240 PFC and the NAc modulate acquisition of CPP behavior, and NFκB2 up-regulation is another
241 molecular marker involved in this circuitry.

242 In conclusion, our study indicates that exposure to cocaine-induced CPP is associated
243 with reduced NFκB1 and increased NFκB2 gene expression in the PFC. This increase in NFκB2
244 levels after exposure to cocaine use is potentiated by the neonatal stress. Thus, this data raise the
245 possibility that the pro-inflammatory effects of cocaine may be enhance the long lasting pro-
246 inflammatory processes derived from early stress exposure, contributing to the effects of stress
247 on higher vulnerability for cocaine addiction and drug-related behaviors during periadolescence.
248 Therefore, it will be interesting in future studies to determine whether manipulation of NFκB2
249 signaling in the PFC could attenuate the effects of early life stress on cocaine-related phenotypes.

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253 LIST OF ABBREVIATIONS

254

255 AFR – Animal Facility Reared

256 ANOVA – Analysis of Variance

257 BAFF – B-cell Activating Factor

258 BDNF – Brain-derived Neurotrophic Factor

259 CEUA – Ethics Committee on Animal Use

260 CPP – Conditioned Place Preference

261 MS – Maternal Separation

262 NAc – Nucleus Accumbens

263 NFkB – Nuclear Factor-kappa B

264 PCR – Polymerase Chain Reaction

265 PFC – Pre-Frontal Cortex

266 PND – Postnatal Day

267 PUCRS – Pontifical Catholic University of Rio Grande do Sul

268 TLRs – Toll-like Receptors

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276 **DECLARATIONS**

277

278 **Ethics approval and consent to participate**

279 The experiments were conducted in accordance with the NIH laboratory animal care guidelines
280 and approved by the Ethical Committee on the Use of Animals of the Pontifical Catholic
281 University of Rio Grande do Sul, Brazil.

282 **Consent for publication**

283 Not applicable.

284 **Availability of data and materials**

285 The datasets used and/or analyzed during the current study are available from the corresponding
286 author on reasonable request.

287 **Competing interests**

288 The authors declare that they have no competing interests.

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292 **Author's contributions**

293 Conceived and designed the experiments: TWV, RGO. Performed the experiments: TWV, RO,
294 KCC, ACS, MSC, MLL, LEWS, RGO. Analyzed the data: TWV. Contributed
295 reagents/materials/analysis tools: RGO. Wrote the paper: TWV, RO, KCC, ACS, MSC, MLL,
296 LEWS, RGO.

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366 **FIGURE TITLES AND LEGENDS**

367

368 Title: Figure 1. Behavioral outcome of CPP protocol in control and stressed animals.

369 Legend: A) AFR group was left undisturbed until weaning, except for routine cage cleaning; MS

370 group was exposed to daily maternal separation from PND2 to PND14 for a period of 180

371 minutes. B) CPP protocol consisted in 3 phases. Habituation: mice had free access to both

372 chambers for 30 minutes. Conditioning: mice were injected alternately with cocaine (striped

373 chamber) and saline solution (dotted chamber) during 10 days; each administration was

374 performed in one chamber. And test: mice were allowed to explore both chambers with no

375 drug/saline administration.

376

377 Title: Figure 2. NFkB1 and NFkB2 mRNA expression in the Hippocampus and Pre-frontal

378 cortex.

379 Legend: A) Significant treatment effect of NFkB1 mRNA levels in the PFC ($p < 0.05$, in a two-

380 way ANOVA). No significant group or interaction effect was detected. B) Significant treatment

381 effect of NFkB2 mRNA levels in the PFC ($p < 0.01$, in a two-way ANOVA). Multiple

382 comparison analyses revealed significant differences between MS-CPP group compared to AFR

383 drug-naïve group (Bonferroni, $p < 0.01$). No significant group or interaction effect was detected.

384 C) No significant group or treatment effect on Nfkb1 mRNA levels in the hippocampus was

385 observed. D) No significant group or treatment effect on Nfkb2 mRNA levels in the

386 hippocampus was detected. AFR drug-naïve $n = 7$, MS drug-naïve $n = 7$, AFR-CPP $n = 7$ —

387 samples collected from the animals with higher CPP scores, MS-CPP $n = 7$ — samples from the

388 animals with higher CPP scores. Data expressed as mean \pm SEM. Single asterisk represents p

389 value below 0.05 in statistical analyses. Double asterisks represent p value below 0.01 in
390 statistical analyses.