

Stress rapidly reorganizes the glutamate synapse in the prefrontal cortex of cocaine-withdrawn adolescent rats

G. Giannotti¹, L. Caffino¹, G. Racagni¹, F. Fumagalli¹

¹Department of Pharmacological and Biomolecular Sciences, University of Milan and Collaborative Center of Department of Antidrug Policies, Presidency of the Council of Ministers

E-mail: giuseppe.giannotti@unimi.it

1. INTRODUCTION

Drug addiction is a major public health issue worldwide characterized by a compulsive drug-seeking and drug-taking behavior. Illicit use of drugs begins and escalates during adolescence, with long-term adverse consequences. The progression from first use to addiction appears to be shorter during adolescence (Clark et al., 1998). The adolescent brain is in a unique state of transition as it undergoes structural and synaptic changes (Crews and Hodge, 2007). The evidence that initiation of drug taking primarily occurs during adolescence suggests a greater addictive potential than at adulthood. The brake against compulsive behaviours is provided by prefrontal cortex that is still developing during adolescence. It is thus possible to hypothesize that if the brake is defective, the chances of risky behaviours become higher. For this reason, adolescence can be considered a crucial period for investigating the development of drug addiction. Recent evidence has implicated, besides dopamine, a role for glutamate at the effects produced by cocaine (Schmidt and Pierce, 2010; Mamei et al., 2011). Using animal models of drug addiction, it has been shown that drug-induced craving, also after withdrawal periods, is accompanied by changes in extracellular dopamine and glutamate concentrations and in glutamatergic receptor expression and phosphorylation in a region- and time-dependent manner (Churchill et al. 1999; Ferrario et al. 2011).

The aim of the present work was 1) to investigate the short-term effects of repeated exposure to cocaine during adolescence on the glutamatergic synapse in prefrontal cortex and 2) to evaluate the dynamic response to a challenging event such as an acute stress as a potential indication of coping ability under a challenging condition.

2. MATERIALS AND METHODS

PROTEIN ANALYSIS

Prefrontal cortex was dissected, frozen on dry ice and stored at -80°C. Tissues were homogenized and membrane fraction was prepared. After being separated on a 10% SDS-polyacrylamide gels, proteins (30 µg) were transferred onto nitrocellulose membrane and blocked 1 hour with 5% milk. Membranes were incubated with antibody against phosphorylated forms of EAAT2 (S1080, S1180, S1191, S1201, S1211, S1221), vGLUT1 (S1080, S1180, S1191, S1201, S1211, S1221) and then detected and revealed with antibody against peroxidase (Dako) (S1080, S1180, S1191, S1201, S1211, S1221). All immunoprecipitates were visualized by chemiluminescence using the ECL Western blotting kit using the Chemidoc MP imaging system (BioRad Laboratories). The abundance of the proteins investigated was expressed as a ratio between the phosphorylated and the respective total forms and analyzed using the Image Lab software from BioRad.

Plasma from each rat was separated by centrifugation and corticosterone was determined by an enzyme-linked immunosorbent assay (ELISA) using a commercial kit according to the manufacturer's instructions (R&D, Wiesbaden, Germany).

STATISTICAL ANALYSIS

Data are presented as mean ± SEM of several independent observations. Changes were analyzed by two-way ANOVA with treatment (saline or cocaine) and sex (male and female) (saline and cocaine groups). Saline and treatment were considered as independent variables. When appropriate, further differences were analyzed by single contrast post hoc test (SNK). An unpaired Student's t test was used to analyze the immobility time, measured during the acute stress. All significant differences were assessed at p < 0.05.

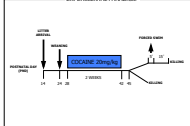
mRNA ANALYSIS

Total RNA was isolated from prefrontal cortex, was treated using RNeasy-quick, the quality of vGLUT1, EAAT2, vGAT1, GAD67, EAAT2, SIRT1, SIRT6, SIRT7, SIRT8, SIRT9, SIRT10 and vGAT1 levels were performed by TaqMan qRT-PCR (Applied Biosystems) real-time system, the β-actin (housekeeping) using the same qRT-PCR kit for primer. All the samples were run in 384-well format to facilitate with a normalizing internal control (NIC). The levels of target genes expression were calculated using a comparative cycle threshold (CT) method.

AIM OF THE WORK



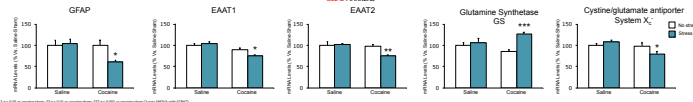
EXPERIMENTAL PARADIGM



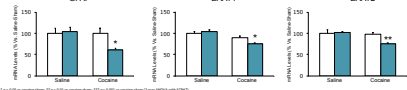
3. RESULTS

GLUTAMATERGIC TRANSMISSION

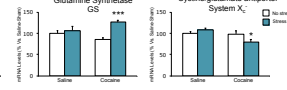
GLIAL MARKERS



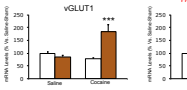
PRE-SYNAPTIC MARKERS



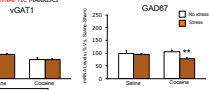
POST-SYNAPTIC MARKERS



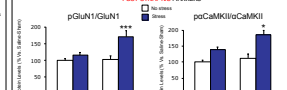
PLASMA CORTICOSTEROID LEVELS



ARC PROTEIN LEVELS



IMMOBILITY TIME



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4. CONCLUSIONS

Stress rapidly reorganizes the glutamate synapse in the prefrontal cortex of cocaine-withdrawn adolescent rats.

Hyper-reactive glutamatergic synapses in mPFC may contribute to explain the hypersensitivity to stress observed in abstinent cocaine users.

Dysregulation of the glutamate homeostasis may contribute to the negative emotional state and stress-induced reinstatement observed in animal models of cocaine abuse.