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Acute Methamphetamine Exposure Inhibits Cardiac Contractile Function

Subat Turdi¹, Robbie M. Schamber^{1,2}, Nathan D. Roe¹, Herbert G. Chew Jr.², Bruce Culver¹, and Jun Ren¹

¹ Center for Cardiovascular Research and Alternative Medicine, University of Wyoming College of Health Sciences, Laramie, WY 82071, USA

² Department of Biology, Western Wyoming College, Rock Springs, WY 82901, USA

Abstract

Methamphetamine, a commonly seen substance of abuse, has been reported to exert detrimental effect on bodily function including the cardiovascular system although its mechanism of action is poorly understood. This study was designed to examine the direct impact of methamphetamine on isolated whole heart and single cardiomyocyte contractile function. Murine hearts and isolated cardiomyocytes from adult FVB mice were exposed to various concentrations of methamphetamine for 30 min prior to the assessment of mechanical function using a Langendorff apparatus and an IonOptix Myocam® system, respectively. Cardiac contractile properties analyzed included maximal velocity of left ventricular pressure development and decline (\pm dP/dt), peak shortening amplitude (PS), maximal velocity of shortening/relengthening (\pm dLdt), time-to-PS (TPS), time-to-90% relengthening (TR₉₀), resting and electrically-stimulated increase of intracellular Ca²⁺ as well as intracellular Ca²⁺ decay. Our results revealed that acute methamphetamine exposure depressed \pm dP/dt, PS and rise of intracellular Ca²⁺ without affecting \pm dLdt, TPS, TR₉₀, resting intracellular Ca²⁺ and intracellular Ca²⁺ decay. Furthermore, methamphetamine nullified the adrenergic agonist norepinephrine-elicited positive cardiomyocyte contractile response, including elevated PS, \pm dLdt and shortened TR₉₀ without affecting TPS. Western blot analysis showed unchanged expression of sarco(endo)plasmic reticulum Ca²⁺-ATPase (SERCA2a) and phospholamban, associated with upregulated Na⁺-Ca²⁺ exchanger levels following acute methamphetamine exposure. In addition, methamphetamine promoted overt cardiomyocyte protein damage evaluated by carbonyl formation. Taken together, these results demonstrate direct cardiac depressant effect of methamphetamine in myocardium and isolated cardiomyocytes, possibly associated with protein damage and dampened adrenergic response.

Keywords

methamphetamine; myocardium; cardiomyocyte; protein damage; adrenergic response

Correspondence to: Dr. Jun Ren, Center for Cardiovascular Research and Alternative Medicine, University of Wyoming College of Health Sciences, Laramie, WY 82071, USA, Tel: (307)-766-6131; Fax: (307)-766-2953; E-mail: jren@uwyo.edu.

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INTRODUCTION

Methamphetamine, a synthetic amine, is derived from amphetamine with profound effect on the central nervous system. It is a potent additive stimulant illegally manufactured, distributed and abused in the United States (NIDA, 1998). It has been usually referred to as “speed”, “crystal”, “crank”, “go”, and “ice”. An estimated US population of 12.3 million are believed to consume the illegal drug somewhere during their life time, representing 5.2% of the American population of age 12 years and older (Office of Applied Studies, 2003). An alarming 54% increase of the emergency department visit was documented between 1995 to 2002 involving use of amphetamines or methamphetamines, with more than half the visits involving young adults aged 18 to 34 (Office of Applied Studies, 2003). The main cardiovascular manifestations of methamphetamine abuse encompass tachycardia, atrioventricular arrhythmias, myocardial ischemia and hypertension (Derlet and Horowitz, 1995; Anglin *et al.*, 2000). Both experimental and clinical examinations especially those from autopsy and individual case reports suggest that methamphetamine exposure is associated with structural and functional changes of hearts, leading to cardiac hypertrophy, disarrangement of myofibers, fibrosis, dilated cardiomyopathy, congestive heart failure and sudden death (Jacobs, 1989; Islam *et al.*, 1995; He *et al.*, 1996; Maruta *et al.*, 1997; Karch *et al.*, 1999; Maeno *et al.*, 2000; Wijetunga *et al.*, 2003). More recently, a case-control study revealed that young methamphetamine users had an alarming 3.7-fold increase in the odds ratio for cardiomyopathy, after adjusting for age, body mass index, and renal failure (Yeo *et al.*, 2007). Nonetheless, the precise mechanism behind methamphetamine exposure-elicited change in cardiac function remains elusive. The aim of the present study was to examine the effect of methamphetamine on cardiac contractile function at the levels of both whole heart and isolated cardiomyocytes. Multicellular cardiac contractile function measured as pre-load recruitable stroke work and dP/dt exhibits significantly decreased myocardial contractile function following methamphetamine treatment (Yu *et al.*, 2002). However, assessment of methamphetamine exposure-induced cardiac electromechanical response in multicellular preparations suffers from the limitation that the heart encompasses a heterogeneous population of cells and may not accurately represent functional changes of the working cardiomyocytes. Thus, the action of methamphetamine on the hearts may be influenced by non-myocyte factors such as interstitial connective tissue or fibrous growth. For example, increased ventricular stiffness may reflect a greater amount of interstitial fibrosis and a shift in collagen content rather than a direct reflection on the mechanical properties of myocytes themselves. To the best of our knowledge, no study has utilized isolated cardiomyocytes to examine the impact of methamphetamine exposure in cardiac contractile function.

MATERIALS AND METHODS

Isolated heart Langendorff study

The experimental procedures described in this study were approved by the institutional animal care and use committee of University of Wyoming (Laramie, WY). In brief, adult male FVB mice (22 – 25 g) were anesthetized with ketamine/xylazine (3:5, 1.32 mg/kg, i.p.). Hearts were removed, the aorta was cannulated, placed on a Langendorff Apparatus (AD Instruments Model ML870B2), and perfused with KHB (95% O₂, 5% CO₂ bubbled; 5 ml/min at 37°C). A micro-balloon tipped catheter was inserted for left ventricular pressure measurements (Verma *et al.*, 2003).

Isolation of murine cardiomyocytes

Single cardiomyocytes were isolated from adult male FVB mice (22 – 25 g) as described (Ren *et al.*, 2008). In brief, hearts were removed and perfused with Krebs-Henseleit bicarbonate (KHB) buffer containing (in mM): 118 NaCl, 4.7 KCl, 1.2 MgSO₄, 1.2 KH₂PO₄, 25

NaHCO₃, 10 HEPES, and 11.1 glucose. Hearts were digested with collagenase D for 20 min. Left ventricles were removed and minced before being filtered. The percentage of the rod-shaped viable cardiomyocytes was ~ 70%.

Myocyte shortening and relengthening

Mechanical properties of cardiomyocytes were assessed by an IonOptix Myocam system (IonOptix Inc., Milton, MA, USA). Cells were placed in a chamber mounted on the stage of an inverted microscope and superfused (at 25°C) with a buffer containing (in mM): 131 NaCl, 4 KCl, 1 CaCl₂, 1 MgCl₂, 10 glucose, 10 HEPES, at pH 7.4. The cells were field stimulated at a frequency of 0.5 Hz. Cell shortening and relengthening were assessed using the following indices: peak shortening (PS), time-to-PS (TPS) and time-to-90% relengthening (TR₉₀), maximal velocities of shortening (+ dLdt) and relengthening (– dLdt) (Ren *et al.*, 2008).

Intracellular Ca²⁺ fluorescence measurement

A cohort of cardiomyocytes was loaded with fura-2/AM (0.5 μM) for 10 min at 25°C, and fluorescence measurements were recorded with a dual-excitation fluorescence photomultiplier system (Ionoptix) as previously described (Ren *et al.*, 2008). Myocytes were imaged through an Olympus IX-70 Fluor oil objective. Cells were exposed to light emitted by a 75W lamp and passed through either a 360 or a 380 nm filter (bandwidths were ± 15 nm), while being stimulated to contract at 0.5 Hz. Fluorescence emissions were detected between 480–520 nm by a photomultiplier tube after first illuminating the cells at 360 nm for 0.5 sec then at 380 nm for the duration of the recording protocol (333 Hz sampling rate). The 360 excitation scan was repeated at the end of the protocol and qualitative changes in intracellular Ca²⁺ concentration was inferred from the ratio of the fura-2 fluorescence intensity (FFI) at the two wavelengths. Fluorescence decay time (single and bi-exponential decay rates) was also measured as an indication of intracellular Ca²⁺ clearing rate.

Western blot analysis

Expression of SERCA2a, Na⁺-Ca²⁺ exchanger (NCX) and phospholamban (PLB) were examined by western blot. Following a 30-min exposure with methamphetamine (1.0 mM), cardiomyocytes were collected and sonicated in a lysis buffer containing 20 mM tris (ph 7.4), 150 mM NaCl, 1 mM EDTA, 1 mM EGTA, 1% triton, 0.1% SDS and protease inhibitor cocktail. The protein concentration of the supernatant was assessed using the protein assay reagent (Bio-Rad Laboratories, Inc, Hercules, CA, USA). The extracted proteins were separated on 10 – 15% SDS-polyacrylamide gels and transferred to polyvinylidene difluoride membranes. After blocking the membrane was incubated with rabbit anti-SERCA2a (1:1000, Affinity Bioreagents Inc., Golden, CO, USA), rabbit anti-NCX polyclonal (1:1000, Swant, Bellinzona, Switzerland) and mouse anti-PLB monoclonal antibody (1:2000, Abcam inc, Cambridge, MA, USA) antibodies overnight at 4°C followed by incubation with second antibodies. The antigens were detected by the luminescence method. Quantification of band density was determined with the Quantity One software (Bio-Rad, version 4.4.0, build 36) and reported in optical density per square millimeter (Ren *et al.*, 2008).

Protein carbonyl assay

Following methamphetamine treatment (1.0 mM, 30 min), proteins were extracted from cardiomyocytes. Nucleic acids were eliminated by treating the samples with 1% streptomycin sulphate for 15 min, followed by a 10 min centrifugation (11,000×g). Protein was precipitated by adding an equal volume of 20% trichloroacetic acid (TCA) to protein (0.5 mg) and centrifuged for 1 min. The TCA solution was removed and the sample resuspended in 10 mM 2, 4-dinitrophenylhydrazine (2,4-DNPH) solution. Samples were incubated at room temperature for 15–30 min. After adding 500 μl of 20% TCA, samples were centrifuged for 3

min. The supernatant was discarded, the pellet washed in ethanol: ethyl acetate and allowed to incubate at room temperature for 10 min. The samples were centrifuged again for 3 min and the ethanol: ethyl acetate steps repeated twice or more times. The precipitate was resuspended in 6 M guanidine solution, centrifuged for 3 min and any insoluble debris removed. The maximum absorbance (360–390 nm) of the supernatant was read against appropriate blanks (water, 2 M HCl) and the carbonyl content was calculated using the molar absorption coefficient of $22,000 \text{ M}^{-1} \text{ cm}^{-1}$ (Ren *et al.*, 2008).

Data analysis

Data were expressed as Mean \pm SEM. Statistical significance ($p < 0.05$) for each variable was estimated by analysis of variance (ANOVA) followed by Tukey's test for *post hoc* analysis.

RESULTS

Effect of methamphetamine on isolated heart function and cardiomyocyte contractile function

Acute exposure of methamphetamine for 30 min did not affect the heart rate although it significantly inhibited the maximal velocity of left ventricular pressure development (+ dP/dt) and decline (– dP/dt). At the end of a 30-min exposure, methamphetamine inhibited + dP/dt by ~30% and 37% at 0.1 mM and 1.0 mM, respectively. Similarly, methamphetamine inhibited – dP/dt by ~20% and 21% at 0.1 mM and 1.0 mM, respectively. The threshold of methamphetamine-induced suppression on heart function was between 0.01 mM and 0.1 mM (Fig. 1). The methamphetamine-induced inhibition in \pm dP/dt was maximal within 15 min of exposure, maintainable up to 60 min and was reversible upon min washout (data not shown). The inhibitory effect of methamphetamine on isolated heart contractility was associated with reduced peak shortening in freshly isolated cardiomyocytes. However, the threshold required for the methamphetamine-elicited inhibition on cardiomyocyte contractile capacity was shifted to a higher concentration range (between 0.1 mM and 1.0 mM). The resting cell length, maximal velocity of shortening and relengthening (\pm dLdt), duration of shortening (TPS) and relengthening (TR_{90}) were unaffected in cardiomyocytes by methamphetamine at the concentration range tested (Fig. 2).

Effect of methamphetamine on norepinephrine-elicited cardiomyocyte contractile response

Methamphetamine is known to interfere with cardiac adrenergic responsiveness (Urabe, 1982), although cardiac norepinephrine content may not be depleted following chronic methamphetamine administration (Ruffoli *et al.*, 2008). To examine the impact of acute methamphetamine exposure on cardiomyocyte adrenergic response, isolated cardiomyocytes were incubated with the adrenergic agonist norepinephrine (1 μM) in the absence or presence of methamphetamine (0.1 mM) for 30 min before mechanical function was assessed. Although this concentration of methamphetamine did not elicit any overt effect on cardiomyocyte contractile function, it effectively blunted norepinephrine-elicited increase in peak shortening (PS) as well as maximal velocities of shortening and relengthening (\pm dL/dt). Furthermore, norepinephrine elicited shortened cardiomyocyte relengthening duration (TR_{90}), which was also abrogated by methamphetamine. The duration of shortening (TPS) was not affected by either norepinephrine or methamphetamine (Fig. 3).

Effect of methamphetamine on intracellular Ca^{2+} transients

To explore the possible mechanism underlying methamphetamine-induced cardiomyocyte mechanical dysfunction, we used the membrane permeable intracellular Ca^{2+} fluorescent dye fura-2 to evaluate intracellular Ca^{2+} homeostasis in cardiomyocytes. Our results shown in Fig. 4 depicted comparable resting intracellular Ca^{2+} levels and intracellular Ca^{2+} transient decay

rates (either single or bi-exponential) associated with a decrease in electrically-stimulated rise in intracellular Ca^{2+} in cardiomyocytes following methamphetamine (1.0 mM) exposure. These data indicated existence of intracellular Ca^{2+} handling defect in cardiomyocytes following methamphetamine exposure.

Protein expression of intracellular Ca^{2+} regulatory proteins and carbonyl formation

To better understand the mechanism of action responsible for methamphetamine-induced intracellular Ca^{2+} dysregulation, Western blot analysis was performed to assess the expression of the key Ca^{2+} regulatory proteins. Our data revealed that acute methamphetamine (1.0 mM) exposure unregulated the expression of NCX without affecting that of SERCA2a and phospholamban. Our data further revealed that significantly elevated formation of protein carbonyl in myocytes following methamphetamine exposure (Fig. 5).

DISCUSSION

Our study provided evidence for the first time that the physically and psychologically addictive substance methamphetamine directly depresses cardiac contractile function, intracellular Ca^{2+} handling and adrenergic response in isolated murine cardiomyocytes, in parallel to its cardiac depressant action on the whole heart Langendorff apparatus setting. The methamphetamine-induced cardiac depression may involve enhanced protein damage evidenced by enhanced carbonyl formation. Intracellular Ca^{2+} homeostasis was disturbed by acute methamphetamine exposure supported by dampened intracellular Ca^{2+} release in response to electrical stimuli and upregulated NCX but not SERCA2a and phospholamban expression. These results favor a direct detrimental effect of methamphetamine on cardiomyocyte mechanical function, which may contribute to the methamphetamine-induced impairment of myocardial contractile function following exposure of the abused substance.

Although the mechanism(s) of action underscoring reduced cardiomyocyte contraction in response to acute methamphetamine exposure is not fully clear at this time, several speculations may be considered. First, the observation that methamphetamine abolished norepinephrine-induced positive cardiac contractile response suggests potential involvement of adrenergic response in methamphetamine-elicited inhibitory effect in the heart. This is consistent with the previous finding that methamphetamine directly interferes with cardiac adrenergic responsiveness (Urabe, 1982). Methamphetamine exposure was reported to promote catecholamine including norepinephrine release (Yu *et al.*, 2002), the later may directly stimulate cardiac adrenergic receptor, improved cardiac contractility (albeit transiently), cardiac remodeling leading to cardiac hypertrophy and fibrosis. Excessive stimulation of catecholamine release in response to methamphetamine exposure may quickly exhaust the catecholamine storage (Ruffoli *et al.*, 2008), leading to a reduced adrenergic response. Catecholamine desensitization is a hallmark of heart failure (Brodde *et al.*, 2006). It has been shown that heart failure may trigger classical catecholamine desensitization to both isoproterenol and norepinephrine in the presence and absence of ganglionic blockade (Brodde *et al.*, 2006). Data from our current study is in line with the desensitized adrenergic response and cardiac contractile function, supporting a role of adrenergic desensitization in methamphetamine-elicited heart anomalies and ultimately heart failure. Nonetheless, it should be mentioned that direct action of methamphetamine on the cardiac adrenergic signaling cascade (such as at the levels of membrane receptor, G protein and G protein coupled effectors) is essentially unknown. Secondly, the observation that methamphetamine inhibits the electrically-stimulated intracellular Ca^{2+} release and upregulates Na^+ - Ca^{2+} exchanger depicts that methamphetamine may interfere with intracellular Ca^{2+} homeostasis. The reduced intracellular Ca^{2+} release in response to electrical stimuli is consistent with the compromised myocardial contractility (\pm dP/dt and PS) following acute exposure of methamphetamine.

Expression of SERCA2a and phospholamban was unchanged following methamphetamine exposure, not favoring a role of the sarco(endo)plasmic Ca^{2+} -ATPase (Ca^{2+} pump) in methamphetamine-induced cardiomyocyte mechanical response. This observation of unchanged SERCA2a/phospholamban expression is in line with the unaffected duration of shortening and relengthening (TPS and TR_{90}) following acute methamphetamine exposure. Upregulated NCX expression/activity has been shown to be associated with compromised myocardial contractile function due to its possible role in reduced cytosolic Ca^{2+} ion (Goldhaber *et al.*, 2005). Homozygous overexpression of NCX has been shown to lead to modest hypertrophy at baseline and are more susceptible to cardiac failure during the stress of pregnancy or aortic banding (Goldhaber *et al.*, 2005).

Data from our current study revealed enhanced protein carbonyl formation, indicative of protein damage, in cardiomyocytes following methamphetamine exposure. Although exactly how methamphetamine induces protein damage warrants further study, methamphetamine has been shown to promote apoptosis and accumulation of reactive oxygen species. Methamphetamine is known to trigger neuronal, splenic and myocardial damage via apoptosis (Iwasa *et al.*, 1996; Davidson *et al.*, 2001). In addition, methamphetamine may facilitate the formation of reactive oxygen species including ONOO^- , which in turn can cause tissue and cell damage (Wang *et al.*, 2001). Although it is beyond the scope of the current study, methamphetamine has been demonstrated to promote p53 expression and necrosis (Imam *et al.*, 2001). These apoptotic and necrotic as well as pro-oxidation properties of methamphetamine are consistent with the detrimental effect of methamphetamine on protein integrity.

Measurement of contractile performance in isolated cardiomyocytes has been established to provide a fundamental assessment of cardiac contractile function in pathological states, in this case acute methamphetamine exposure. However, as in any study of this nature, caution needs to be taken when correlating the cellular findings to whole heart function, as the latter is composed of heterogeneous cell types, including nerve terminals and fibroblasts. These heterogeneous cell types may have contributed to the subtle discrepancies between our cardiomyocyte and Langendorff heart studies (e.g., in the threshold of methamphetamine).

In conclusion, our study provides the first laboratory evidence at the level of cardiomyocytes to confirm the clinical findings that methamphetamine use is linked to a distinctive form of dilated cardiomyopathy (Jacobs, 1989; Wijetunga *et al.*, 2003; Yeo *et al.*, 2007). Our results demonstrated that direct cardiomyocyte contractile depression by methamphetamine, possibly through protein damage and intracellular Ca^{2+} dysregulation. These findings not only reveal an essential role of the basic myocardial working element, cardiomyocytes, in methamphetamine-induced cardiac anomalies but also provide new insights for better therapeutic remedy against methamphetamine-induced global cardiac pathology. Nonetheless, the precise nature behind methamphetamine-induced depression on cardiomyocyte contractile function is still unclear. Awaiting future studies include cardiac excitation-contraction coupling and membrane ion channels in response to methamphetamine exposure. These approaches will be crucial to further our understanding of the pathological profiles of methamphetamine.

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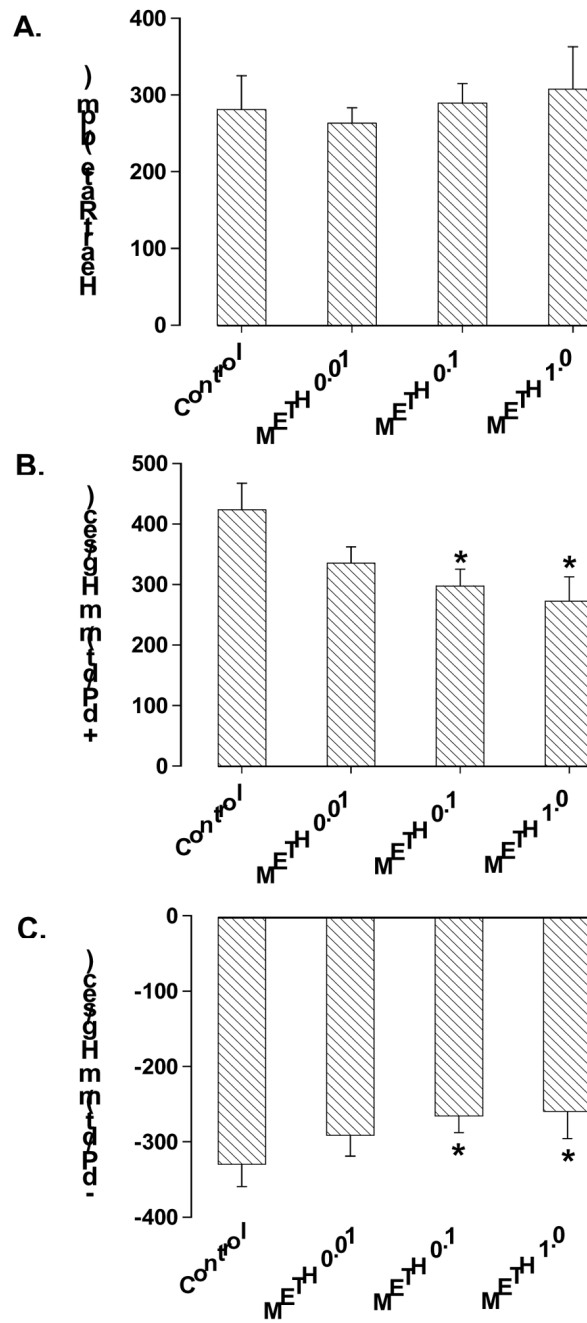


Fig. 1. Influence of acute methamphetamine (METH, 0.01 – 1.0 mM for 30 min) exposure on heart rate (panel A), maximal velocity of left ventricular pressure development (+ dP/dt, panel B) and decline (- dP/dt, panel C) in isolated mouse hearts. Mean \pm SEM, n = 6, * p < 0.05 vs. control value.

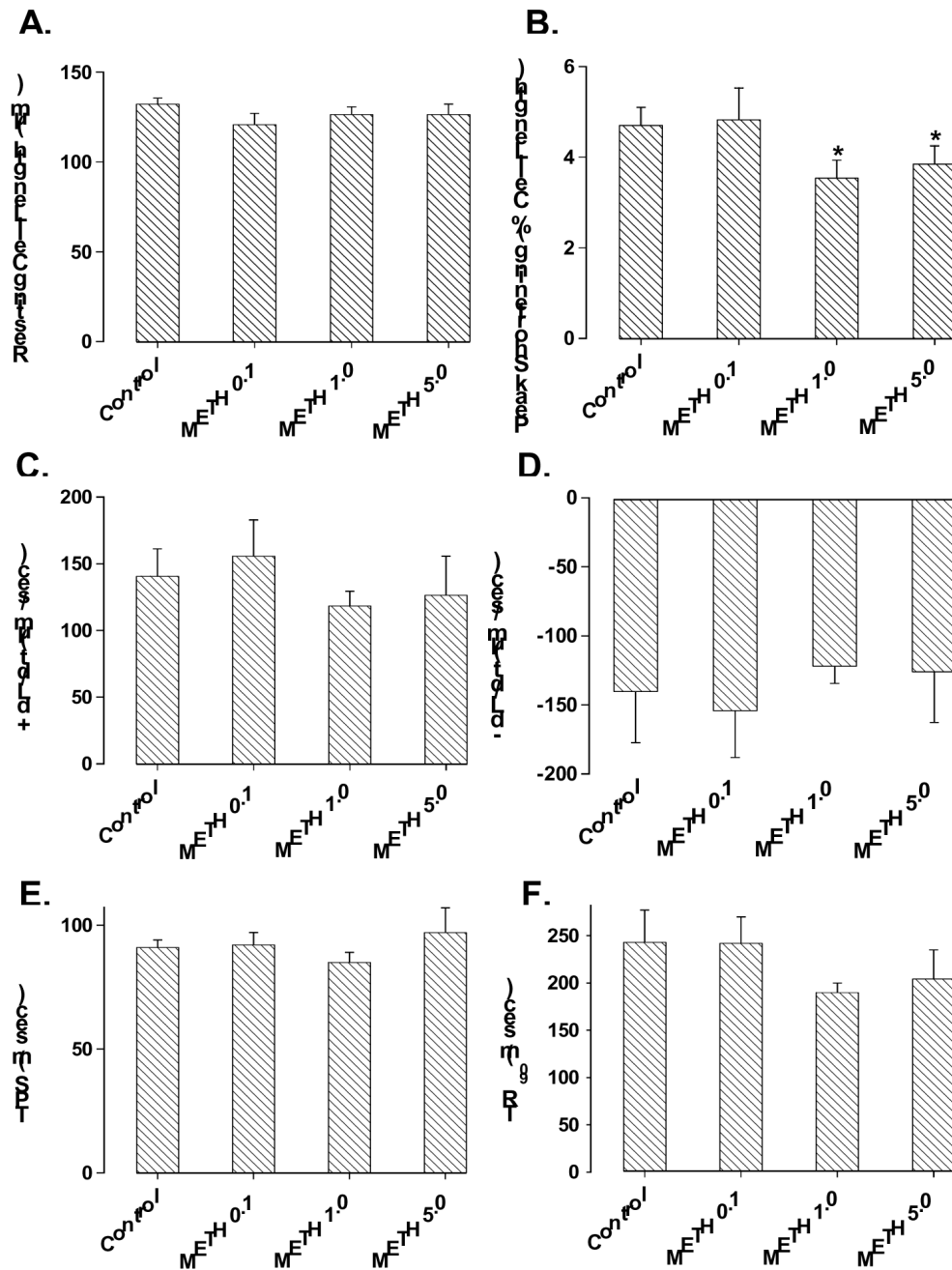


Fig. 2. Murine cardiomyocyte contractile function in response to acute methamphetamine (METH, 0.1 – 5.0 mM for 30 min) exposure. Panel A: Resting cell length; Panel B: peak shortening (% of cell length); Panel C: Maximal velocity of shortening (+ dL/dt); Panel D: Maximal velocity of relengthening (– dL/dt); Panel E: Time-to-PS (TPS); and Panel F: Time-to-90% relengthening (TR₉₀). Mean ± SEM, n = 28 –35 cells per group, * p < 0.05 vs. control value.

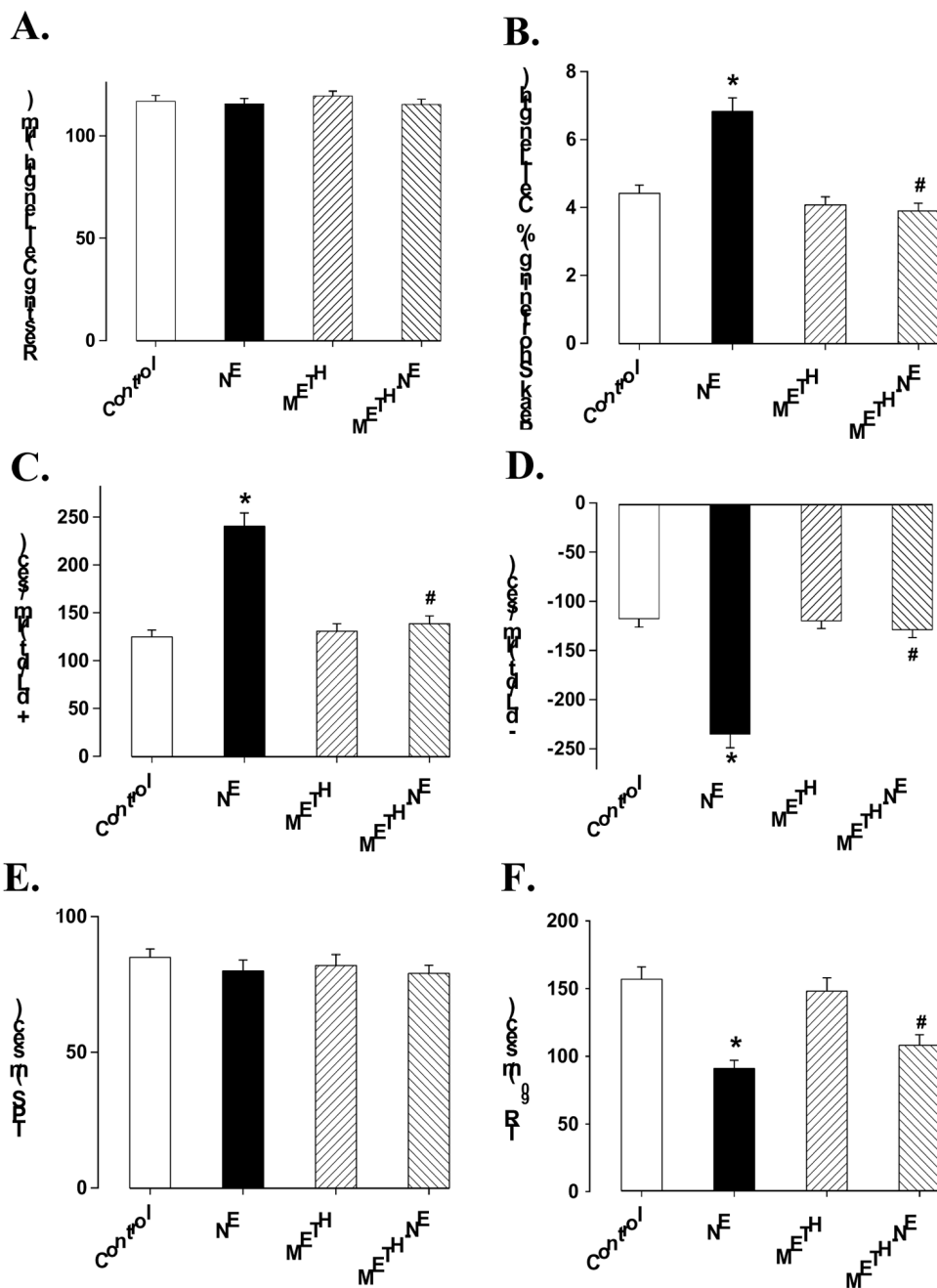


Fig. 3. Influence of methamphetamine (METH) on norepinephrine (NE)-induced cardiomyocyte contractile response. Isolated murine cardiomyocytes were incubated with NE (1 μ M), METH (0.1 mM) or both for 30 min prior to assessment of mechanical function. Panel A: Resting cell length; Panel B: peak shortening (% of cell length); Panel C: Maximal velocity of shortening (+ dL/dt); Panel D: Maximal velocity of relengthening (- dL/dt); Panel E: Time-to-PS (TPS); and Panel F: Time-to-90% relengthening (TR₉₀). Mean \pm SEM, n = 107 cells per group, * p < 0.05 vs. control group, # p < 0.05 vs. NE group.

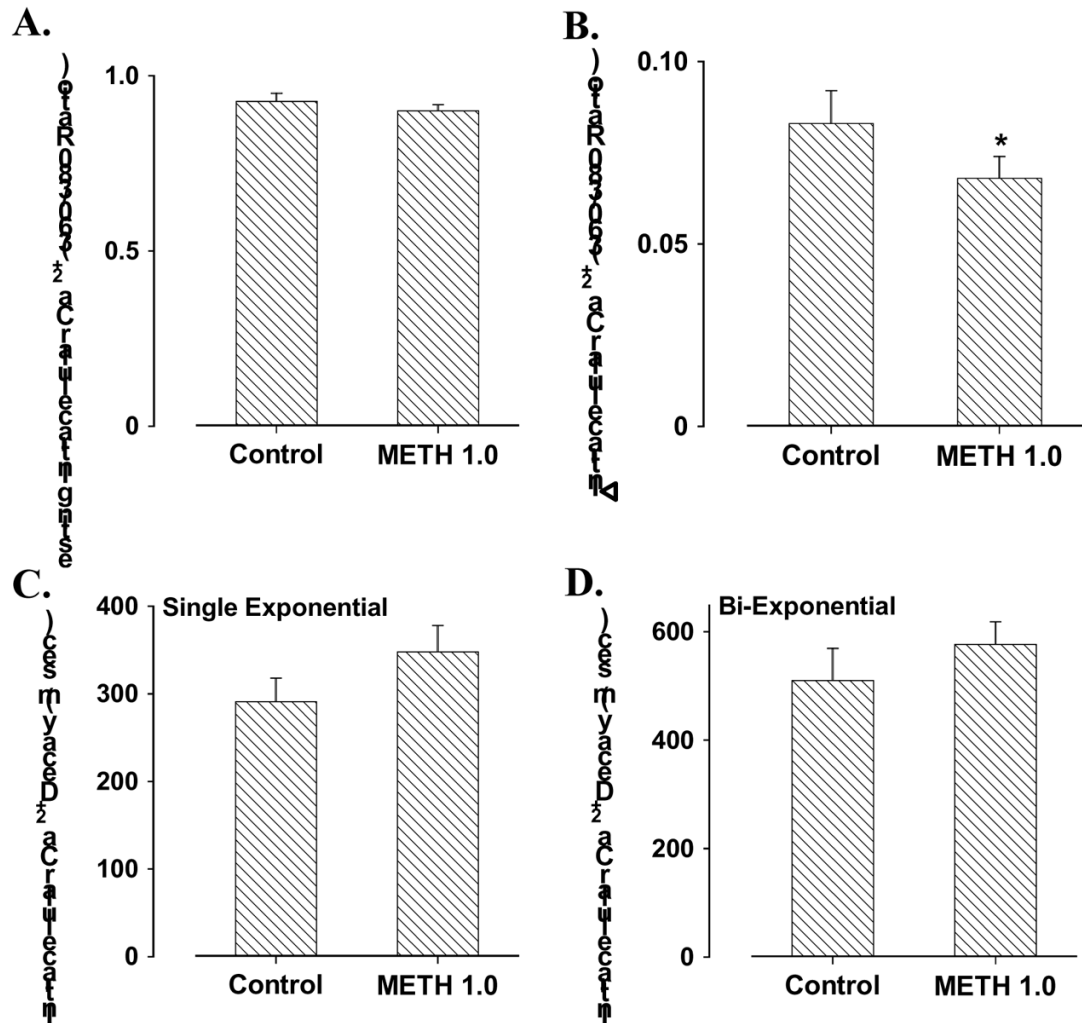


Fig. 4. Murine cardiomyocyte intracellular Ca²⁺ property in response to acute methamphetamine (METH, 1.0 mM for 30 min) exposure. Panel A: Resting intracellular Ca²⁺ levels; Panel B: electrically-stimulated increase in intracellular Ca²⁺ levels; Panel C: Intracellular Ca²⁺ transient decay rate (single exponential); and Panel D: Intracellular Ca²⁺ transient decay rate (Bi-exponential). Mean ± SEM, n = 47–50 cells per group, * p < 0.05 vs. control value.

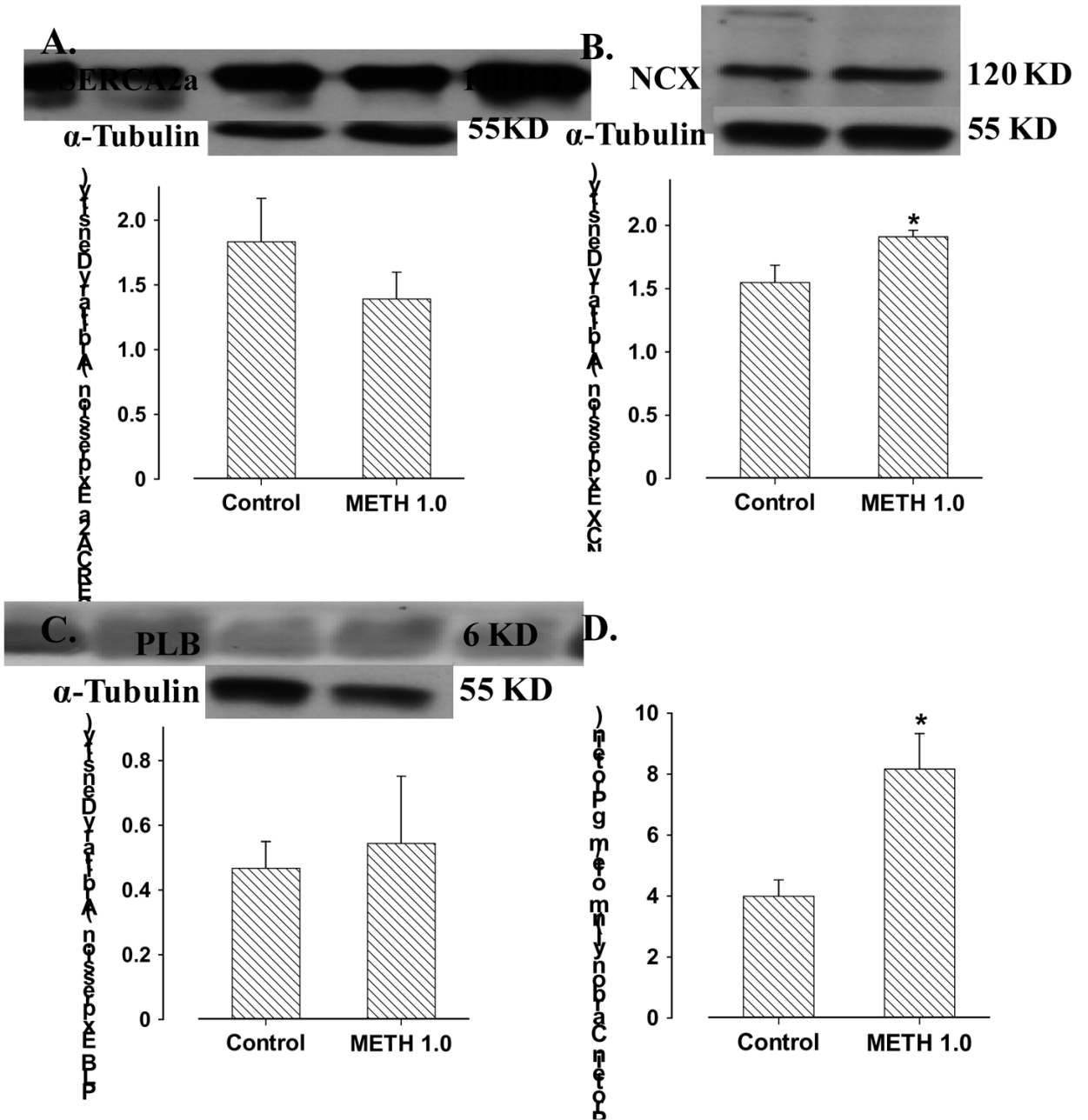


Fig. 5. Levels of cardiomyocyte Ca^{2+} regulatory proteins and myocardial protein carbonyl formation in response to acute methamphetamine (METH, 1.0 mM for 30 min) exposure. Panel A: SERCA2a; Panel B: Na^{+} - Ca^{2+} exchanger (NCX); Panel C: Phospholamban (PLB); and Panel D: Protein carbonyl formation. Inset: Representative gel blots depicting the expression of SERCA2a, NCX and PLB using specific antibodies. Mean \pm SEM, n = 6 – 7 cell isolation or myocardium per group, * p < 0.05 vs. control value.