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***“MOLECULAR MECHANISMS OF NCX1 AND  
NCX3-INDUCED NEUROPROTECTION IN  
ANOXIC BRAIN PRECONDITIONING”***

**TUTORS:**

Prof. Gianfranco Di Renzo

Prof. Antonella Scorziello

**PhD STUDENT:**

Dr. Maria Josè Sisalli

**COORDINATOR:**

Prof. Lucio Annunziato

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## INTRODUCTION

The term *ischemia* (Greek *iskhein* to keep back + *haema* or *hema* blood) means a restriction in blood supply to a bodily organ or tissues. Interruption of blood flow to the brain, or part(s) of the brain, is known as cerebral ischemia or stroke. There are two different kinds of stroke: haemorrhagic or ischemic. Hemorrhagic stroke is due to the rupture of a cerebral artery because of the increase of blood pressure; ischemic stroke, that is far more common, is due to a deficit of cerebral blood flow.

The stroke is a medical emergency, which can cause permanent neurological damage, complications, and death. Stroke causes 9% of all deaths around the world, is the third most common cause of death after ischemic heart disease and cancer, and may soon become the leading cause of death worldwide. It is the leading cause of adult disability, since 76% of people in the United States and Europe survive their stroke (American Heart Association; 2008).

Representing stroke one of the three leading causes of death and the major cause of long-term disability makes it and its therapy, one of the major challenges facing contemporary medicine. In order to reduce the consequences of this disease, and to develop therapeutic strategies aimed to reduce neuronal damage, the medical research has to conduct experiments to highlight the cellular, biochemical and molecular mechanisms involved in

neuronal damage. Nowadays, there are no effective therapeutic protocols able to induce significant improvement of ischemia-induced brain damage, and all clinical trials conducted in the eighties and nineties were ineffective in the clinical treatment of stroke. The only FDA-approved treatment for acute cerebral ischemia is rTPA, which has demonstrated to cause a improvement in long-term outcomes. However, only a small percentage of patients with ischemic stroke are treated with rTPA due to its narrow therapeutic window and contraindications to thrombolytic therapy.

Thrombolytic therapy does not directly interfere with the ischemic cascade, but changes the cellular and molecular environment that induces it. Neuronal damage often continues with restored blood flow to areas. New therapies based on the understanding of the complex interactions of the ischemic cascade may eventually increase the therapeutic window for salvage of the penumbra and provide direct neuroprotection. Combination treatment with a neuroprotective drug or multiple drugs and thrombolytic therapy might theoretically improve stroke outcomes, but this hypothesis requires confirmation in clinical trials. Potentially, patients at risk of ischemic stroke could receive prophylactic treatment with a neuroprotective drug, either extending the window for thrombolytic efficacy or decreasing the damaged area. The prevention of oxidative damage with novel free radical trapping agents may be effective in improving outcome by reducing disability caused by ischemic stroke. Such agents may be efficacious both as a stand-alone

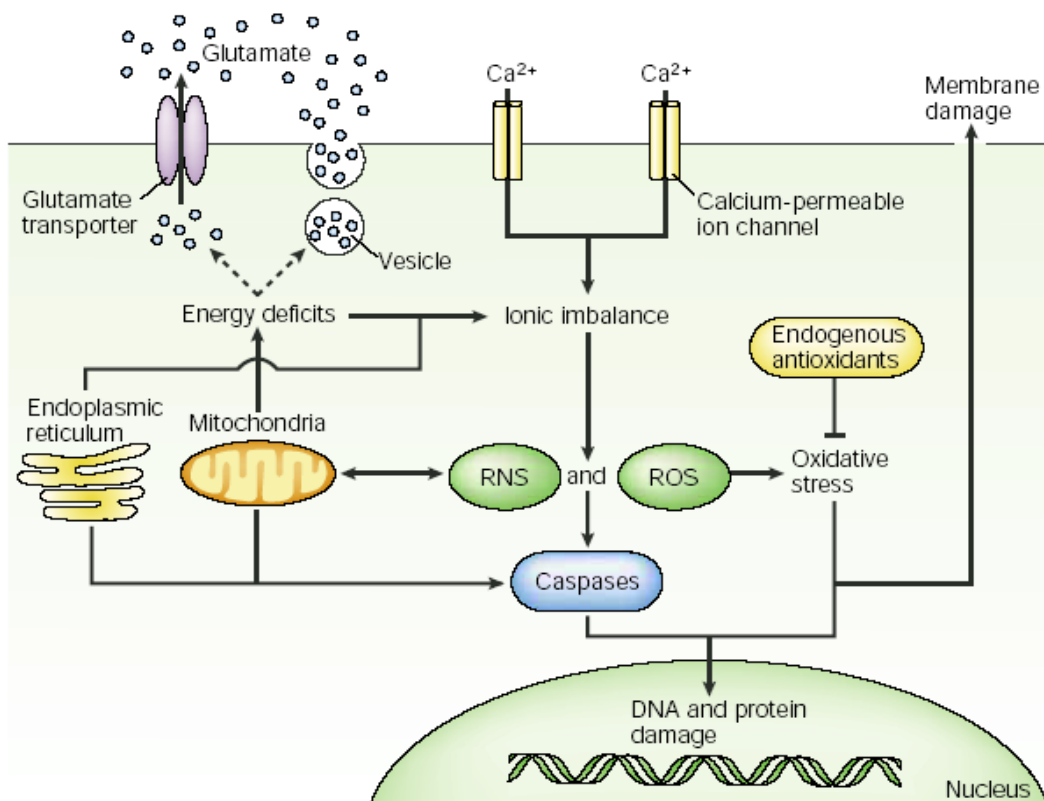
treatment in view of the efficacy seen in models of permanent ischemia in several species, as well as when used as adjuncts to thrombolytic therapy. A combination treatment approach may, in addition to reducing the damage due to ischemia, also reduce reperfusion injury, which may be caused by the deleterious effects of oxygen radical species produced when oxygenation is restored to the ischemic brain (Weinberger, 2006).

## ***PHYSIOPATHOLOGY OF ISCHEMIC DAMAGE***

The ischemic process is due to a complex series of pathophysiological events triggered by a decreased supply of oxygen and glucose in the cerebral tissue which requires, for a smooth running, large amounts of these two molecules. The cerebral energy production, given the scarcity of glucose deposits and other macromolecules available for anaerobic pathways of ATP production, depends almost entirely on the good operating of the mitochondrial oxidative chain.

The main pathogenetic mechanism involved in the genesis of ischemic injury seems to be excitotoxicity, peri-infarct depolarization, oxidative stress, inflammation, intracellular ionic dysregulation. In fact, altered  $\text{Ca}^{2+}$  dysregulation, triggering  $\text{Ca}^{2+}$ -dependent bio-polymer degradation and mitochondrial and bioenergetic failure, ultimately culminates in the activation of reactions leading to necrotic/apoptotic cell death. The ischemic cell death is due to changes directly related to the inhibition of oxidative phosphorylation. These modifications include pH reduction, ATP decrease, production of free radicals from the mitochondrial chain, increase in  $[\text{Na}^+]_i$ . The energetic depletion determines the block of  $\text{Na}^+/\text{K}^+$  ATPase and  $\text{Ca}^{2+}$  ATPase lying to a reduction of neuronal and glial membrane potential (Katsura et al., 1994) (Figure 1).





**Figure 1: Physiopathology of ischemic damage**

This kind of depolarization, called “anoxic depolarization” as it occurs some minutes after the stimulus anoxic/ischemic, determines both the activation of voltage gated  $Ca^{2+}$  channels (VOCC) and an inversion in the operation mode of the  $Na^+$  depending transporters, such as the  $Na^+/Ca^{2+}$  exchanger activated in *reverse mode* of operation. These modifications determine a massive entrance of  $Ca^{2+}$  into the cell with a consequent increase of its concentration, critical for the genesis and perpetuation of the ischemic damage. Indeed, this ion is able to activate enzyme and metabolic pathways potentially

detrimental, to induce the release of neurotransmitter such as glutamate, and to inhibit oxidative mitochondrial chain.

Glutamate is the most common excitatory neurotransmitter in the central nervous system and many functions of the brain occur through the release of this mediator in the synapses. Therefore, an alteration of glutamatergic neurotransmission seems to be responsible for different diseases, including them the ischemic neuronal death. In neurons, glutamate is stored in synaptic vesicles by a vesicular transporter which uses, as energy source, a  $H^+$  gradient created by the vesicular proton pump. This excitatory neurotransmitter is released from the presynaptic terminal after the  $Ca^{2+}$  entrance and acts both on presynaptic and postsynaptic receptors, which are ionotropic as AMPA, NMDA e kainate, and metabotropic.

The glutamate extracellular concentrations briskly rise during acute brain injury since its presynaptic  $Na^+$  dependent transporter is blocked by the anoxic depolarization and consequent rising of  $[Na^+]_i$  induced by ischemic insult.

The metabotropic glutamate receptors are coupled to a  $G_q$  protein and their stimulation activate phospholipase-C that, through the production of inositol triphosphate (IP3), mobilizes calcium from intracellular stores contributing to further increasing on its concentrations. Therefore, there is a continuous "*release to release*" of glutamate, followed by entry of  $Ca^{2+}$  and  $Na^+$  into the cell and consequent damage and cell death.

These findings have led to the elaboration of the paradigm of glutamate excitotoxicity that explained ischemic neuronal cell death as a mere consequence of  $\text{Na}^+$  and  $\text{Ca}^{2+}$  influx through glutamate receptors. Although this theory has been guiding basic research in the field of neurodegeneration for almost 3 decades, more recently it has become the object of serious criticism and reassessment. What has aroused such skepticism among researchers has been the fact that although first, second, and third generation glutamate receptor antagonists have long yielded promising results in animal models of brain ischemia, they have failed to elicit a neuroprotective action in stroke and traumatic brain injury in humans. Therefore, the theory of excitotoxicity, though a fascinating paradigm, can only explain some of the events occurring in the acute phase of anoxic insult but cannot be seen as a major target for developing new therapeutic avenues for brain ischemia.

Increasing evidence suggests that there may exist mechanisms of neurotoxicity that operate independently of, or in parallel with, excitotoxicity. Indeed, during ischemia, neuronal calcium channels and transporters (including NCX, TRM2, TRPM7, ASICS, CaV1.2 and hemichannels) as well as glutamate receptors (NMDAR, AMPAR, KAR) are overactivated. The increased activity of plasma membrane  $\text{Ca}^{2+}$  channels can then trigger the entry of  $\text{Ca}^{2+}$  into the cytosol of neurons, leading to larger than usual increases in the cytosolic calcium concentration.  $\text{Ca}^{2+}$  ions may gain access

to the neuronal cytoplasm via ion channels or  $\text{Ca}^{2+}$  transport systems, or through the release of  $\text{Ca}^{2+}$  ions from intracellular stores (Figure 2) (Szydłowska et al., 2009).

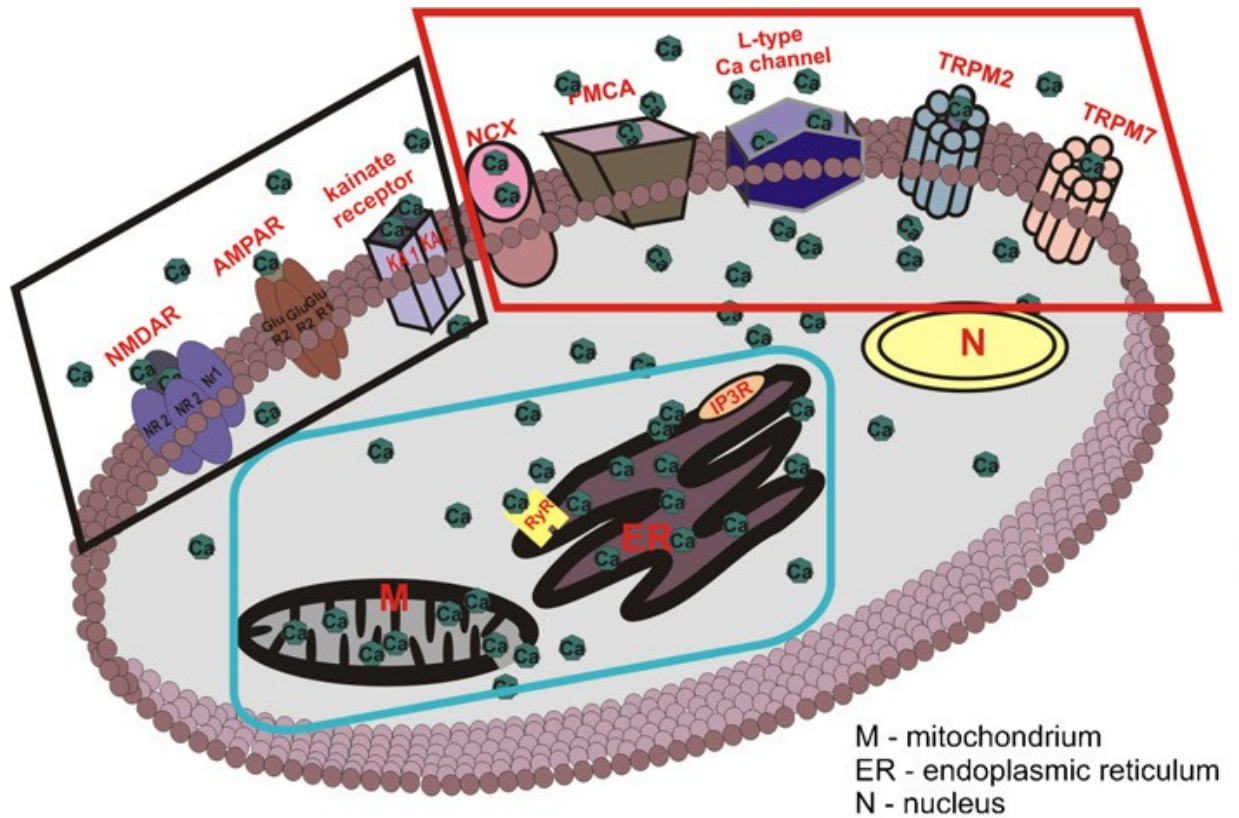


Figure 2: Routes of calcium entry into neurons after ischemic injury

Depletion of calcium ions from the endoplasmic reticulum (ER) has been suggested as an initial signal for ER dysfunction in ischemic neurons (Paschen et al., 1999). Many studies indicate that a strong release of calcium ions from ER is associated with damage to cells, including damage to neurons after ischemia (Pisani et al., 2000). Chen et al. reported that dysregulation of  $\text{Ca}^{2+}$ ER homeostasis following ischemia involves two phases: accumulation of  $\text{Ca}^{2+}$  in ER stores and subsequent release of  $\text{Ca}^{2+}$  from ER following ischemia/reoxygenation (REOX) (Chen et al., 2008). NKCC1 and NCXrev are involved in  $\text{Ca}^{2+}$  ER dysregulation, which includes overload of  $\text{Ca}^{2+}$  into ER during oxygen and glucose deprivation (OGD) and release of  $\text{Ca}^{2+}$  from ER during REOX (Chen et al., 2008).  $\text{Ca}^{2+}$  ER release is largely mediated by IP3R activation during early REOX (Bruno et al., 2001). Inhibition of NKCC1 during OGD prevented  $\text{Ca}^{2+}$  ER accumulation. This occurs because NKCC1-mediated  $\text{Na}^+$  flux may trigger  $\text{Ca}^{2+}$  influx through NCXrev and contributes to  $\text{Ca}^{2+}$  ER loading (Chen et al., 2008). It has been suggested that IP3R-mediated ER  $\text{Ca}^{2+}$  release can enter the adjacent mitochondria and trigger cytochrome c release (Rizzuto et al., 2006).

Consistent with an elevation of NCX activity and in accordance with normal cytosolic calcium concentrations, Sirabella et al. found that during OGD an increased refilling of  $\text{Ca}^{2+}$  into ER occurred (Sirabella et al., 2009). NCX1 upregulation occurred following OGD and may play a fundamental role in ER  $\text{Ca}^{2+}$  refilling thus preventing ER stress and cell death during OGD.

As well as endoplasmic reticulum, mitochondria sense and shape cytosolic  $\text{Ca}^{2+}$  signals by taking up and subsequently releasing  $\text{Ca}^{2+}$  ions during physiological and pathological  $\text{Ca}^{2+}$  elevations.

Mitochondria contain two membranes, an outer membrane permeable to solutes and an inner membrane impermeable to solutes that harbours the respiratory chain complexes. The respiratory chain pumps protons against their concentration gradient from the matrix of the mitochondrion into the inter-membrane space, generating an electrochemical gradient in the form of a negative inner membrane potential and of a pH gradient, the matrix being more alkaline than the cytosol (Bernardi, 1999; Poburko et al., 2011). The electrical and chemical components of the proton-motive force add up to energize the back-flux of protons down their electrochemical gradient across the ATP synthase, the enzyme that generates ATP. The negative mitochondrial membrane potential ( $\Delta\Psi_m$ ) used to drive the entry of protons also favors the entry of calcium, a divalent cation, into the mitochondrial matrix. As a result, mitochondria can accumulate large amounts of calcium through a  $\text{Ca}^{2+}$ -selective channel known as the mitochondrial  $\text{Ca}^{2+}$  uniporter (MCU) (Kirichok et al., 2004; Nicholls et al., 2000). The MCU has a relatively low  $\text{Ca}^{2+}$  affinity,  $K_d \sim 10$  mM in permeabilized cells (Bernardi, 1999), but  $\text{Ca}^{2+}$  uptake can be readily detected in intact cells because a significant fraction of mitochondria are located close to calcium release or calcium entry channels and therefore exposed to microdomains of high calcium concentrations

(Giacomello et al., 2010; Rizzuto et al., 1993; Pacher et al., 2002). Electrophysiological recordings of mitoplasts, small vesicles of inner mitochondrial membrane, revealed that the MCU is a highly  $\text{Ca}^{2+}$ -selective inward-rectifying ion channel (Kirichok et al., 2004).

Instead,  $\text{Ca}^{2+}$  efflux from mitochondria is catalyzed by antiporters that drive  $\text{Ca}^{2+}$  out of the mitochondrial matrix in exchange with  $\text{Na}^+$  or  $\text{H}^+$  (Nicholls and Crompton, 1980). Two types of exchangers have been functionally characterized in the 1970s, the  $\text{Na}^+/\text{Ca}^{2+}$  and the  $\text{H}^+/\text{Ca}^{2+}$  exchangers; the former is particularly abundant in excitable tissues (Carafoli, 2003). These two pathways have been defined as  $\text{Na}^+$ -independent pathway for  $\text{Ca}^{2+}$  efflux ("NICE") and  $\text{Na}^+$ -dependent pathway for  $\text{Ca}^{2+}$  efflux ("NCE") respectively. They have different kinetics of activation and calcium affinity (Harris, 1979; Lehninger et al., 1978; Ramachandran and Bygrave, 1978). Another mechanism responsible for mitochondrial calcium efflux is represented by mPTP.

Although the  $\text{Na}^+$ -dependent  $\text{Ca}^{2+}$  efflux is characteristic of excitable cells, recent observations indicate that the mitochondrial  $\text{Na}^+-\text{Ca}^{2+}$  exchanger (mNCX) plays an important role in the control of  $[\text{Ca}^{2+}]_c$  also in non-excitable cells, since the mNCX inhibitor CGP37157 blunts oscillations induced by G-protein coupled agonists (Hernández-SanMiguel 2006). Located in the inner mitochondrial membrane, the mNCX mediates the efflux of  $\text{Ca}^{2+}$  from the mitochondria coupled to the influx of  $\text{Na}^+$ , with a

stoichiometry of  $3\text{Na}^+/2\text{Ca}^{2+}$  (Pfeiffer et al., 2001; Saris and Carafoli, 2005). This implies that the exchanger is electrogenic because it transports an excess of positive charge into the matrix. In respiring mitochondria, the relative membrane depolarization, caused by the activity of the exchanger, is compensated by proton pumping through the respiratory chain. As for most ion transporters of the inner mitochondrial membrane, identity of these proteins has been long searched. Only recently has the NCLX exchanger, a member of the NCX family, been hypothesized to be expressed and localized in the inner membrane of mitochondria (IMM) (Palty et al., 2010). This  $\text{Li}^+$  sensitive protein localized within the cristae, is both phylogenetically and functionally distinct from NCX and NCKX family members (Palty et al., 2010). These authors also showed that NCLX participates to the mitochondrial  $\text{Na}^+/\text{Ca}^{2+}$  exchanger activity. However, the molecular mechanisms involved in NCLX expression, transport, localization, as well as its role in mitochondrial activity regulation still remain unidentified. On the other hand, the analysis of mitochondrial DNA does not reveal the presence of genes encoding for proteins belonging to the family of the  $\text{Na}^+/\text{Ca}^{2+}$  exchanger, suggesting that OMM and the IMM do not possess their own endogenous proteins operating as a  $\text{Ca}^{2+}$  efflux/influx pathway. Therefore, it is conceivable that a member of the NCX family, coded by nuclear DNA, might be present on mitochondria, as it occurs for the majority of mitochondrial proteins (Stojanovski, 2003). Indeed, 90% of mitochondrial proteins are coded by nuclear genes,



synthesized in the cytosol, and, subsequently, imported into mitochondria through protein translocation machineries of the outer and inner membranes (Stojanovski, 2003). More recently, our group provided evidence that the nuclear encoded NCX3 is the only isoform of the  $\text{Na}^+/\text{Ca}^{2+}$ -exchanger localized within the OMM, where it plays a relevant role in the control of mitochondrial  $\text{Ca}^{2+}$  homeostasis both under basal and under hypoxic conditions. This novel finding is not in contrast with the results of Palty et al, since it possible to hypothesize that the  $\text{Na}^+/\text{Ca}^{2+}$  exchanger in mitochondria requires two consecutive steps. The first, operated by the  $\text{Na}^+$ -sensitive NCLX, mediates  $\text{Ca}^{2+}$  transport from the matrix to the intermembrane space, and the second, operated by mNCX3, that promotes  $\text{Ca}^{2+}$  efflux from the intermembrane space to the cytosol. This interpretation is in line with the recent physiological role attributed to the OMM in the control of mitochondrial  $\text{Ca}^{2+}$  cycling. Indeed, although the outer surface of the membrane is not a passive permeable membrane, it does constitute a permeability barrier not only to  $\text{Ca}^{2+}$  influx but also to  $\text{Ca}^{2+}$  efflux (Szabadkai and Duchen, 2008). On the other hand, evidence that mNCX3 is involved in  $\text{Ca}^{2+}$  extrusion is that the  $\text{Ca}^{2+}$  lowering effect found in cells transfected with NCX3 was completely prevented by the benzothiazepine compound CGP-37157, which is considered a selective mitochondrial  $\text{Na}^+/\text{Ca}^{2+}$  exchanger inhibitor (Szabadkai and Duchen, 2008).

The Na<sup>+</sup>-independent Ca<sup>2+</sup> efflux is the main mitochondrial Ca<sup>2+</sup> efflux system in non-excitabile cells and since no specific cations have been found to be exchanged with Ca<sup>2+</sup> it is believed to be a Ca<sup>2+</sup>-H<sup>+</sup> exchanger (Saris et al., 2005). This transport mechanism requires transmembrane potential, since it is not observed in non-energized mitochondria, thus indicating that it is not an electroneutral passive 1Ca<sup>2+</sup>-2H<sup>+</sup> exchanger (Gunter et al., 1991). Indeed, this system is able to extrude Ca<sup>2+</sup> against a gradient that is much higher than predicted from thermodynamics for an electroneutral H<sup>+</sup>/Ca<sup>2+</sup> exchanger, which indicates that it uses a component of the electrochemical gradient for its activity. A characteristic of this transporter is that it saturates at low calcium loads and its kinetics is extremely slow (Bernardi, 1999). This emphasizes a feature of the mitochondrial Ca<sup>2+</sup> machinery: it is equipped with high V<sub>max</sub> uptake transport systems coupled to slow and easily saturable release systems, increasing the risk of Ca<sup>2+</sup> overload.

As discussed above, cytosolic Ca<sup>2+</sup> elevations are rapidly transmitted to the mitochondrial matrix, where they amplify the activity of Krebs cycle enzymes and of the ATP synthase, thereby increasing the production of ATP (Denton et al., 2009; Jouaville et al., 1999). During physiological Ca<sup>2+</sup> elevations, the boost of ATP enables PMCA to extrude the cytosolic calcium and to sustain neuronal activity. During ischemia however, the levels of oxygen and glucose drop rapidly, impairing the production of ATP by mitochondria and by cytosolic glycolysis. As a result, ATP dependent calcium extrusion

mechanisms progressively come to a halt because the intracellular reservoir of ATP is depleted by the continuous activity of the  $\text{Na}^+/\text{K}^+$  ATPases. The importance of the  $\text{Na}^+/\text{K}^+$  ATPases in “stealing “ ATP from PMCA could be directly demonstrated as PMCA activity, which collapsed during metabolic depletion, could be rescued by inhibition of the  $\text{Na}^+/\text{K}^+$  ATPases (Castro et al., 2006). PMCA inhibition amplifies the cytosolic calcium elevations that are transmitted to the mitochondrial matrix, and can then triggers a vicious sequence of mitochondrial calcium overload, mitochondrial dysfunction, release of mitochondrial pro-apoptotic factors, and the activation of death signals (Starkov et al., 2004; Duchen, 2004; Kristian et al., 1998). Mitochondria located near the plasma membrane (subplasmalemmal mitochondria) are more exposed to calcium overload due to their proximity to plasma membrane voltage-sensitive calcium channels and to the functionally incapacitated PMCA (Frieden et al., 2002; Frieden et al., 2005; Jousset et al., 2007). Intracellular mitochondria are also at risk however, and calcium release from the endoplasmic reticulum has been associated to ischemia induced-cell damage (Paschen et al., 1999; Pisani et al., 2000; Chen et al., 2008). Mitochondria are embedded within sheets of endoplasmic reticulum and the two organelles are maintained in very close proximity by linker proteins (Csordas et al., 2010; Csordas et al., 2006). Because of this proximity, the release of calcium ions through IP3 receptor of the endoplasmic reticulum readily triggers an entry of calcium in adjacent

mitochondria (Giacomello et al., 2010; Rizzuto et al., 1993). Thus, neuronal mitochondria are exposed both to  $\text{Ca}^{2+}$  ions entering across membrane channels and to  $\text{Ca}^{2+}$  released from endoplasmic reticulum  $\text{Ca}^{2+}$  stores.

Currently, it is unclear whether the mitochondrial matrix  $\text{Ca}^{2+}$  elevations occurring during ischemia are causally related to the neuronal cell death that occurs after cerebral ischemia. The best established link between mitochondrial  $\text{Ca}^{2+}$  and cellular toxicity is the opening of a  $\text{Ca}^{2+}$ -activated channel located in the inner mitochondrial membrane, the permeability transition pore (PTP) (Halestrap, 2009; Rasola et al., 2007, Zoratti et al., 2010), responsible for the so-called mitochondrial permeability transition (MPT). The MPT (Zoratti et al., 1995) is a phenomenon induced by high levels of matrix  $\text{Ca}^{2+}$  accumulation and oxidative stress, responsible for a sudden increase in the mitochondrial inner membrane permeability to solutes with molecular masses up to 1500 Da. The PTP is a  $\text{Ca}^{2+}$ , ROS (reactive oxygen species), voltage-dependent and CsA sensitive high-conductance channel, located in the inner mitochondrial membrane. Despite the great interest generated by this channel, which has been extensively characterized at the pharmacological and biophysical level, the molecular identity of the PTP is not known. The classical model envisions a supramolecular complex spanning the double membrane system of mitochondria, localized at contact sites (Brdiczka et al., 1998). Proteins of all mitochondrial compartments have been proposed to be part of the PTP (Halestrap, 2009; Rasola et al., 2007;

Zoratti et al., 2005), in particular cyclophilin D (CypD) in the mitochondrial matrix, the adenine nucleotide translocator in the inner membrane and mitochondrial porin VDAC in the outer membrane. Surprisingly, genetic studies have demonstrated that MPT can still be observed in mitochondria devoid of each of these proteins (Basso et al., 2005; Kokoszka et al., 2004; Krauskopf et al., 2006). Some other proteins, including inter-membrane and cytosolic proteins and even the proapoptotic Bcl-2 family protein Bax, have been proposed to be part of the PTP under particular conditions (Marzo et al., 1998; Narita et al., 1998), but  $\text{Ca}^{2+}$ -dependent MPT was shown to be independent of Bax (De Marchi et al., 2005). Although the proteins responsible for this important mitochondrial process have not yet been identified, the generation of CypD-knockout mice has now established an unquestionable role for CypD in facilitating PTP opening (Basso et al., 2005; Baines et al., 2005; Nagakawa et al., 2005; Schinzel et al., 2005; De Marchi et al., 2006). The biophysical properties (conductance, voltage-dependence, selectivity) of PTP were indistinguishable in mitochondria isolated from isogenic wild-type and engineered mice lacking CypD (De Marchi et al., 2006), but CsA only inhibited the PTP in wild-type mice (Basso et al., 2005; De Marchi et al., 2006), demonstrating that CypD represents the target for PTP inhibition by CsA. That the permeability transition is triggered by an elevation in the free  $\text{Ca}^{2+}$  concentration within the mitochondrial matrix was discovered early (Haworth et al., 1979; Crompton et al., 1988; Petronilli et al.,

1993). Chelation of matrix  $\text{Ca}^{2+}$  induces a rapid closure of PTP and divalent cations such as  $\text{Mg}^{2+}$ ,  $\text{Mn}^{2+}$ ,  $\text{Ba}^{2+}$  and  $\text{Sr}^{2+}$ , instead of inducing PTP opening, can act as inhibitors of  $\text{Ca}^{2+}$  trigger sites (Bernardi et al., 1992). The relationship between MPT and cerebral ischemia inferred from the effects of CsA was nicely confirmed in CypD-knockout mice, by measuring the infarct size after cerebral ischemia/reperfusion injury induced by the occlusion of the middle cerebral artery (Schinzel et al., 2005). In this study, Schinzel and colleagues demonstrated that isolated, CypD-deficient mitochondria showed an increased capacity to retain calcium and were resistant to  $\text{Ca}^{2+}$ -induced MPT in swelling experiments. When they induced ischemia/reperfusion, a dramatic decrease in infarct size (62%) was recorded in the brains of CypD deficient mice, suggesting an essential role for CypD in cell death in the brain. A correlation between gene dosage and the extent of injury was elegantly established, by recording a partial protection in heterozygous mice (37% of reduction in infarct size). These data proved that conditions required for the activation of PTP were present during ischemia/reperfusion, as suggested by earlier pharmacological studies (Shiga et al., 1992; Khaspekov et al., 1999; Matsumoto et al., 1999). Although the causal relationship between mitochondrial  $\text{Ca}^{2+}$  accumulation and PTP opening is well established, and despite the fact that MPT invariably leads to neuronal cell death, these relationships do not necessarily imply that matrix  $\text{Ca}^{2+}$  accumulation is directly responsible for the injuries related to cerebral

ischemia. The group of Lemasters, for example, has proposed that mitochondrial  $\text{Ca}^{2+}$  overload is a consequence, rather than a cause, of the bioenergetic failure that follows MPT onset. In this view, the mitochondrial  $\text{Ca}^{2+}$  elevation is only a signature of diseased mitochondria and is not involved in the induction of the MPT, which occurs after reperfusion (Kim et al., 2006). In this study of adult rat myocytes, ROS but not  $\text{Ca}^{2+}$  overload has been suggested to trigger pH- and MPT-dependent death after ischemia/reperfusion. Another important parameter to take into account is the timing of the PTP opening during ischemia/reperfusion. In the heart, there is a broad consensus that during ischemia the factors favoring PTP opening (increased matrix  $\text{Ca}^{2+}$  and depolarization) are balanced by PTP antagonists (intracellular acidosis, high levels of  $\text{Mg}^{2+}$  and ADP) that prevent PTP opening during ischemia (Di Lisa et al., 2011). Upon reperfusion, oxygen and substrate supplies are restored to the tissue, mitochondria re-energize, take up the  $\text{Ca}^{2+}$  that has accumulated in the cytosol during ischemia, and produce a burst in ROS. The combination of these factors provides ideal conditions for triggering PTP opening (Halestrap et al., 2010; Di Lisa et al., 2011). Direct methods to assess PTP opening in intact hearts support the concept that PTP is more likely to open upon reperfusion (Griffiths et al., 1995; Di Lisa et al., 2001). Whether the same sequence of events also occurs in ischemic brain is not known, and further studies are needed to determine the precise timing of the PTP opening during cerebral ischemia.

The recent identification of the proteins involved in mitochondrial  $\text{Ca}^{2+}$  uptake and release provides new opportunities to study the role of mitochondrial  $\text{Ca}^{2+}$  in neuronal death during cerebral ischemia. The role of calcium in MPT activation and cell death can now be directly tested by modulating the expression levels of mitochondrial transport proteins. Targeting the proteins that control the fluxes of  $\text{Ca}^{2+}$  should reveal whether altered mitochondrial  $\text{Ca}^{2+}$  handling is causally related to ischemic neuronal death, and can potentially increase the repertoire of therapeutic tools to treat ischemic brain diseases.

It is not yet clear which sources of  $\text{Ca}^{2+}$  and which pathways are involved, however the defective cross-talk between intracellular stores is presumed in the etiopathogenesis of the injury (Bano D., 2007).

The activation of intracellular second messengers  $\text{Ca}^{2+}$ -related, the increase of reactive oxygen species, but also the same hypoxia, trigger the expression of many pro-inflammatory genes that induce synthesis of transcription factors such as the nuclear factor  $\kappa\text{B}$  (NF $\kappa\text{B}$ ) (O'Neill L.A. et al., 1997) and the hypoxia inducible factor (HIF) (Ruscher, K. et al., 1998). Moreover, the inflammation mediators, such as platelet activating factor (PAF), tumoral necrosis factor (TNF- $\alpha$ ) and the interleukin-1, are produced by impaired neurons and glial cells. The activation of inflammatory process after stroke can induce many consequences and seems to be related to the apoptosis as well. Antibodies against adhesion molecules, as well as limits in the



development of inflammatory process, are able to reduce the number of cell dead in ischemic brain (Chopp, M. et al., 1996).

In light of what has been said so far it is clear that the pathogenesis of ischemic injury is complex and depends on multifactorial events and that intracellular ionic homeostasis alterations, impaired mitochondrial function and the production of free radicals play a major role leading to neuronal death consequent to anoxic damage.

This complexity of events explains the great discrepancy between the increasing number of cerebral ischemic accidents and the difficulty with which, unfortunately, are made studies in the development of effective treatments able to inhibit or slow the neuronal death following an ischemic episode. Hence the urgent need to identify new potential targets for the development of therapeutic strategies able to defend the ischemic brain.

In this context, one approach that has received particular attention in recent years has been an important aspect of the ischemic process referred as ischemic tolerance. This phenomenon, also known as ischemic preconditioning, is a sub-lethal anoxic insult that increases the tissue tolerance to a subsequent and potentially lethal ischemia. The cerebral ischemic preconditioning is very important because give us the possibility to study the endogenous mechanisms activated in neurons to promote cell survival after this sublethal stimulus, and thus to develop new therapeutic

strategies more selective for the treatment of the ischemic disease. The great interest in the cerebral ischemic preconditioning and in the tolerance evoked by itself, also comes from the similarity of this phenomenon with some clinical situations encountered in the human brain. Indeed, the tolerance evoked by brief ischemic episodes reminds the transient ischemic attacks (TIAs) that don't cause structural damage but appear to protect against a subsequent "stroke" (Weih et al., 1999a; Moncayo et al., 2000) and in clinical practice that precede a subsequent stroke.

## ***ISCHEMIC PRECONDITIONING***

Ischemic preconditioning or ischemic tolerance of the brain, heart, and other organs refers to a natural adaptive process that can be induced by a variety of sublethal insults (e.g., transient hypoxia, spreading depression, oxidative stress, hyperthermia, heat shock), and which increases the tissue tolerance to a subsequent, potentially lethal ischemia. This adaptive cytoprotection is a fundamental capability of living cells, allowing them to survive exposure to potentially recurrent stressors. Clearly identified in the heart by Murry et al. in 1986, preconditioning and subsequent ischemic tolerance were then demonstrated in the brain (Kitagawa et al., 1990), and attracted rapidly the interest of clinical and basic neuroscientists for several reasons. First, this biological process became widely recognized as a pertinent and effective experimental probe to understand how the brain protects itself against ischemia, thereby providing an innovative approach for the discovery of novel cerebroprotective strategies. Second, retrospective case-control studies showed a clinical correlate of the phenomenon discovered experimentally.

Preconditioning induces two different time windows of tolerance. The first phase, named “rapid preconditioning”, occurs rapidly after the stimulus and is characterized by post-translational and metabolic modifications, ionic channel activation but is not dependent on the synthesis of new proteins; the second phase, named “delayed preconditioning”, corresponds to a robust state of

tolerance that is usually detectable 24 h after preconditioning induction, peaks at 3 days, fades after 7 days and requires the synthesis of new proteins. In the brain, conversely in the heart, the tolerance induced by ischemic preconditioning is generally of the delayed type (Kitagawa et al., 1990; Liu et al., 1992; Nishi et al., 1993; Gidday et al., 1994; Goldberg et al., 1998).

The induction of ischemic tolerance is accompanied by substantial changes in gene expression, suggesting that preconditioning stimulates a fundamental genomic reprogramming of cells that confers cytoprotection and survival (Stenzel-Poore et al., 2007). The genomic response after ischemic preconditioning is a signature of the complex interplay of multiple signalling pathways. These highly specialized pathways in different cell types of the brain seem to refine the cellular and systemic response to combat the noxious stimulus. Hundreds of genes are either upregulated or downregulated in response to ischaemic preconditioning stimuli (Bernaudin et al., 2002; Stenzel-Poore et al., 2003; Tang et al., 2006). Changes in gene expression differ between harmful ischemia and ischemic preconditioning. Preconditioning seems to attenuate the response to ischemia (Stenzel-Poore et al., 2003) and the tolerance induced by it changes the expression of genes involved in the suppression of metabolic pathways, immune responses, ion-channel activity, and blood coagulation (Stenzel-Poore et al., 2003). Gene expression is regulated by transcription factors but also depends on

epigenetic mechanisms such as DNA methylation and histone modification, which modify the chromatin structure that controls access of transcription factors to regulatory loci. Inhibitions of DNA methylation and increased histone acetylation have neuroprotective effects in experimental models of stroke (Endres et al., 2000).

## ***EXPERIMENTAL MODELS OF ISCHEMIC PRECONDITIONING***

Starting from these assumptions many subsequent studies, both *in vivo* and *in vitro*, have confirmed the presence of the ischemic brain tolerance, and, in order to clarify the intracellular mechanisms underlying this phenomenon, several experimental models have been developed to reproduce *in vitro* and *in vivo* this condition.

### ***In vivo models***

A large number of studies in animal model of cerebral ischemia have shown that brief periods of non-lethal focal or global ischemia may have a protective effect against further episodes of brain ischemia.

About three decades ago, was proposed a model in which Mongolian gerbils were subjected to bilateral carotid occlusion for 5 min followed by processing of their brains for light and electron microscopy. The authors reported that there occurred rapid ischemic changes in the CA4 region, with development of slower abnormalities in the CA2 area and even still slower changes in the CA1 region (Kirino et al., 1982). Using that model, others groups (Kitagawa et al., 1990) showed that exposure to 2 min of ischemia caused depletion of high-energy phosphate compounds and perturbation of protein synthesis without any evidence of neuronal death. In addition, they found that single episodes of 2-min ischemia were able to provide protection against cell death

caused by 5 min ischemic events performed 1 or 2 days after the pretreatment. Moreover, two 2-min ischemic pretreatments performed one day apart provided complete protection against cell death induced by 5 minutes of ischemia (Kitagawa et al., 1990). The preconditioning with 2 min of ischemia prevented damage to the hippocampal CA1 area following 3 min of ischemia induced 3 days later (Kato et al., 1992). In addition, the pretreatment with unilateral occlusion of the middle cerebral artery (MCA) could induce ischemic tolerance in the hippocampal CA1 neurones of gerbils (Miyashita et al., 1994). Moreover, two minutes of preconditioning ischemia also protected gerbil hippocampal neurons against damage caused by a 3.5 min ischemia performed 3 days later (Ohtsuki et al., 1996). Because previous investigators had focused their attention on younger animals and because most ischemic strokes occur in older individuals, other groups (Dowden et al., 1999) used 18- to 20-month-old gerbils to test if they could observe preconditioning-induced protection in these animals. They found that two episodes of global ischemia, each lasting 1.5-minutes, separated by 24 hours protected animals who underwent a 5-minute occlusion of both carotid arteries 72 hours later. They also reported normal microtubule-associated protein-2 expression in the hippocampal CA1 region of preconditioned animals in contrast to the complete loss of microtubule-associated protein-2 staining observed in ischemic animals.

The ischemic preconditioning was reproduced using rats to test if sub-lethal ischemia could protect against neuronal death caused by subsequent lethal ischemic insults (Liu et al., 1992). The forebrain ischemia for 3 min was able to protect against hippocampal CA1 neuronal damage caused by 6 and 8 min of ischemia but not against damage caused by 10 min of ischemia (Liu et al., 1992). Other groups (Simon et al., 1993) also showed that two brief periods of global cerebral ischemia, separated by 24 h of normoxia, which did not cause significant cell death in the brain, were able to significantly reduce the size of an infarct caused by permanent MCA occlusion. It was also reported that global ischemia can provide substantial protection against strokes caused by MCA occlusion in the rat brain (Matsushima et al., 1995). Moreover, in another very interesting study, the possibility that ischemic preconditioning applied only 30 min before a lethal insult could still offer protection against histological damage in the rat brain was assessed. In that study, the conditioning ischemic insult lasted 2 min. After only 30 min of reperfusion, the rats were subjected to 10-min of ischemia. This procedure caused a significant degrees of protection in the hippocampal CA1 subfield and in the cortex (Perez-Pinzon et al., 1997).



### ***In vitro models***

Cell culture systems have also been used extensively to investigate the effects of preconditioning. The model of combined oxygen and glucose deprivation (OGD) that recapitulates the cascades of changes that occur during ischemia-induced injuries in the mammalian brain (Goldberg et al., 1993; Martin et al., 1994) has been largely used to investigate the potential bases of preconditioning-induced neuroprotection. In that model, transient deprivation of rodent cortical cell cultures of both oxygen and glucose causes neuronal swelling, followed by severe neuronal degeneration over a period of several hours even when the cells were returned to normal media (Goldberg et al., 1993). Acute and delayed injuries were inhibited by combined removal of extracellular  $\text{Ca}^{2+}$  in conjunction with  $\text{Na}^{+}$  or  $\text{Cl}^{-}$  substitution. The toxic effects of OGD appear to be mediated by excitotoxic damage because it causes a large increase in the levels of extracellular glutamate and because N-methyl-D-aspartate (NMDA) receptor antagonists provided protection against both the early and the late damage caused by OGD. Different groups used the OGD model to study the effects of preconditioning in vitro. They found that cellular damage was significantly reduced in cells pre-exposed to different short times (from 5 minutes to 90 minutes) of OGD with an interval of 24-48 hours before a second exposure to 3 hours of OGD. Interestingly, it was also reported that the preconditioned cultures were not protected against NMDA, kainate-, or glutamate-induced neuronal death (Grabb et al., 1999).

This is in contrast with a study in which it was reported that subjecting rat cortical cultures to a preconditioning OGD paradigm provided cross-tolerance to NMDA-induced cell death. It was also reported that NMDA receptor blockade during preconditioning by OGD eliminated tolerance (Tauskela et al., 2001). Another study comparing various preconditioning manipulations to OGD found that cycloheximide, heat stress, MK801 were as protective as OGD preconditioning in both acute and delayed models of ischemia-induced neuronal death (Meloni et al., 2002). Similar results have been reported using hippocampal slice cultures (Hassen et al., 2004), after combined deprivation of glucose and amino acids (Gaspar et al., 2006), and after exposure to neuronal cultures to various agents including anesthetic agents and diazoxide (Bickler et al., 2005; Kis et al., 2003), mild hypoxic insults (Khaspekov et al., 1998), as well as stimulation of adenosine receptors (Heurteaux et al., 1995). Interestingly, the OGD model was reproduced in organotypic cultures in which provided protection against cell death and the transient loss of evoked potential responses (Badaut et al., 2005).

### ***Other models of ischemic preconditioning***

In addition to ischemia preconditioning, there are other paradigms that have been shown to protect against ischemic injuries. These include spreading depression, hyperoxia and oxidative stress, prolonged hypoperfusion, and, as alluded to above, hyperthermia or heat shock. Therefore, one stressor can

promote 'cross-tolerance' to another. The sheer variety of stimuli capable of inducing an ischemia-resistant phenotype in the brain indicates that the signalling pathways activated by these different triggers converge downstream on some common, fundamental mechanisms that ultimately account for the protection. Many exogenously delivered chemical preconditioning agents (for example, inflammatory cytokines, anaesthetics and metabolic inhibitors) can also induce ischemic tolerance, raising the hope that in the future it will be possible to pharmacologically activate these distal pathways in the human brain (Gidday, 2006). The efficacy of cross-tolerance, relative to the classic ischemic preconditioning (i.e., brief, sublethal cerebral ischemia, with subsequent test ischemia that would damage the naive brain) may be more modest, and it appears to vary with the nature and intensity of the first challenge. The window of ischemic tolerance may be also shifted or different; for example, the "tolerizing" effect of LPS needed more time to be manifest, starting at 48 h after LPS injection and reaching full potential at 72 h as described in the study of Tasaki et al. (Tasaki et al., 1997). Cross-tolerance is an important feature, but potentially misleading by suggesting that different preconditioning stimuli result ultimately in similar mechanisms of tolerance. That may be only partially true, and dependent on the nature of the stimuli considered. Comparison of the genomic profiles of brain responses to ischemia from mice preconditioned with either brief ischemia or low-dose LPS revealed that a substantial subset of the

differentially-expressed genes were unique to each preconditioning stimulus (Stenzel-Poore et al., 2007). This suggests that the nature of the preconditioning stimulus may determine a specific neuroprotective phenotype. When a specific variable (gene expression, level of a specific protein, etc.) is found to be altered in animals subjected to both preconditioning stimulus and test insult, this change may be causally related to the mechanism of protection, and this interpretation is often favored by authors. However, as the fundamental nature of preconditioning is to protect the brain against a subsequent insult, an alternative and valid interpretation may be that the identified change simply reflects, or is an epiphenomenon of, the reduced damage.

## ***INTRACELLULAR MECHANISMS ACTIVATED BY ISCHEMIC PRECONDITIONING***

Recently, three phases temporally consecutives of the preconditioning phenomenon have been identified: induction, transduction, and tolerance.

To induce tolerance, the preconditioning stimulus must be recognized by molecular sensors as a sign of something potentially much more severe to come. So far, numerous types of sensor have been identified, including neurotransmitter, neuromodulator, cytokine and toll-like receptors (Kariko et al., 2004) as well as ion channels and redox-sensitive enzymes. In turn, these sensors activate transduction pathways that initiate the adaptive response. Although dependent in part on the nature of the preconditioning stimulus, members of these transduction pathways for which there is strong general support include mitogen-activated protein kinases (MAPKs) and their phosphorylated Ras, Raf, MEK and ERK subfamilies (Dawson et al., 2000; Gonzalez-Zulueta et al., 2000; Jones et al., 2004), mitochondrial ATP-sensitive K<sup>+</sup> (KATP) channels (Heurteaux et al., 1995; Yoshida et al., 2004) Akt (also known as protein kinase B) (Yano et al., 2001; Wick et al., 2001; Hashiguchi et al., 2004) and the protein kinase C- $\epsilon$  isoform (Raval et al., 2003). The possibility that the nitric oxide-based adaptive response to hypoxia in *Drosophila* (Wingrove et al., 1999) is evolutionarily conserved suggests that this multifunctional modulator might be a logical choice as an autocrine and paracrine mediator of preconditioning stress. Indeed,

pharmacological and genetic evidence supporting the involvement of nitric oxide in the transduction process is continuing to mount. Given the redox sensitivity of many kinases and transcription factors, reactive oxygen species might also serve as transducers (Puisieux et al., 2004; Zhang et al., 2004; Ravati et al., 2001). Adenosine, another prototypical paracrine mediator and 'retaliatory metabolite', the production of which is linked to ATP degradation, seems integral to tolerance induction in some models (Heurteaux et al., 1995; Yoshida et al., 2004; Plamondon et al., 1999; Nakamura et al., 2002). Finally, caspases might be essential induction catalysts, given that cyclic AMP responsive element-binding protein (CREB), the p50 and p65 subunits of nuclear factor- $\kappa$ B (NF- $\kappa$ B), and protein kinase C and other kinases are caspase substrates (McLaughlin et al., 2003; Algeciras-Schimnich et al., 2002). Notably, some of the aforementioned molecular transducers and signalling intermediates also serve as post-ischaemic effectors of the ischaemia-tolerant phenotype (Dirnagl et al., 2003).

### ***Role of nitric oxide***

Nitric oxide (NO) is synthesized from L-arginine and oxygen by the oxidoreductase nitric oxide synthase, an enzyme present in three isoforms: neuronal NOS (nNOS), endothelial NOS (eNOS) and inducible or 'immunological' NOS (iNOS) (Alderton et al. 2001). NO exerts its biological effects by reacting with oxygen, superoxide, or transitional metal centres

(Gross & Wolin, 1995; Pacher et al. 2007). The cytoprotective effects of NO typically occur through direct post-translational modification of proteins (S-nitrosylation), through activation of the cGMP second messenger system and through reaction with superoxide to form peroxynitrite. S-Nitrosylation, produced by the direct reaction of NO with cysteine thiol groups, has been linked to neuroprotection through inhibition of NMDA receptor activity (Lei et al. 1992; Lipton et al. 1993; Kim et al. 1999). In addition, S-nitrosylation may confer neuroprotection by inhibition of apoptosis, particularly through inactivation of caspases, proteases that degrade key intracellular proteins (Melino et al. 1997; Budihardjo et al. 1999; Zhou et al. 2005). S-Nitrosylation also leads to activation of the mitogen-activated protein kinase/extracellular-signal-regulated kinase (MAPK/ERK) pathway (Yun et al. 1998), which triggers the expression of pro-survival genes in cortical neurons. S-Nitrosylation can also play a role in gene expression through modification of transcription factors involved in neuronal survival, such as CREB (cyclic adenosine monophosphate response element binding; Lonze & Ginty, 2002). Recently, a study in cortical neurons found that the brain-derived neurotrophic factor (BDNF), well known for its role in cell survival, induced CREB binding to target DNA (Ricchio et al. 2006). These effects of NO on CREB-DNA binding are mediated by S-nitrosylation of histone deacetylase 2 (HDAC2) (Ricchio et al. 2006). In addition, S-nitrosylation of the p50 subunit negatively regulates the pro-inflammatory transcription factor NF- $\kappa$ B (nuclear

factor  $\kappa$  light chain enhancer of activated B cells), possibly contributing to an anti-inflammatory environment (Colasanti & Persichini, 2000). A main biological target of NO is the haem-containing enzymesoluble guanylyl cyclase (sGC), which catalyses the conversion of GTP to cGMP. Numerous studies suggest that NO-dependent elevations in cGMP and consequent activation of protein kinase G (PKG) inhibit cell death and activate prosurvival pathways. A role for NO in the expression of specific neuroplasticity-associated proteins, including BDNF, was recently demonstrated in cortical neurons and in the whisker barrel cortex, an effect that involves cGMP/PKG and ERK signalling (Gallo & Iadecola, 2011). Similarly, cGMP improves synaptic function and CREB activation in a mouse model of Alzheimer's disease (Puzzo et al. 2009). A glimpse into the cellular mechanisms of NO protection downstream of PKG is provided by studies in cardiomyocytes. Following ischaemia–reperfusion, NO-dependent PKG activity leads to activation of protein kinase C $\epsilon$ , which phosphorylates mitochondrial ATP-sensitive K<sup>+</sup> (mitoKATP) channels (Costa & Garlid, 2008). The subsequent opening of the channels results in cytoprotection by inhibiting the mitochondrial transition pore (Costa & Garlid, 2008). Opening of mitoKATP channels is also associated with neuroprotection through mechanisms related to stabilization of mitochondrial function and prevention of apoptosis (Liu et al. 2002; Mayanagi et al. 2006). NO rapidly reacts with superoxide to form peroxynitrite (Beckman et al. 1990). Besides its well-established its toxic



potential at high concentrations (Calabrese et al. 2009), at low concentrations peroxynitrite can act as a signalling molecule (Liaudet et al. 2009). For instance, in the vasculature, peroxynitrite is able to induce vascular relaxation and inhibition of platelet aggregation (Liu et al. 1994; Moro et al. 1994). Peroxynitrite protects cardiomyocytes from ischaemia–reperfusion injury (Lefer et al. 1997), an effect associated with inhibition of leukocyte adhesion to endothelial cells. In addition, peroxynitrite leads to redox modulation of proteins within signalling cascades that mediate neuroprotection (Liaudet et al. 2009). For example, peroxynitrite activates ERK and Akt signalling in different cells promoting their survival (Jope et al. 2000; Pesse et al. 2005; Li et al. 2006). These observations suggest that NO has powerful protective effects that involve not only pro-survival signalling pathways, but also blood flow and vascular inflammatory events that play a prominent role in ischaemic brain injury. These aspects of NO biology are critical for understanding the protective effect that NO exerts in the context of ischaemic preconditioning. Indeed, NO has been implicated both in early and late preconditioning in the newborn as well in the adult brain. NO derived from eNOS and nNOS is essential for induction of tolerance in models of early preconditioning. For example, the lesion produced by neocortical injection of the glutamate receptor agonist NMDA was markedly reduced 1 h after administration of the pro-inflammatory mediator LPS in mice (Orio et al. 2007). The effect was not abolished by the protein synthesis inhibitor

anisomycin, attesting to the fact that the protection did not require new protein synthesis, a hallmark of early preconditioning. The protection conferred by LPS was abolished by inhibitors of nNOS, and could be reconstituted by neocortical injection of a NO donor. Furthermore, the tolerance was not observed in nNOS or eNOS null mice, but was preserved in iNOS null mice and in mice treated with the iNOS inhibitor aminoguanidine (AG) (Orio et al. 2007). In agreement with LPS preconditioning in NMDA lesions, the early tolerance to focal ischaemic injury produced by a transient episode of focal cerebral ischaemia was not observed in eNOS or nNOS null mice (Atochin et al. 2003), suggesting that eNOS and nNOS-derived NO is also involved in other preconditioning and injury modalities. These observations implicate eNOS and nNOS, but not iNOS, as sources of NO. There is evidence that the mechanisms by which NO induces early tolerance involve the second messenger cGMP. The mechanisms by which cGMP leads to tissue protection may involve PKG and mitoKATP channels. These observations, collectively, indicate that NO is critical for the establishment of ischaemic tolerance induced by transient ischaemia or by the proinflammatory mediator LPS, an effect involving activation of sGC and cGMP Production. Mounting evidence suggests that NO is also involved in delayed preconditioning. In hypoxic preconditioning in a neonatal rat model of ischaemic–hypoxic damage, the protection is abolished by the non-selective NOS inhibitor nitro-L-arginine, but not by the nNOS inhibitor 7-nitroindazole

or the iNOS inhibitor AG implicating eNOS as the source of NO. Similar results were obtained in another model of preconditioning in neonatal rats (Lin et al. 2010). With some exceptions (Hashiguchi et al. 2004; Vellimana et al. 2011), the majority of studies on the role of NO in delayed preconditioning in the adult have indicated that the source of NO is iNOS. Studies in adult mice have revealed a key role of iNOS in the preconditioning produced by LPS or transient forebrain ischaemia. iNOS null mice do not develop tolerance to focal ischaemia or NMDA lesions after treatment with LPS or transient forebrain ischaemia (Cho et al. 2005; Kawano et al. 2007; Kunz et al. 2007). Similarly, AG prevents the development of tolerance in these models. The preconditioning is associated with an increase in mitochondrial resistance to calcium-mediated depolarization (Cho et al. 2005), supporting the involvement of mitochondria also in the mechanisms of delayed tolerance. There is increasing evidence that preconditioning stimuli, in addition to changes in gene expression and mitochondrial function, also improve cerebrovascular function (Gidday, 2006). Therefore, the mechanisms of ischaemic tolerance may also include vascular effects that lead to an improvement of cerebral perfusion and blood–brain barrier function.

Moreover, it is well known that NO generated by nNOS activates the G-protein Ras (Lander et al., 1995; Yun et al., 1998; Kolch et al., 2000). Ras, in turn, also stimulates the PI3 kinase/Akt pathway. This signaling pathway is now largely

recognized as one of the most relevant pathways in regulating neuronal survival (Song et al., 2005). Indeed, it has various functions: 1) it regulates glucose metabolism; 2) it inactivates the mitochondrial death pathway (i.e., BAD and caspase 9) and FOXO pathway; 3) it activates CREB phosphorylation, which regulates the expression of genes critical for survival, such as those encoding for cytokines and brain derived neurotrophic factor; and 4) it activates the antiapoptotic factor nuclear factor- $\kappa$ B through the activation of I $\kappa$ B kinase complex (Fukunaga and Kawano, 2003). In addition, Akt regulates Ca<sup>2+</sup> homeostasis through the potentiation of L-type voltage-gated calcium channels (Blair et al., 1999), sarco(endo)plasmatic Ca<sup>2+</sup> ATPase (Kim et al., 2003), and intracellular ligand-gated ion channels (Barac et al., 2005), thus enhancing neuronal survival. Moreover, it has been demonstrated that the overexpression of constitutively active Akt1, the most highly diffused and expressed isoform in the brain (Hanada et al., 2004; Hui et al., 2005), induces an up-regulation of NCX1 and NCX3. Indeed, the overexpression of NCX1 and NCX3 isoforms induced by this serine-threonine kinase was only partially responsible for the 80% reduction of hypoxia induced cell death by Akt1-positive mutants, because siRNA-NCX1 and siRNA-NCX3 reduced this neuroprotective effect to values of 26 and 28%, respectively. This result suggested that Akt1, besides activating several neuroprotective pathways, produces an overexpression of NCX1 and NCX3 that may contribute to the reduction of hypoxia-induced cell death through an

improvement of intracellular  $\text{Ca}^{2+}$  and  $\text{Na}^{+}$  dysregulation. Together, these findings indicate that NCX1 and NCX3 constitute novel and additional targets of Akt neuroprotective action.

Overall, these downstream effector mechanisms of Akt1 result in a program of cell survival.

### ***Role of inflammatory cytokines***

Inflammatory cytokines are known to have an important role in acute stroke. Cytokines such as interleukin-1 (IL-1), and TNF- $\alpha$  are important mediators of the inflammatory reactions seen in cerebral ischemia (Kariko et al., 2004). TNF- $\alpha$  is one of the pro-inflammatory cytokines and is expressed in the ischemic brain. The effects of pretreatment with TNF administered intracisternally in mice that were subjected to MCAO 48 h later were studied (Nawashiro et al., 1997). A significant reduction in infarct size was noted in mice pretreated with TNF at the dose of 0.5 microgram/mouse. Immunohistochemical analysis of brains subjected to 24 h of MCAO revealed a significant decrease in CD11b immunoreactivity after TNF pretreatment compared with the MCAO control. Therefore, TNF induces significant protection against ischemic brain injury and is likely to be involved in the signaling pathways that regulate ischemic tolerance. Liu et al (Liu et al., 2000) demonstrated that preconditioning of rat cortical neurons with mild hypoxia protected them from hypoxia and oxygen and glucose deprivation

(OGD) injury 24 h later (50% protection). Interestingly, TNF- $\alpha$  pretreatment could be substituted for hypoxic preconditioning (HP). HP was attenuated by TNF- $\alpha$ -neutralizing antibody. The role of IL-1 in the induction of tolerance to global ischemia in Mongolian gerbils was also demonstrated (Ohtsuki et al., 1996). Arterial IL-1 $\alpha$  and IL-1 $\beta$  became elevated between 1 and 3 days after a 2-min ischemic exposure. Recombinant human IL-1 receptor antagonist (IL-1ra) blocked ischemic tolerance induction by 2-min preconditioning ischemia. The possible mechanisms of IL-1 action include release of arachidonic acid, enhancement of NMDA activation and stimulation of nitric oxide synthase (Huang et al., 2006).

### ***Role of heat shock proteins (HSPs)***

Although most protein synthesis is inhibited after cerebral ischemia, the remaining active protein synthesis may have an important role in maintaining cell viability. HSPs are molecular chaperones and among the proteins synthesized during ischemia. HSPs are expressed both constitutively (cognate proteins) and under stressful conditions (as inducible forms). In addition to heat shock, a variety of stressful situations including environmental (ultraviolet radiation or heavy metals), pathological (infections or malignancies), or physiological (growth factors or cell differentiation) stimuli induce a marked increase in HSPs synthesis, known as the stress response (Tsan et al., 2004). Inducible HSPs are thought to assist the

maintenance of cellular integrity and viability by preventing protein denaturation and improper polypeptide aggregation during exposure to physiochemical insults (Ravati et al., 2001).

HSP70 is the most abundant HSP found in cells. It is expressed constitutively and is only mildly inducible. Many researchers have shown the extensive links between HSP70 overexpression and tolerance in ischemic brain injury. Some of them have reported an increase in HSP70 expression in an *in vivo* model of preconditioning (McLaughlin et al., 2003). Furthermore, HSP70 messenger RNA (mRNA) was compared in the cases of a 15-min ischemia 2 days after sham treatment and a 15-min ischemia 2 days after 10-min preconditioning (Sakurai et al., 1998). The authors reported that HSP70 mRNA in the motor neurons was strong at 8 h after preconditioning with 10-min ischemia, mild at 1 or 2 days, and not observed at 7 days after preconditioning. Others reported that thermal preconditioning with 44°C body temperature protected rat cerebellar granule neurons by modulating HSP70 expression. HSP70 mRNA was detected after thermal preconditioning at 30, 60, and 90 min and increased gradually with time, whereas HSP70 antisense oligodeoxynucleotides inhibited the protective effects of thermal preconditioning against apoptosis (Chen et al., 2004). All these results implicate that HSP70 is an important player in the preconditioning process.

The underlying mechanisms of ischemic brain tolerance HSP70-mediated are not only related to its important functions in protein refolding and

transport. Emerging evidence suggests that the HSP70 family is also capable of binding and sequestering activated caspases, such as Apaf and AIF (Yenari et al., 1999). Overexpression of HSP70 inhibits the activation of NF- $\kappa$ B, which is persistently activated during ischemia and appears to promote apoptotic cell death. On the contrary, the deletion of the HSP70 gene notably increases cytochrome *c* release into the cytoplasm and subsequent caspase-3 activation, thereby exacerbating apoptosis and increasing infarction volume after focal cerebral ischemia (Lee et al., 2004). Furthermore, HSP70 expression is regulated by transcription factors, the activity of which is increased by ERK phosphorylation (Minet et al., 2000), a process which has been implicated in preconditioning.

### ***Role of the anti-apoptotic protein Bcl-2***

The protein Bcl-2 is an anti-apoptotic protein that resides in the outer mitochondrial membranes and the membranes of the endoplasmic reticulum. Overexpression of Bcl-2 is known to block the release of cytochrome *c*, which contributes to a signaling pathway leading to apoptosis. Indeed, two minutes of bilateral carotid artery occlusion in gerbils produced an increase in Bcl-2 at 30 h and peaked at 96 h, suggesting that the expression of Bcl-2 is involved in ischemic preconditioning (Kato et al., 1991). In addition, 20 min of transient focal ischemia, representing a sub-lethal insult, produced ischemic tolerance (attenuated infarction volume) (Shimizu et al., 2001). The results of Western



blot analysis from tolerant caudate-putamen demonstrated an increase in Bcl-2 expression 3–7 days after preconditioning. Immunocytochemical examination also found that Bcl-2 was expressed in cells with both neuronal and non-neuronal morphology in striatum. Bcl-2 antisense oligodeoxynucleotides (ODNs) treatment reduced expression of Bcl-2 in the striatum and blocked the induction of tolerance by preconditioning ischemia. Therefore, Bcl-2 appears to be a major determinant in this model of induced tolerance to focal ischemia. The transcription factors driving the induction of Bcl-2 during tolerance include CREB, indeed it has been described a cAMP-responsive element in Bcl2 promoter. As mentioned above, CREB is regulated by NMDA receptor activation, and phosphorylated robustly in the penumbral region of the preconditioned rats (Nakajima et al., 2002). Moreover, multiple protein kinases can activate transcription via the CRE, by phosphorylation of CREB, further implicating that the interrelation of these different molecular regulatory pathways could be involved in ischemic preconditioning.

### ***Role of adenosine A<sub>1</sub> receptors and ATP-sensitive K<sup>+</sup> channels***

In 1995, was demonstrated the essential role of adenosine, adenosine A<sub>1</sub> receptors, and ATP-sensitive K<sup>+</sup> channels in ischemic brain preconditioning. Adenosine is an endogenous neuroprotectant that can inhibit the release of excitatory amino acids. When ischemia occurs, adenosine can increase conspicuously. Adenosine inhibits synaptic transmission, decreases K<sup>+</sup>-

stimulated glutamate release, and inhibits presynaptic calcium fluxes via adenosine A1 receptors. The response to calcium influx is also important for the development of protection against ischemia, because calcium influx has been linked to the production of reactive oxygen species and the initiation of a number of signaling cascades leading to cell damage (Kristian et al., 1998). Many reports have shown that preconditioning-induced neuroprotection is dependent on adenosine A1 receptors. In rats, ischemic preconditioning increased adenosine A1 receptor immunoreactivity in the hippocampal CA1 region at days 1, 3, and 7 after preconditioning induction, within the window of ischemic tolerance (Zhou et al., 2004). Some unspecific and specific adenosine A1 receptor antagonists abolished the ischemic tolerance. Moreover, it was demonstrated that the tolerance isoflurane-induced against focal cerebral ischemia model was attenuated by adenosine A1 receptor antagonists in a rat middle cerebral artery occlusion (MCAO) (Liu et al., 2006). Activation of the ATP-sensitive potassium channels (K<sup>+</sup> ATP channels) also has a role in ischemic brain preconditioning. Blockade of the K<sup>+</sup> ATP channels abolished preconditioning and the protection afforded by adenosine and R-PIA (an adenosine A1 receptor agonist). By contrast, a K<sup>+</sup> ATP channel opener (RP-52891, aprikalim) induced ischemic tolerance. Meanwhile, recent evidence showed that transient infusion of the K<sup>+</sup> ATP channel antagonist sulfonylurea tolbutamide prior to ischemia could block preconditioning induced protection after forebrain ischemia, whereas

pinacidil, a K<sup>+</sup> ATP channel agonist, can emulate this phenomenon in hippocampal slices (Pérez-Pinzón et al., 1999). Although the precise K<sup>+</sup> ATP channels involved in the tolerance remain undefined, two K<sup>+</sup> ATP channels have been described recently. One of these channels resides in the mitochondrial inner membranes; the other resides in the plasma membranes. The mitochondrial K<sup>+</sup> ATP (mK<sup>+</sup>ATP) has been suggested to be the key channel involved in ischemic preconditioning, because the mK<sup>+</sup> ATP blocker 5-hydroxydecanoate (5-HD) prevented BIP-induced neuroprotection. It has been hypothesized that opening the mK<sup>+</sup> ATP channels may depolarize mitochondrial membrane potential and promote an increase in the electron transport chain rate and thus increase ATP production (Schultz et al., 1999).

### ***Role of opioid receptors***

There are three types of opioid receptor:  $\delta$ ,  $\kappa$ ,  $\mu$  opioid receptors. The activation of opioid receptors is neuroprotective when the body encounters ischemia, hypoxia and cold. Zhang and collaborators showed that the  $\delta$  opioid receptors were involved in hypoxia preconditioning in cultured rat cortical neurons (Zhang et al., 2001). They observed that  $\delta$  opioid receptor activation protected cortical neurons from hypoxia injury, whereas the  $\delta$  opioid receptor antagonist naloxone blocked such protection. In addition, another group found that morphine (an agonist for  $\delta$ ,  $\kappa$ ,  $\mu$  opioid receptors) and Tan-67 (a selective  $\delta$  receptor agonist) induced a delayed

preconditioning effect both in vivo and in vitro. The morphine preconditioning-induced neuroprotection was inhibited by  $\beta$ -funaltrexamine, a  $\mu$ -opioid receptor antagonist, but not by 7-benzylidenenaltrexone, a  $\delta$ -receptor antagonist, or norbinaltorphimine, a  $\kappa$ -receptor antagonist. The Tan-67 preconditioning-induced neuroprotection was inhibited by 7-benzylidenenaltrexone. These results suggest that the delayed phase of morphine preconditioning may involve  $\mu$  opioid receptors and  $\delta$  opioid receptors. Morphine and Tan-67 may activate a shared intracellular signalling pathway to induce the delayed preconditioning effects in the brain (Zhao et al., 2006).

### ***Role of the ubiquitin-proteasome pathway***

There are two protein degradation systems in mammalian cells, the autophagy/lysosomal pathway and the ubiquitinproteasomal pathway. The ubiquitin-proteasomal pathway has been studied in cerebral ischemia and ischemic preconditioning. In rat forebrain ischemia models, transient cerebral ischemia followed by reperfusion leads to delayed selective neuronal death in hippocampal CA1 pyramidal neurons. Under electron microscopy (EM), visible protein aggregates progressively accumulate in some CA1 neurons and accumulation of the aggregates seem to occur primarily in neurons destined to undergo delayed neuronal death after brain ischemia.

Further evidence (Yoneda et al., 2004; De Gracia et al., 2007) showed that the endoplasmic reticulum (ER), mitochondria and cytoplasm all respond to the accumulation of unfolded proteins by compartment-specific signaling pathways to participate in neuronal injury, whereas ubiquitin-proteasome as well as beneficial chaperones function to prevent protein aggregation and assist in protein folding (Ghaemmaghami et al., 2003; Hebert et al., 2007). Furthermore, by utilizing a rat transient cerebral ischemic preconditioning model, was demonstrated that ischemic preconditioning significantly reduced protein aggregation in CA1 neurons after ischemia. Biochemical analyses revealed that ischemic preconditioning decreased accumulation of ubiquitin-conjugated proteins (ubi-proteins) and reduced free ubiquitin depletion after brain ischemia. Ischemic preconditioning also reduced redistribution of heat shock cognate protein 70 and Hdj1 (HSP40) from cytosolic fraction to protein aggregate-containing fraction after brain ischemia (Liu et al., 2005). Indeed, the proteasome comprises multiple protein subunits and degrades cytosolic proteins as well as misfolded proteins that fail to pass protein quality control in the ER (Wu et al., 2006). Misfolded proteins in the ER are recognized by ER-specific E3 ligases that mediate polyubiquitination of the misfolded protein on the cytosolic side of the ER and are subsequently degraded by the proteasome (Chen et al., 2008). Proteasome degradation of ubi-proteins is strictly ATP-dependent. Ubi-proteins serve as signals to activate heat shock transcription factors to induce expression of molecular chaperones (Voellmy

et al., 2004), which then shield hydrophobic surfaces of proteins in non-native states, thereby blocking their aggregation. Major cellular chaperones are ATPases and assist protein folding through numerous cycles of binding and release of unfolded protein substrates by hydrolysis of ATP. Therefore, ubiquitin–proteasome and molecular chaperons may be concerned with the neuroprotection of ischemic preconditioning and may together prevent protein aggregation in lethal ischemia.

### ***Role of autophagy/lysosomal pathway***

Recently, the roles of autophagy and the lysosomal pathway in cerebral ischemia have attracted intensive attention. Studies have reported the activation of autophagy following ischemic insults, but the contribution of autophagy to neuronal death/ survival is still under debate (Adhami et al., 2006; Carloni et al., 2008; Koike et al., 2008; Rami et al., 2008; Balduini et al., 2009). In an earlier study, we found profound activation of autophagy and lysosomes after permanent middle cerebral artery occlusion (pMCAO) and an autophagic mechanism may contribute to ischemic neuronal injury (Wen et al., 2008). However, autophagy is a double-edged sword (Shintani et al., 2004). Although massive autophagy is associated with cell death through excessive self-digestion and degradation of cellular constitutions, activation of autophagy can also protect the neurons by degrading unfolded proteins and damaged organelles. A study carried out in primary cultured

cardiomyocytes (Dosenko et al., 2006) showed that inhibition of autophagy with 3-methyladenine (3-MA, an autophagy inhibitor) during anoxia-reoxygenation caused an increase in the number of necrotic cells and a decrease of the living cell population. Thus, the process of autophagy during anoxia-reoxygenation was proposed to provide some protective effects. In addition, a recent in vitro study in PC12 cells also showed that autophagy was associated with neuroprotective effects induced by IPC (Park et al., 2009). Inhibition of autophagy, especially during reperfusion or lethal oxygen-glucose deprivation periods, ameliorated the neuroprotective effects of IPC, and also attenuated HSP70 upregulation induced by IPC. Furthermore, our recent observations showed that autophagy was also induced in a rat ischemic preconditioning model. Pretreatment with 3-MA before the onset of ischemic preconditioning completely suppressed the neuroprotective effects of preconditioning. These results strongly supported the hypothesis that induction of autophagy in ischemic preconditioning may play a part in protecting against the sequential lethal cerebral ischemia (Sheng et al, manuscript submitted).

### ***Role of hypoxia inducible factor-1 (HIF-1)***

Activation of hypoxia-inducible factor-1 (HIF-1) has been proposed as a crucial event in the regulation of hypoxia-inducible gene expression in mammalian cells (Semenza, 2000). HIF-1 is a member of the basic helix-

loop-helix family of transcription factors and requires the dimerization of HIF-1 $\alpha$  and HIF-1 $\beta$  subunits for its activity (Wang and Semenza, 1995). Hypoxia-inducible factor-1 heterodimer binds to a specific DNA consensus sequence in the enhancer and promoter regions of many hypoxia-inducible genes (Ebert et al., 1996; Semenza et al., 1994, 1996). Under normoxic conditions, HIF-1 $\alpha$  protein is negatively regulated by ubiquitination and proteasomal degradation (Huang et al., 1998; Kallio et al., 1999) such that constitutive levels of HIF-1 $\alpha$  protein are generally low or undetectable (Wiener et al., 1996; Bergeron et al., 1999b, 2000). During hypoxia, HIF-1 $\alpha$  expression and concomitant HIF-1 DNA binding activity are markedly increased in several cell lines (Wang and Semenza, 1995; Wang et al., 1995; Jiang et al., 1996a) and primary neuronal cultures (Ruscher et al., 1998). In the brain, increased HIF-1 $\alpha$  expression has been observed in rats exposed to systemic hypoxia (Wiener et al., 1996; Bergeron et al., 2000) or subjected to global (Jin et al., 2000) and focal ischemia (Bergeron et al., 1999b; Marti et al., 2000). Interestingly, *in vitro* exposure to the divalent metal cobalt chloride (CoCl<sub>2</sub>) or the iron chelator desferrioxamine during normoxic conditions triggers transcriptional changes that mimic a hypoxic response that is characterized by increased HIF-1 $\alpha$  and HIF-1 $\beta$  expression, up-regulation of HIF-1 binding activity, and consequent HIF-1 target gene expression (Ebert et al., 1996; Semenza et al., 1994, 1996; Zaman et al., 1999). The hypoxia-mimetic effect of these compounds may involve the interaction with a ferroprotein oxygen sensor. Recent evidence



has shown that HIF-1 $\alpha$  and HIF-1 $\beta$  mRNA and protein are induced after preconditioning with hypoxia or CoCl<sub>2</sub> in neonatal rats, suggesting that HIF-1-regulated gene expression may be involved in the hypoxia-induced tolerance to ischemic injury in immature brain (Bergeron et al., 2000).

Recently, we demonstrated that the gene coding for the ubiquitous Na<sup>+</sup>/Ca<sup>2+</sup> exchanger NCX1 protein can be included in the family of genes activated by the transcription factor HIF-1. This regulation assumes particular relevance in light of the recently emerged role of HIF-1 and NCX1 in several pathophysiological aspects of cerebral ischemia. In fact, neuron specific knocking-out of HIF-1 $\alpha$  exacerbates brain damage after tMCAO, suggesting its neuroprotective action (Baranova et al., 2007). Similarly, NCX1 knocking-down by oligodeoxynucleotide antisense strategy increases infarct damage after permanent middle cerebral artery occlusion (Pignataro et al., 2004).

Interestingly, *ncx1* transcript levels have been demonstrated to be up-regulated in regions surrounding the ischemic core (Boscia et al., 2006). Therefore, HIF-1-dependent up-regulation of *ncx1* expression provides an additional mechanism involved in the pro-survival role attributed to HIF-1.

### ***Cellular ionic homeostasis and energy metabolism***

As ischemic preconditioning activates intracellular biological responses prior to a potential lethal insult, it is expected that an increase of energy metabolism or a latency in anoxic depolarization after the onset of ischemic insult might represent the mechanisms by which organs strengthen their tolerance when exposed to a sublethal insult. In this regard, several experiments have been performed, both *in vivo* and *in vitro*, in order to demonstrate that a reduction in energy demand and in the activity of ion channels represent determinant factors for ischemic tolerance (Stenzel-Poore et al., 2003). In fact, impairment in voltage-gated potassium channels has been observed in cortical neurons exposed to brief non injurious oxygen and glucose deprivation. Similarly, *in vivo* experiments demonstrated that ischemic preconditioning prevented the inhibition of Na<sup>+</sup>/K<sup>+</sup>-ATPase activity after brain ischemia in hippocampal and cortical neurons of rats exposed to global forebrain ischemia (De Souza et al., 2000). In the last years, it has been shown that some integral plasma-membrane proteins, involved in the control of Ca<sup>2+</sup> and Na<sup>+</sup> ion influx or efflux and, therefore, responsible for maintaining the homeostasis of these two cations, might function as crucial players in the brain ischemic process (Pignataro et al., 2004). These proteins, by regulating Na<sup>+</sup> and Ca<sup>2+</sup> homeostasis may represent more suitable molecular targets for therapeutic intervention. Indeed, during ischemic preconditioning, the results of *in vivo* experiments in gerbils showed

an increase in  $\text{Ca}^{2+}$ -ATPase activity and an enhancement in mitochondrial calcium sequestration in CA1 hippocampal neurons after preconditioning (Ohta et al., 1996). In line with this result, intracellular calcium imaging performed in hippocampal neurons of preconditioned gerbils showed that the increase in  $[\text{Ca}^{2+}]_i$  occurring after anoxic and aglycemic episodes was markedly inhibited in the ischemic tolerant animals (Shimazaki et al., 1998). The molecular mechanisms underlying this effect are still under investigation. A possible explanation could be increased expression of the  $\text{Ca}^{2+}$ -ATPase isoform 1 [plasma membrane calcium ATPase 1 (PMCA-1)] as recently demonstrated by Kato et al. (Kato et al., 2005). Also, mitochondria co-operate in the regulation of intracellular  $\text{Ca}^{2+}$  homeostasis, both in physiological and in pathological conditions. Although mitochondria have a high capacity for  $\text{Ca}^{2+}$  sequestration, excessive amounts of  $\text{Ca}^{2+}$ , as it occurs during ischemia, impair mitochondrial function. This induces a massive uncoupling of oxidative phosphorylation and reduction of mitochondrial membrane potential with consequent mitochondrial permeability transitional pore (MPTP) opening and reversal of the action of ATPase, which in turn hydrolyses, rather than synthesizes, ATP. As a consequence of ATP depletion and increased permeability of the inner mitochondrial membrane, cell death occurs. Studies performed both in the heart and in the brain suggested that the inhibition of MPTP opening and its signalling cascade represent crucial events responsible for cytoprotection observed in ischemic preconditioning

(Halestrap et al., 2007; Dirnagl et al., 2008). The molecular mechanisms underlining these effects are still the objects of investigation. Nitrite and protein kinases have been proposed as possible MPTP regulators (Zhao et al., 2006; Shiva et al., 2007).

The hypothesis that a modulation of the expression and activity of the sodium calcium exchanger (NCX) might play a role in the regulation of calcium and sodium homeostasis during ischemic tolerance cannot be ruled out. Indeed, the  $\text{Na}^+/\text{Ca}^{2+}$  exchanger, in parallel with selective ion channels and ATP dependent pumps, maintains the physiological cytosolic concentrations of these ions (Blaustein and Lederer, 1999). In the brain, unlike other tissues, this exchanger is present in 3 different gene products, named NCX1, NCX2, and NCX3, with a distinct distribution pattern in different brain regions (Canitano et al., 2003). Under physiological conditions, its primary role is to extrude  $\text{Ca}^{2+}$  through a *forward mode* of operation in response to a depolarization or to an increase in intracellular  $\text{Ca}^{2+}$  concentrations coupled to receptor stimulation. However, during hypoxic conditions, owing to the compromission of the 2 plasma-membrane ATP-dependent pumps  $\text{Na}^+/\text{K}^+$  ATPase and  $\text{Ca}^{2+}$  ATPase, NCX assumes a relevant role in controlling the intracellular homeostasis of these two cations since is able to operate in the *reverse mode*, thus extruding  $\text{Na}^+$  ions while promoting  $\text{Ca}^{2+}$  influx (Annunziato et al., 2004). Although this reverse mode of operation in the early phase of anoxia does undoubtedly elicit an increase in  $[\text{Ca}^{2+}]_i$ , its effect

could be beneficial for neurons because it contributes to decrease  $[Na^+]_i$  overload, a phenomenon which would otherwise lead to cell swelling and, thus, sudden necrotic neuronal death. Conversely, in the later phase of neuronal anoxia, when  $[Ca^{2+}]_i$  overload takes place, NCX working in the forward mode can contribute to the lowering of  $Ca^{2+}$  concentrations, and thus can protect neurons from  $Ca^{2+}$  intracellular overload, neurotoxicity and subsequent cell death. A new emerging concept is that NCX1, NCX2, and NCX3 may exert different roles during *in vitro* and *in vivo* anoxic conditions. Indeed, in cells singly and stably transfected with NCX3, this isoform contributes more significantly to the maintenance of  $[Ca^{2+}]_i$  homeostasis during experimental conditions mimicking ischemia, thereby preventing mitochondrial  $\Delta\Psi$  collapse and cell death (Secondo et al., 2007). In addition, in *in vivo* experiments, the selective knocking down of NCX1 and NCX3, but not of NCX2, by antisense oligodeoxynucleotide strategy, (Pignataro et al., 2004) or the disruption of the *ncx3* gene, renders the brain more susceptible to the ischemic insult (Molinaro et al., 2008). Moreover, the induction of permanent middle cerebral artery occlusion in rats, correlates with NCX1 mRNA upregulation in the peri-infarct area thus suggesting the possibility that this isoform could be a new druggable target for the treatment of cerebral ischemia. In line with this hypothesis, we recently demonstrated that NCX1 transcript and protein were up-regulated by ischemic cerebral preconditioning. This effect was mediated by the transcriptional factor HIF1 a

and was accompanied by a relevant neuroprotective effect (Valsecchi et al., 2011).

## **AIM OF THE STUDY**

Since several players are involved in the complex molecular events implicated in ischemic tolerance, the aim of this study was to identify whether the sodium-calcium exchanger might play a role in the neuroprotection induced by cerebral ischemic preconditioning through the regulation of intracellular ionic homeostasis.

To address this issue a model of hypoxia/reoxygenation was set up in vitro by exposing cortical neurons to 3 hours of combined oxygen and glucose deprivation (OGD) followed by subsequent 24 hr exposure to normoxic conditions (reoxygenation), which represented the severe insult. The sub-lethal preconditioning stimulus was reproduced by exposing neurons to 30 minutes of OGD, 24 hours before the lethal insult.

After these treatments, experiments were conducted to detect the rate of neuronal survival by using propidium iodide (PI) and fluorescein diacetate (FDA) assay. The amount of cell death was determined measuring the ratio between the number of PI positive- dead cells compared to the total number of cells (PI and FDA positive cells) (Wei et al. 2000). The results showed that the preconditioning stimulus did not impair neuronal survival and prevented neuronal death induced by OGD/reoxygenation. Once demonstrated the neuroprotective effect of ischemic preconditioning, a further aim of the study was to investigate the molecular and cellular events involved in the regulation

of this phenomenon. In particular, the attention was addressed to the role of  $\text{Na}^+/\text{Ca}^{2+}$  exchanger (NCX). Therefore, the expression and activity of the three NCX isoforms was evaluated by Western Blot analysis, confocal microscopy and single cell microfluorimetry.

The results obtained demonstrated that NCX1 and NCX3 protein expression increased in neurons exposed to the sub-lethal insult within 48 hours, an effect that was observed even when ischemic preconditioning was applied to neurons exposed to OGD/reoxygenation, suggesting the possibility that this effect could be positively correlated with the development of tolerance. Interestingly, the rise in NCXs protein expression occurring during IPC was related to the over-activity of the nNOS synthase and NO production which in turn stimulated Akt activation. Indeed, the treatment with L-NAME and LY294002, well known inhibitors of nNOS and PI3 kinase respectively prevented NCX3 IPC-induced overexpression. Conversely, the effect of IPC on NCX1 expression was counteracted by LY294002 pretreatment, but was not affected by L-NAME. On the other hand, the activation of Akt observed after IPC was inhibited by the treatment with L-NAME, thus supporting the hypothesis that PI3K/akt/NO pathway could play a role in the regulation of NCX1 and NCX3 protein expression in the IPC phase.

To understand the functional implication of NCXs protein expression in the IPC-induced neuroprotection, the activity of the exchanger was evaluated by single cell microfluorimetry. The results showed an increase in NCX activity



in the reverse mode of operation in neurons exposed to IPC that was still detected in preconditioned neurons exposed to OGD/Reoxygenation. This raised activity did not affect cytosolic calcium concentration but promoted ER-calcium refilling. Interestingly, the treatment with siRNA against NCX1 and L-NAME abolished this effect, suggesting the importance of this isoform in the regulation of ER calcium content in a NO dependent manner. Conversely, the treatment with siRNA against NCX3 induced an increase in ER calcium content detectable 30 minutes, 24, and 48 hrs after the IPC stimulus. However, as a reduction in NCX activity was observed after the treatment with siNCX3 as well, the role of NCX3 as possible player in the regulation of mitochondrial calcium concentration ( $[Ca^{2+}]_m$ ) during IPC was also investigated. To address this issue,  $[Ca^{2+}]_m$  was measured, by confocal microscopy and single cell microfluorimetry, in neurons treated with siNCX3 and exposed to IPC and IPC followed by OGD/reoxygenation. The results obtained with this experimental approach suggested that NCX3 during IPC contributes to the regulation of  $[Ca^{2+}]_m$  which results lower compared with that observed in control neurons or in neurons exposed to OGD/reoxygenation. This effect was more pronounced 48 hrs after IPC stimulus and was not affected by the treatment of neurons with siRNA against NCX1. Moreover, during IPC, the treatment of neurons with CGP37157 slightly modifies NCX3 activity on mitochondria inducing an increase in mitochondrial calcium concentration. Interestingly, this effect was

related to an impairment of mitochondrial oxidative capacity during IPC whereas, in the absence of the mNCX inhibitor this sub-lethal insult did not affect mitochondrial function.

Finally, to correlate the role of the NCX isoforms 1 and 3 with the neuroprotection occurring in the ischemic brain tolerance, we assessed neuronal survival in the presence of siRNAs against NCX1 and NCX3. These treatments resulted in an increase of neuronal death after preconditioning insult whereas such stimulation in the absence of treatment did not affect cell viability.

## **MATERIALS AND METHODS**

### ***Primary culture of cortical neurons***

Cortical neurons were prepared from brains of 16-day-old Wistar rat embryos (Charles River) and used at 7 to 10 days in vitro for protein expression analysis. Briefly, dissection and dissociation were performed in  $\text{Ca}^{2+}/\text{Mg}^{2+}$ -free phosphate-buffered saline containing glucose (30 mmol/L). Tissues were incubated with papain for 10 minutes at 37°C and dissociated by trituration in Earl's balanced salt solution containing DNase (0.16 U/mL), bovine serum albumin (10 mg/mL), and ovomucoid (10 mg/mL). Cells, plated in plastic Petri dishes 100 mm diameter with a confluence of  $15 \times 10^6$  (Falcon; Becton-Dickinson) precoated with poly-D-lysine (20  $\mu\text{g}/\text{mL}$ ), were grown in MEM/F12 containing glucose, 5% of deactivated fetal bovine serum, and 5% horse serum, glutamine (2 mmol/L), penicillin (50 U/mL), and streptomycin (50  $\mu\text{g}/\text{mL}$ ; Invitrogen). To carry out single cell videoimaging and confocal microscopy, cultures of cortical neurons were prepared as described previously (Abramov et al., 2007) with modifications, from Wistar rat pups 2–4 days postpartum. The tissue was minced and trypsinized (0.1% for 15 min at 37°C), trituated, and plated on poly-D-lysine-coated coverslips and cultured in Neurobasal medium (Invitrogen) supplemented with B-27 (Invitrogen) and 2mM L-glutamine. Cells were plated at concentration of

1.8x10<sup>6</sup> on 25-mm glass coverslips. Cultures were maintained at 37°C in a humidified atmosphere of 5%CO<sub>2</sub> and 95% air, fed twice a week, and maintained for a minimum of 10 days before experimental use.

### ***Combined oxygen and glucose deprivation (OGD)***

In cortical neurons, OGD insult was reproduced in vitro by exposing cells to a medium previously saturated with 95% N<sub>2</sub> and 5% CO<sub>2</sub> for 20 min and containing (in mmol/L) 116 NaCl, 5.4 KCl, 0.8 MgSO<sub>4</sub>, 26.2 NaHCO<sub>3</sub>, 1 NaH<sub>2</sub>PO<sub>4</sub>, 1.8 CaCl<sub>2</sub>, 0.01 glycine, and 0.001 wt/vol phenol red (Scorziello et al., 2004). Hypoxic conditions were maintained using a hypoxia chamber (Billups Rothemberg Inc; temperature 37°C, atmosphere 5% CO<sub>2</sub>, and 95% N<sub>2</sub>) for 3 hours. Then, following their removal from the hypoxic chamber, they were challenged with a culture medium containing glucose and previously saturated with a mixture of 95% O<sub>2</sub> and 5% CO<sub>2</sub> for 10 min. Thus, reoxygenation (Rx) was achieved by returning neurons to normoxic conditions (5% CO<sub>2</sub> and 95% air) for 24 h.

Instead, the preconditioning stimulus was reproduced by exposing neurons to 30 minutes of OGD.

### ***Western Blot***

Protein samples (50  $\mu\text{g}$ ) were analyzed on 8% sodium dodecyl sulfate polyacrilamide gel with 5% sodium dodecyl sulfate stacking gel (SDS-PAGE) and electrotransferred onto Hybond ECL nitrocellulose paper (Amersham). Membranes were blocked with 5% nonfat dry milk in 0.1% Tween 20 (TBS-T; 2 mmol/l Tris-HCl, 50 mmol/l NaCl, pH 7.5) for 2 h at RT and subsequently incubated overnight at 4 °C in the blocked buffer with the 1:1000 antibody for NCX1 (polyclonal rabbit antibody, Swant); 1:5000 antibody for NCX3 (polyclonal rabbit antibody, Philipson laboratory); 1:1000 antibody for Akt and pAKT (polyclonal rabbit antibody); 1:1000 antibody for  $\beta$ -actin (monoclonal antibody, Sigma). The membranes were washed with 0.1% Tween-20 and incubated with the secondary antibodies for 1 h (1:5000 NCX3 and 1:1000 NCX1; Amersham). Immunoreactive bands were detected with the ECL (Amersham). The optical density of the bands was determined by Chemi Doc Imaging System (Biorad).

### ***Imaging mitochondrial $\text{Ca}^{2+}$ and mitochondrial membrane potential***

Mitochondrial calcium concentration ( $[\text{Ca}^{2+}]_m$ ) was assessed using the fluorescent dye X-Rhod1. Cells were loaded with X-Rhod1 0.2  $\mu\text{M}$  for 15 min in a medium containing 156 mM NaCl, 3 mM KCl, 2 mM  $\text{MgSO}_4$ , 1.25 mM  $\text{KH}_2\text{PO}_4$ , 2 mM  $\text{CaCl}_2$ , 10 mM glucose, and 10 mM HEPES. The pH was adjusted to 7.35 with NaOH. At the end of the incubation, cells were washed

3 times in the same medium. An increase of mitochondrial localized intensity of fluorescence was indicative of mitochondrial  $\text{Ca}^{2+}$  overload.

Mitochondrial membrane potential was assessed using the fluorescent dye tetramethyl rhodamine ethyl ester (TMRE) in the “redistribution mode”. Cells were loaded with TMRE (20 nM) for 30 min in the above described medium. At the end of the incubation, cells were washed in the same medium containing TMRE (20 nM) and allowed to equilibrate. A decline of mitochondria-localized intensity of fluorescence was indicative of mitochondrial membrane depolarization.

Confocal images were obtained using a Zeiss inverted 510 confocal laser scanning microscopy and a 63X oil immersion objective. The illumination intensity of 543 Xenon laser, used to excite X-Rhod-1 fluorescence, was kept to a minimum of 0.5% of laser output to avoid phototoxicity.

### ***[Ca<sup>2+</sup>]<sub>i</sub> measurment***

[Ca<sup>2+</sup>]<sub>i</sub> was measured by single cell computer-assisted videoimaging (Secondo et al., 2007). Briefly, cells, grown on glass coverslips, were loaded with 5 μM Fura-2 acetoxymethyl ester (Fura-2AM) for 30 min at room temperature in normal Krebs solution containing (in mM): 5.5 KCl, 160 NaCl, 1.2 MgCl<sub>2</sub>, 1.5 CaCl<sub>2</sub>, 10 glucose, and 10 HEPES–NaOH, pH 7.4. At the end of the Fura-2AM loading period, the coverslips were placed into a perfusion chamber (Medical System, Co. Greenvale, NY, USA) mounted onto a Zeiss

Axiovert 200 microscope (Carl Zeiss, Germany) equipped with a FLUAR 40X oil objective lens. The experiments were performed as previously reported (Secondo et al., 2007). The amount of  $\text{Ca}^{2+}$  extruded in the cytoplasm upon FCCP exposure was measured as  $[\text{Ca}^{2+}]_i$  increase. This release is widely considered as an index of mitochondrial  $\text{Ca}^{2+}$  efflux.

***Determination of cell death: Propidium iodide/Fluorescein assay***

Neuronal death was evaluated by measuring the ratio between dead and living cells. To quantify cell death after the experimental procedures, the cells were washed with normal Krebs and double stained with  $36\mu\text{M}$  Fluorescein Diacetate (FDA) and  $7\mu\text{M}$  Propidium Iodide (PI) for 5 min at  $37^\circ\text{C}$  in a phosphate buffer solution. Stained cells were examined immediately with a standard inverse fluorescence microscope at 480 nm and 546 nm (Secondo et al 2007). PI- and FDA-positive cells were counted in three representative high power fields of independent cultures and cell death was determined by the ratio of the number of PI positive cells/PI + FDA-stained positive cells (Wei et al 2000).

***Determination of mitochondrial function: MTT assay***

Neuronal survival was assessed by using 3-(4,5-dimethylthiazol-2-yl)-2,5, diphenyltetrazolium bromide (MTT) (Hansen et al. 1989; Amoroso et al. 1999). The assay was based on the red-ox ability of living mitochondria to

convert dissolved MTT into insoluble formazan. Briefly, after treatments, the medium was removed and cells were incubated in 2 mL of MTT solution (0.5 mg/mL) for 1 h in a humidified 5% CO<sub>2</sub> incubator at 37°C. To stop incubation, MTT solution was removed and 1 mL dimethyl sulfoxide was added to solubilize the formazan product. The absorbance was monitored at 540 nm with a Perkin-Elmer LS 55 luminescence spectrometer (Perkin-Elmer Ltd, Beaconsfield, England). The data are expressed as percentage of cell viability compared to sham-treated cultures.

### ***Small interfering RNA against NCX1 AND NCX3***

The mammalian expression vector, pSUPER.retro.puro (Oligo-Engine), was used to express siRNA against NCX1 (siRNA-NCX1; NM\_019268) in cortical neurons. To prepare siRNA-NCX1, a 60-base oligonucleotide and another oligonucleotide with the complementary sequence were annealed and inserted into pSUPER.retro.puro as previously reported by Formisano et al. A mismatch sequence cloned in the same vector was used as an experimental control. To express siRNA against NCX3, the gene-specific insert contained a 19-nucleotide sequence corresponding to the coding region +124 to +142 relative to the first nucleotide of the start codon of rat NCX3 (GenBank accession no. U53420), whose specificity was verified by BLAST, was used. A mismatch sequence cloned in the same vector and pSUPER.retro.puro vector itself were used as experimental controls.



After 72 hours' plating, cortical neurons were transfected with siRNA-NCX1, siRNA-NCX3 or siRNA mismatch in OPTIMEM by means of lipofectamine LTX (Invitrogen). After 5 hours, it was replaced with fresh medium.

### ***Materials***

All the reagents were purchased from Sigma Chemicals (Milan Italy), unless otherwise specified. LY294002 was supplied by Calbiochem (Inalco, Milan, Italy). L-NAME (500  $\mu\text{mol/L}$ ) and PD98059 (25  $\mu\text{mol/L}$ ) were present in the medium 30 min before exposure to IPC and during the IPC phase.

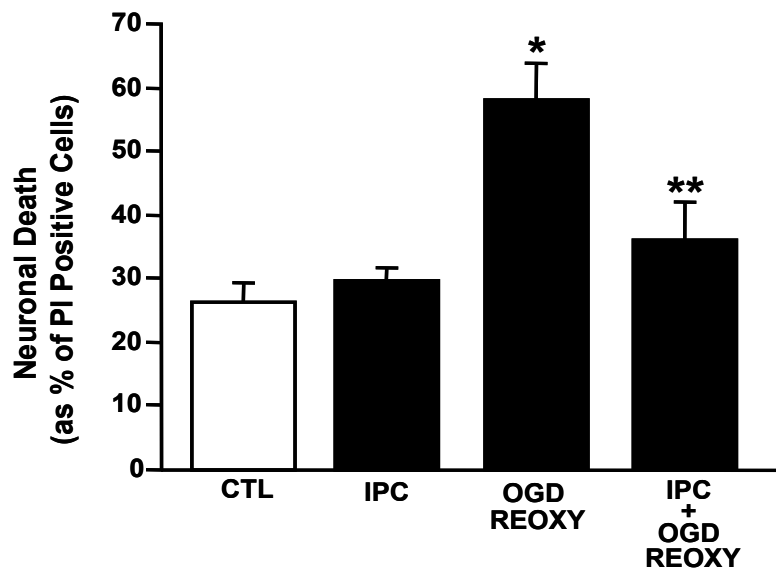
### ***Statistical analysis***

Western blot analyses were repeated three times. All the other experiments were performed in triplicate and repeated at least three times. Data were expressed as mean  $\pm$  SEM values and as percentages of basal values. Statistical significance among the means was determined by the ANOVA followed by the Newman–Keuls test. A p-value  $<0.05$  was considered statistically significant.

## RESULTS

### ***Effect of ischemic preconditioning on neuronal survival of cortical neurons exposed to OGD/REOXYGENATION***

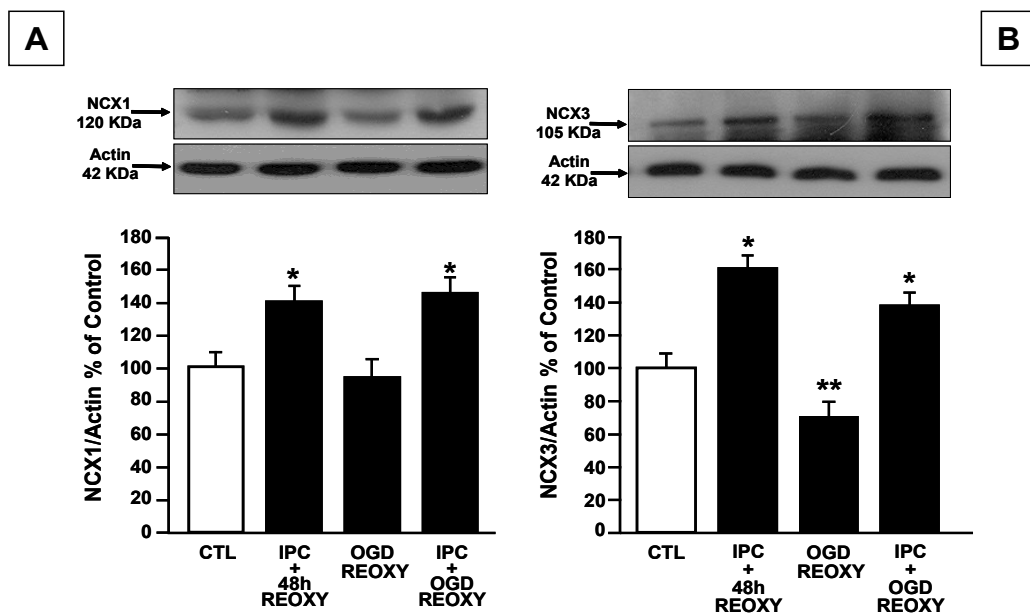
The exposure of cortical neurons to 30 minutes of OGD, a condition that is comparable to a sublethal ischemic insult and commonly defined as ischemic preconditioning (IPC), did not affect neuronal survival. 3 hours of OGD followed by 24 hours of reoxygenation, representing the lethal insult, caused an increase in neuronal death. However, this effect was thwarted when the cells were pre-exposed, 24 hours before the lethal insult, to IPC followed by 24 hours of reoxygenation. (Figure 1).



**Figure 1: Effect of ischemic preconditioning on neuronal survival of cortical neurons exposed to OGD/REOXY:** cortical neurons (12 DIV) were exposed to IPC (30 min di OGD) followed by 24 hrs of normoxia, and then to 3 hrs of OGD followed by 24 hrs of reoxygenation. Cell viability was measured by propidium iodide and fluorescein diacetate assay at the end of reoxygenation. \*P<0.05 vs CTL and IPC. \*\*P<0.05 vs OGD/Reoxy.

## ***NCX1 and NCX3 protein expression after ischemic preconditioning in cortical neurons exposed to OGD/REOXYGENATION***

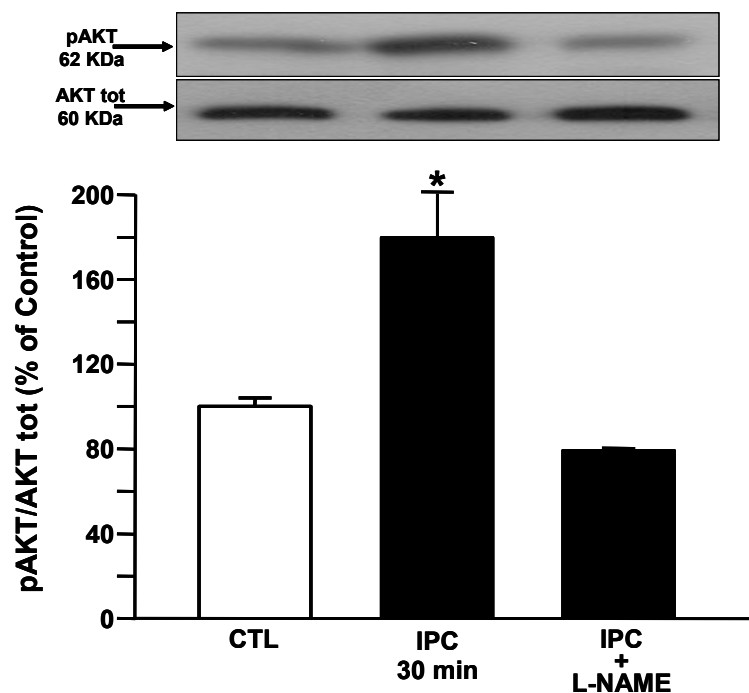
The exposure of cortical neurons to IPC followed by 48 hours of reoxygenation determined a significant increase in NCX1 (Figure 2A) and NCX3 (Figure 2B) protein expression. The same increase was observed in neurons exposed to 3 hrs of OGD preceded, 24 hours before, by a sublethal insult followed by 24 hours of normoxia. These results create the conditions to investigate first the molecular mechanisms underlying the NCX1 and NCX3 increased expression, and to understand if this increase is positively related to the neuroprotection induced by cerebral ischemic preconditioning.



**Figure 2: NCX1 e NCX3 expression after IPC in cortical neurons exposed to OGD/REOXY:** cortical neurons exposed to IPC followed by 24 hrs of normoxia, and to 3 hrs of OGD followed by 24 hrs of reoxygenation. Protein expression was evaluated by western blotting at the end of reoxygenation (A) NCX1 expression. (B) NCX3 expression. \*P<0.05 Vs CTL and OGD/REOXY.

### ***Akt activation in cortical neurons exposed to IPC***

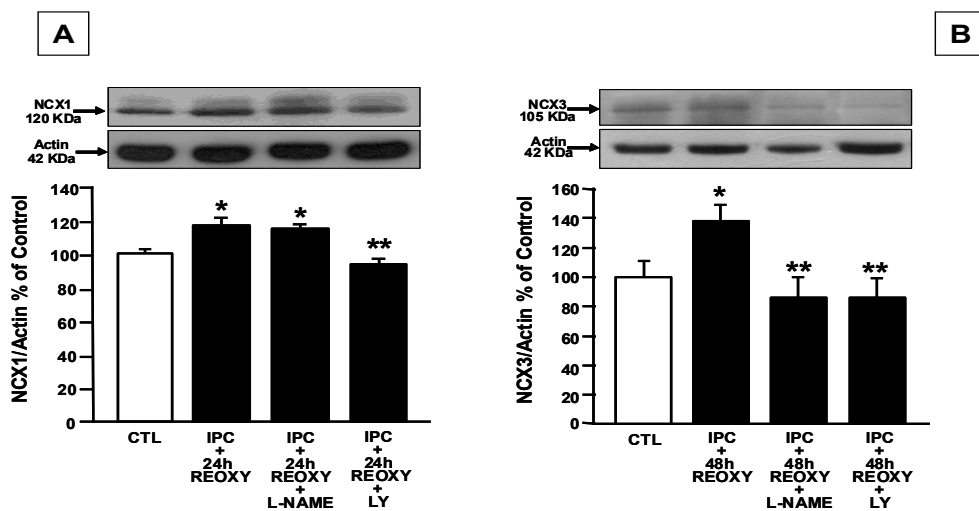
The exposure of cortical neurons to IPC yielded a significant increase in phosphorylated, therefore activated, Akt expression. In order to understand if the biochemical mechanisms responsible for this rise were related to the increased nitric oxide (NO) production induced by IPC, Akt activation was explored after the treatment with L-NAME, a well known NOS inhibitor (Knepper and Kurylo 1998). The results demonstrated that this treatment was able to induce a significant decrease in pAkt expression (Figure 3), suggesting a role of NO in the regulation of Akt activity in ischemic tolerance.



**Figure 3: Effect of IPC and NOS inhibition on Akt activation:** cortical neurons (12 DIV) were exposed to IPC (30 min OGD) with and without L-NAME. Protein expression was evaluated by western blotting at the end of IPC stimulus. \*P<0.05 Vs CTL and IPC+L-NAME.

### **NCX1 and NCX3 expression after LY294002 and L-NAME treatment in cortical neurons exposed to IPC**

To correlate the increase in the expression of NCX1 and NCX3 observed in cortical neurons exposed to IPC with the rise in NO production and the activation of Ras/IP3/Akt pathway, the expression of the above mentioned proteins was investigated in cortical neurons treated with LY294002 (8 $\mu$ mol/L), a well known IP3-kinase inhibitor, and L-NAME (500  $\mu$ mol/L) before the exposure to a sublethal insult. The data confirmed that LY294002 was able to reduce significantly the NCX1 expression, whereas L-NAME did not affect its expression (Figura 4A). Instead, both LY294002 and L-NAME were able to reduce NCX3 expression (Figura 4B), suggesting that the two different isoforms of NCX might be target of the transductional Ras/IP3/Akt pathway activated by NO, probably through different mechanisms.



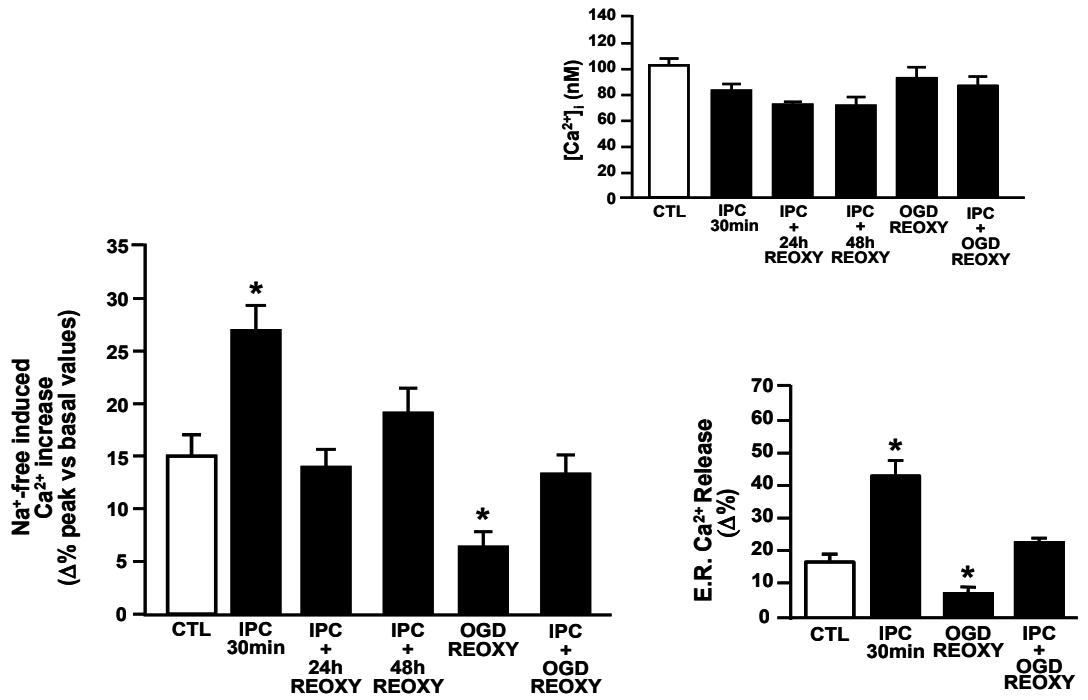
**Figure 4: NCX1 and NCX3 expression after the treatment with LY294002 and L-NAME:** cortical neurons exposed to IPC and treated with LY294002 and L-NAME. Protein expression was evaluated by western blotting.(A) NCX1 expression. (B) NCX3 expression. \*P<0.05 Vs CTL; \*\*P<0.05 Vs IPC.

### ***NCX activity after ischemic preconditioning in cortical neurons exposed to OGD/REOXYGENATION***

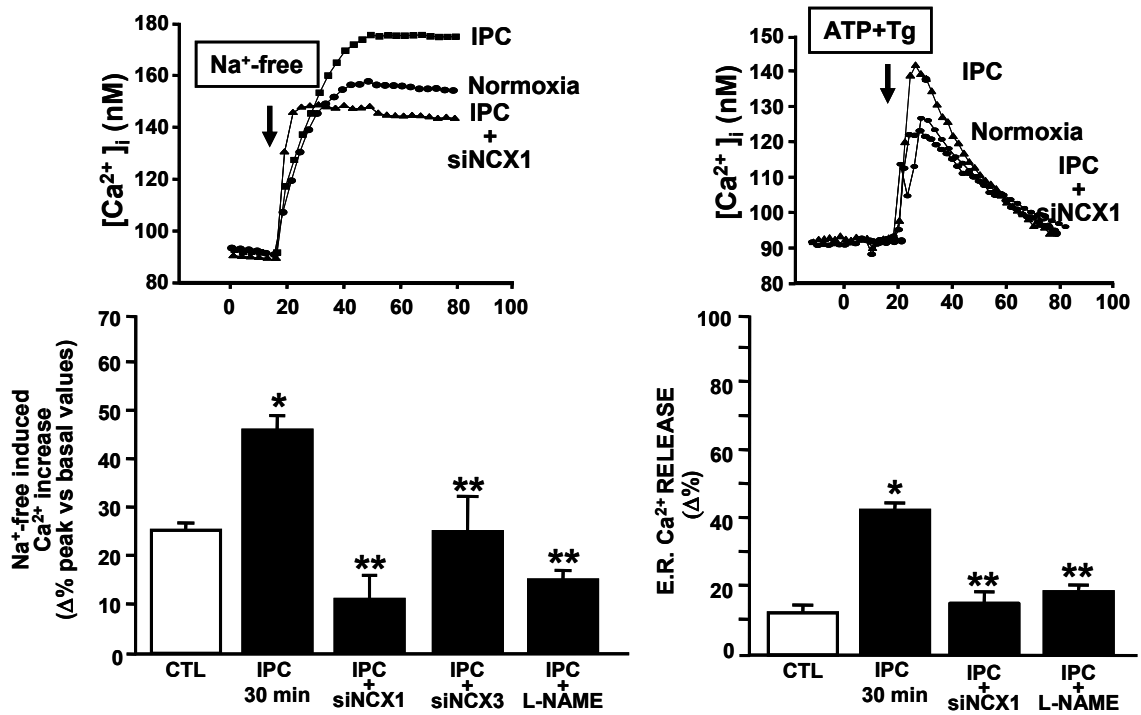
Parallel studies have been conducted to assess NCX activity during ischemic tolerance. The experiments were carried out by using single cell microfluorimetry with the fluorescent probe Fura-2AM. The results obtained demonstrated an early increase in NCX activity in the reverse mode of operation in cortical neurons exposed to IPC and IPC followed by OGD/Reoxygenation. Since this increased activity did not affect  $[Ca^{2+}]_i$ , it has been hypothesized that the increase in NCX activity observed during IPC could be related with endoplasmic reticulum (ER)  $Ca^{2+}$  refilling. To this aim, cortical neurons exposed to 30 min of IPC were loaded with Fura-2AM and stimulated with a solution containing thapsigargin to induce a massive calcium release from intracellular IP3 sensitive calcium stores. By this approach, it was possible to verify that the calcium content in the ER is higher in cortical neurons exposed to IPC compared to non preconditioned neurons (Figure 5A). Since it has been demonstrated that NCX1 plays a role in ER  $Ca^{2+}$  refilling and this is considered a protective mechanism able to prevent apoptotic death associated to ER stress (Sirabella et al. 2009), the hypothesis that NCX1 might regulate the entry of calcium into endoplasmic reticulum during the preconditioning was investigated. To verify this hypothesis the experiments described above were performed in presence of siRNA against NCX1 and NCX3. The results demonstrated that only the

siNCX1 was able to prevent ER  $\text{Ca}^{2+}$  refilling during ischemic preconditioning. Moreover, the same effect was observed after the pretreatment of neurons with L-NAME (Figure 5B), thus suggesting again that NCX1 has a crucial role to prevent neuronal death during ischemic preconditioning by stimulating ER calcium refilling, probably in a NO dependent manner. The involvement of NCX1 in the regulation of ER calcium content is an early event since it was detectable within 24hr by the IPC stimulus time in which the amount of calcium into ER is higher when compared with that observed in control neurons. Conversely, 48 hrs after IPC the amount of calcium into the ER was similar to that observed in the control cells. Interestingly, siRNA against NCX3 induced an increase in ER calcium content both after 30 minutes and after 24 and 48 hrs after the IPC stimulus. In these conditions it is possible to speculate that the impairment of NCX3 activity working in the forward mode of operation, leads to the increase of intracellular calcium content, further contributing to NCX1-induced ER calcium refilling. The treatment of cortical neurons with CGP37157, the inhibitor of the mitochondrial sodium calcium exchanger, during IPC further supports this hypothesis. Indeed, in these experimental conditions the amount of calcium content in the ER was comparable to that observed in the untreated cells, thus excluding the involvement of NCX3 in the regulation of ER calcium refilling (Figure 5C).

**A**

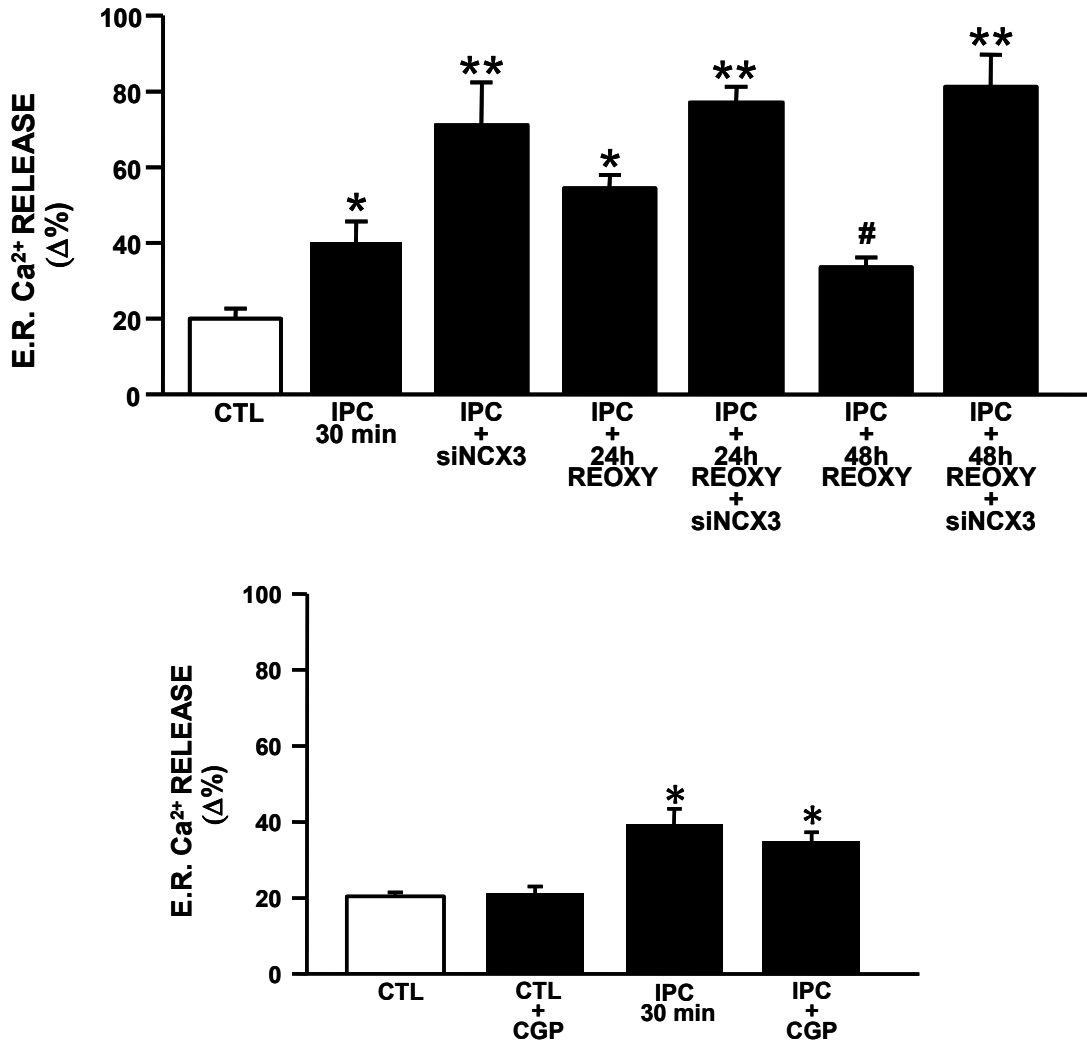


**B**





C

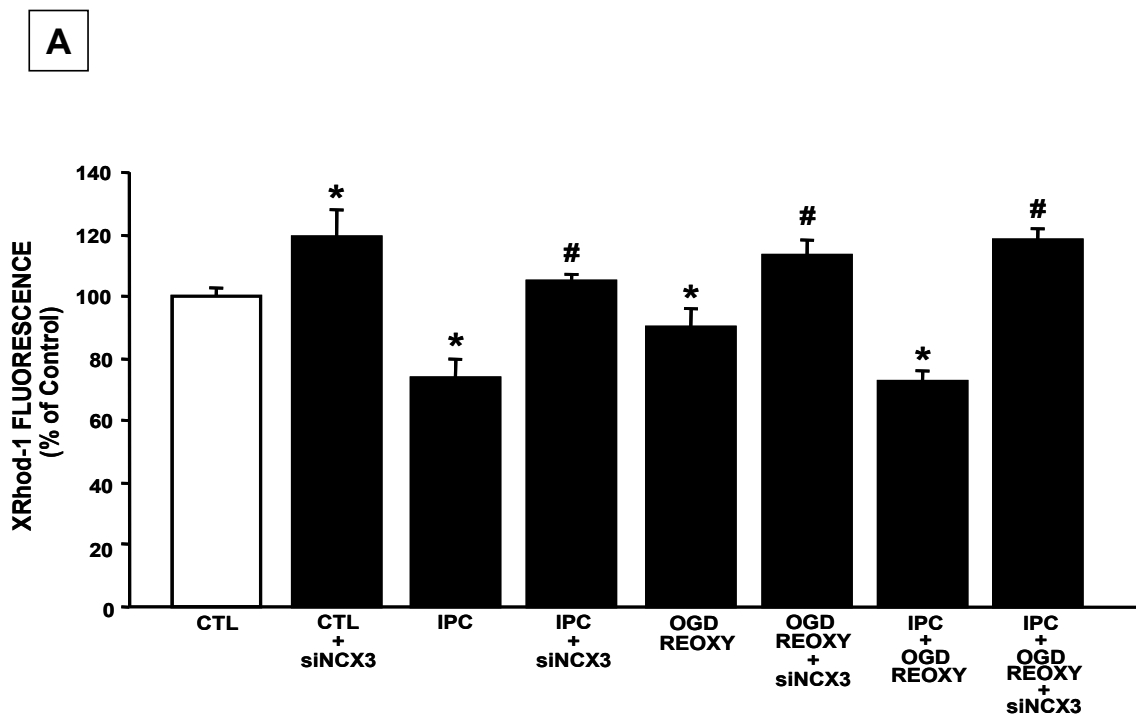


**Figure 5: Effect of IPC on NCX activity and ER Ca<sup>2+</sup> refilling:** (A) cortical neurons exposed to IPC, followed by 24 and 48 hrs of normoxia and to 3 hrs of OGD followed by 24 hrs of Reoxygenation. [Ca<sup>2+</sup>]<sub>i</sub>, NCX activity and ER calcium release, after thapsigargin stimulation, were evaluated at the end of the treatments. \*P<0.05 Vs CTL; (B) effect of siNCX1 and L-NAME on NCX activity and Ca<sup>2+</sup> release from ER. \*P<0.05 Vs CTL; \*\*p<0.05 Vs IPC 30 min. (C) effect of siNCX3 and CGP on Ca<sup>2+</sup> release from ER. \*P<0.05 Vs CTL; \*\*p<0.05 Vs not silenced neurons; \*\*p<0.005 Vs IPC+24h reoxy and IPC+48h reoxy.

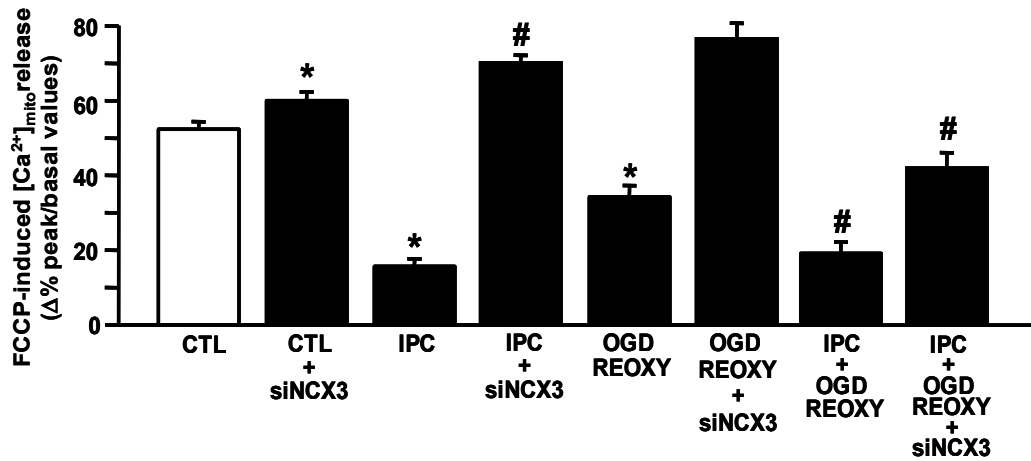
***[Ca<sup>2+</sup>]<sub>m</sub> in cortical neurons treated with siRNA against NCX3 and exposed to OGD/REOXYGENATION***

Since the previous experiments demonstrated that NCX3 knocking down impairs NCX activation during IPC and stimulate ER calcium refilling, the next experiment was addressed to understand how NCX3 could contribute to this phenomenon. On the basis of data obtained from our research group showing that NCX3 is the only isoform present on mitochondria where it plays a key role in the mitochondrial calcium handling, and considering that mitochondria work in concert with ER in the regulation of intracellular calcium homeostasis, we performed experiments aimed to evaluate the contribution of NCX3 in the regulation of mitochondrial calcium concentration in cortical neurons exposed to ischemic preconditioning. The results showed an increase in mitochondrial calcium concentrations after the silencing of NCX3, whereas the exposure of neurons to the only IPC stimulus was able to reduce significantly the calcium levels, even when exposed to 3 hours of OGD followed by 24 hours of reoxygenation (Figure 6A). These data, obtained by using confocal microscopy and the fluorescent probe X-Rhod1, were confirmed with single cell microfluorimetry experiments. To this aim, neurons exposed to IPC were loaded with Fura 2 and then treated with FCCP uncoupler (250 nM) to induce mitochondrial depolarization and Ca<sup>2+</sup> extrusion. The amount of Ca<sup>2+</sup> extruded in the cytoplasm is considered an index of mitochondrial Ca<sup>2+</sup> content. The results obtained with this

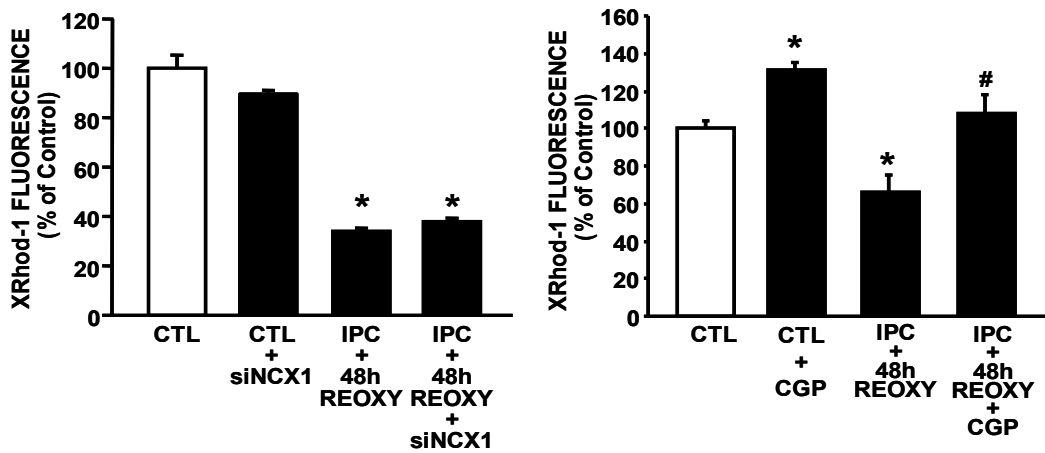
experimental approach suggested that NCX3 during IPC contributes to the regulation of  $[Ca^{2+}]_m$  which results lower when compared with that observed in control neurons or in neurons exposed to OGD/reoxygenation (Figure 6B). Interestingly, this effect was not affected by the treatment of neurons with siRNA against NCX1 (Figure 6C). To further support the hypothesis that NCX3 on the outer mitochondrial membrane participate to mitochondrial calcium handling during IPC, neurons were treated with CGP37157 during IPC and mitochondrial calcium concentrations was measured by confocal microscopy. The results of this experiments demonstrated that CGP37157 significantly affects NCX3 activity inducing an increase in mitochondrial calcium concentration (Figure 6C).



**B**



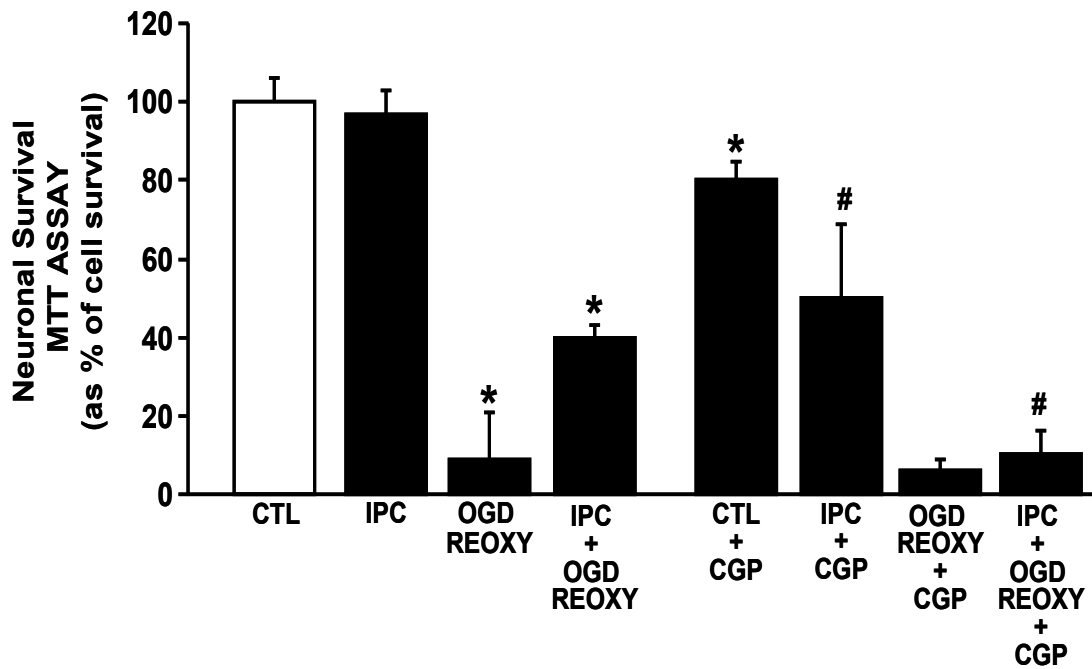
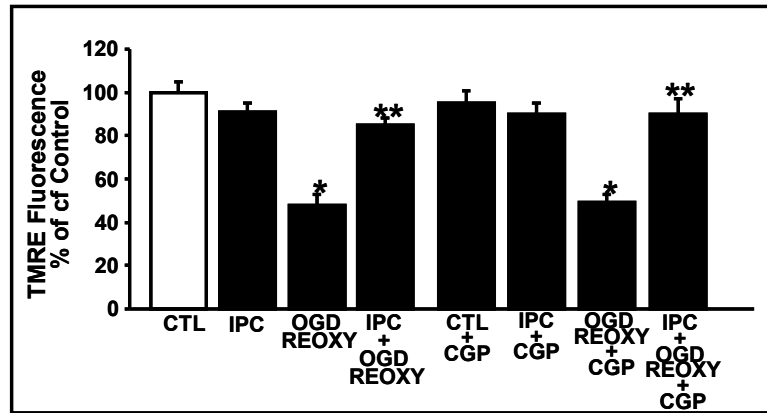
**C**



**Figure 6:  $[Ca^{2+}]_m$  in cortical neurons treated with siNCX3 and exposed to IPC/OGD followed by reoxygenation:** (A)  $[Ca^{2+}]_m$  evaluated by using confocal microscopy with the fluorescent probe X-Rhod1 in neurons (12 DIV) exposed to IPC, followed by 24 and 48 hrs of normoxia and to 3 hrs of OGD followed by 24 hrs of reoxygenation. (B)  $[Ca^{2+}]_m$  evaluated by using single cell microfluorimetry with the probe Fura-2AM after the treatment with FCCP in cortical neurons exposed to IPC, followed by 24 e 48 hrs of normoxia and then to 3 hrs of OGD followed by 48 hrs of reoxy.  $[Ca^{2+}]_m$  were measured at the end of the treatments. \* $P < 0.05$  Vs CTL; #  $P < 0.05$  Vs its not treated CTL. (C) effect of siNCX3 and CGP on  $[Ca^{2+}]_m$  in neurons exposed to IPC followed by 48 hrs of normoxia and treated with siNCX1 and CGP. \* $P < 0.05$  Vs CTL; #  $P < 0.05$  Vs IPC+48h reoxy.

***Mitochondrial function in cortical neurons treated with CGP and exposed to IPC followed by OGD/REOXYGENATION***

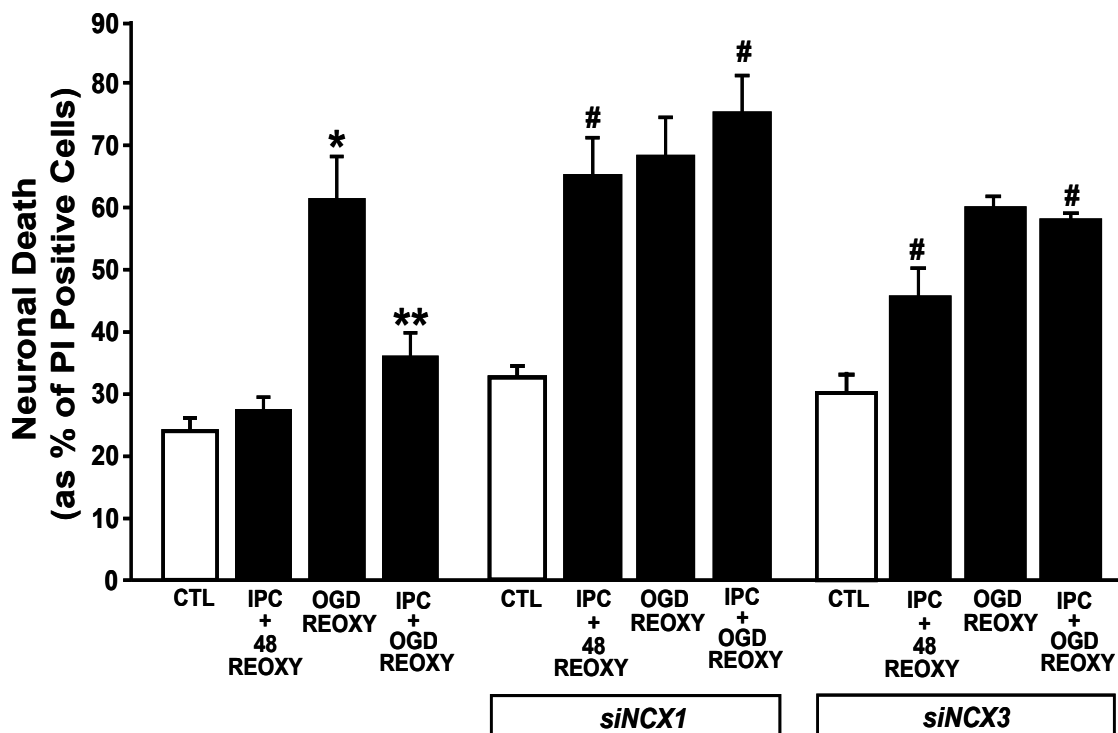
To understand whether the variations observed in mitochondrial calcium concentrations could be responsible for the neuroprotection induced by cerebral ischemic preconditioning, cell viability was assessed in terms of mitochondrial function after treatment with CGP, in cortical neurons exposed to IPC. The data showed a worsening in the mitochondrial function in neurons exposed to IPC insult after this treatment, whereas the sublethal insult, in the absence of CGP, did not affect mitochondrial function. Moreover, the significant improvement in mitochondrial activity observed in preconditioned neurons exposed to OGD followed by reoxygenation was abolished by treatment of cells with CGP (Figure 7). To rule out the possibility that CGP effect might be related to a modification of mitochondrial membrane potential and not to a real inhibition of mitochondrial exchanger, mitochondrial membrane potential was measured by using TMRE fluorescent probe in CGP-treated neurons exposed to IPC and IPC followed by OGD/Rexoygenation. The results demonstrated that CGP did not affect mitochondrial membrane (Figure 7, inset).



**Figure 7: Mitochondrial function in cortical neurons treated with CGP and to IPC followed by OGD/Reoxygenation:** cortical neurons exposed to PC followed by OGD/REOXY after the treatment with CGP. Mitochondrial function was measured by using MTT assay at the end of reoxygenation. \*P<0.05 Vs CTL e IPC; # P<0.05 Vs no CGP treatment. (7 Inset) Mitochondrial membrane potential in cortical neurons exposed to IPC/OGD reoxygenation after the treatment with CGP \*P<0.05 Vs CTL; \*\*P<0.05 Vs OGD reoxy.

***Effect of ischemic preconditioning on neuronal viability of cortical neurons treated with siRNA against NCX1 and NCX3 and exposed to OGD/REOXYGENATION***

To further support the hypothesis that both NCX1 and NCX3, by regulating ER and mitochondrial calcium content, may be among the effectors responsible for the tolerance induced by cerebral ischemic preconditioning, neuronal viability was evaluated after the silencing of the two Na<sup>+</sup>/Ca<sup>2+</sup> exchanger isoforms. The data obtained demonstrated that after these treatments there was a significant increase in cell death in cortical neurons exposed to the only IPC stimulus, whereas this sublethal insult did not affect neuronal survival but rather was able to protect neurons from death induced by OGD/Reoxygenation. Moreover, the treatment with siRNA against both NCX1 and NCX3 prevented the neuroprotective effect of ischemic preconditioning in cortical neurons exposed to a subsequent OGD followed by 24 hours of reoxygenation (Figure 8).



**Figure 8: Effect of IPC on neuronal survival of cortical neurons treated with siNCX1 and siNCX3 and exposed to OGD/Reoxy:** cortical neurons (12 DIV) were exposed to IPC (30 min di OGD) followed by 24 hrs of normoxia, and then to 3 hrs of OGD followed by 24 hrs of reoxygenation with and without the silencing against NCX1 and NCX3. Cell viability was measured by propidium iodide and fluorescein diacetate assay at the end of reoxygenation. \* $P < 0.05$  Vs CTL and IPC; \*\* $P < 0.05$  Vs OGD/REOXY; # $p < 0.05$  Vs cx not silenced.



## DISCUSSION

The results of the present study demonstrated that the two isoforms of the  $\text{Na}^+/\text{Ca}^{2+}$  exchanger, NCX1 and NCX3, which are involved in several pathophysiological aspects of cerebral ischemia, can be included among the members of the growing family of the mediators involved in the ischemic brain tolerance due to their ability to regulate neuronal calcium homeostasis. In particular, we proved evidence that neuroprotection observed in preconditioned neurons exposed to OGD/reoxygenation was correlated to the increase in NCX1 and NCX3 protein expression. Indeed, the treatment with siRNA against NCX1 and NCX3 was able to prevent this effect. These data are in accordance with results recently observed *in vivo* in an animal model of ischemic preconditioning (Pignataro et al., 2011). Consistently with these results we found that the upregulation of NCX1 and NCX3 protein expression in neurons exposed to IPC was dependent on PI3K/Akt activation, since the treatment with LY294002 was able to abolish this increase. Interestingly, in our experimental model, we demonstrated that NO plays a key role in the triggering PI3K/Akt pathway as the increase in the phosphorylated form of Akt observed within 30 minutes after IPC was completely abolished by the treatment with L-NAME. More importantly, the treatment with L-NAME was able to inhibit NCX3 but did not affect NCX1 protein expression. We recently demonstrated that in neurons NCX3, apart

its localization on the plasmamembrane, is also expressed on the outer mitochondrial membrane where it contributed to the extrusion of calcium from mitochondria (Scorziello et al., submitted paper). It is well known that mitochondria, in addition to the generation of cellular energy, play an important role in regulating cellular calcium homeostasis (Babcock et al., 1997; Budd and Nicholls, 1996; Werth and Thayer, 1994; Jouaville et al., 1995) in concert with the sarco-endoplasmic reticulum  $\text{Ca}^{2+}$ -ATPase, the plasma membrane  $\text{Ca}^{2+}$ -ATPase and  $\text{Na}^+/\text{Ca}^{2+}$  exchanger (Saric and Carafoli, 2005). On the other hand, the maintenance of mitochondrial calcium homeostasis is an important requirement ensuring mitochondrial function. In fact,  $\text{Ca}^{2+}$ -sensitive dehydrogenases can regulate oxidative phosphorylation and ATP synthesis during times of high cellular demand (McCormack and Denton, 1980). Therefore it is possible to hypothesize that the increased expression of NCX3, within 48hrs from the sub-lethal insult, might exert neuroprotective effects regulating calcium handling and improving mitochondrial oxidative capacity. This finding is in line with the results previously obtained in an *in vitro* model of IPC and demonstrating that the increase in nNOS expression and  $\text{NO}^{\bullet}$  production through the activation of Ras/ERK1/2 pathway stimulated mitochondrial MnSOD. These effects were associated with a reduction in free radical production, cytochrome c release from mitochondria to cytosol, and in turn to an improvement of neuronal survival (Scorziello et al., 2007). On the basis of the tight correlation among

mitochondrial calcium content, oxidative metabolism and neuronal survival, the results of the present study add new insight to the molecular mechanisms involved in the ischemic brain preconditioning. Indeed, the over-expression of NCX3 might help mitochondria to preserve their energetic capacity making them less vulnerable to the subsequent lethal insult represented by OGD/reoxygenation. This hypothesis was strongly supported by the results obtained measuring the activity of NCX during IPC. In fact, we demonstrated that IPC induced an increase of NCX activity in the reverse mode of operation that was still observed in preconditioned neurons exposed to OGD/Reoxygenation. This is an early event since it occurred within 30 minutes after IPC stimulus and is due to the contribution of NCX1 and NCX3 isoforms and was promoted by NO because was abolished by the treatment with L-NAME. Intriguingly, this effect was associated with an increase in ER calcium content. Consistently with data previously published (Sirabella et al., 2009), the experiments of the present study suggested that NCX1, working in the reverse mode of operation, play a key role in the regulation of ER calcium refilling in the early phase of IPC. Indeed, the treatment with siNCX1, but not siNCX3, was able to prevent this phenomenon. The novel aspect of the study is the demonstration that NO promoted NCX1-induced ER refilling that was hampered by L-NAME pretreatment. This finding is in accordance with data recently published by Secondo et al., and demonstrating that NO was able to stimulate NCX1 to work in the reverse mode of operation, whereas NO did

not affect NCX3 activity (Secondo et al. 2011). The possibility that NCX1 activation in the early phase of IPC could affect mitochondrial calcium content promoting mitochondrial calcium uptake could not be excluded. However, in the late phase of IPC mitochondrial calcium handling is mainly regulated by NCX3 that is able to promote mitochondrial calcium extrusion. This data is supported by the finding that NCX3 expression increased 48 hrs after the IPC insult. We have previously demonstrated that NCX3 is distributed also at mitochondrial level, therefore, it is possible to speculate that during IPC the increased expression of NCX3 on mitochondria might contribute to the efflux of calcium from mitochondria thus protecting neurons by the subsequent lethal OGD/reoxygenation. The finding that the treatment with CGP and siNCX3 counteracted the effect of IPC on mitochondrial calcium content leading to the lack of IPC-neuroprotection further supports this hypothesis.

Collectively, we proposed a model (Figure 9) in which a functional interplay between NCX1 and NCX3 occurs during IPC. This phenomenon is tightly dependent on NO and Akt activation and, by contributing to the modulation of intracellular ionic homeostasis, could represent the mechanism responsible for neuroprotection induced by ischemic preconditioning.

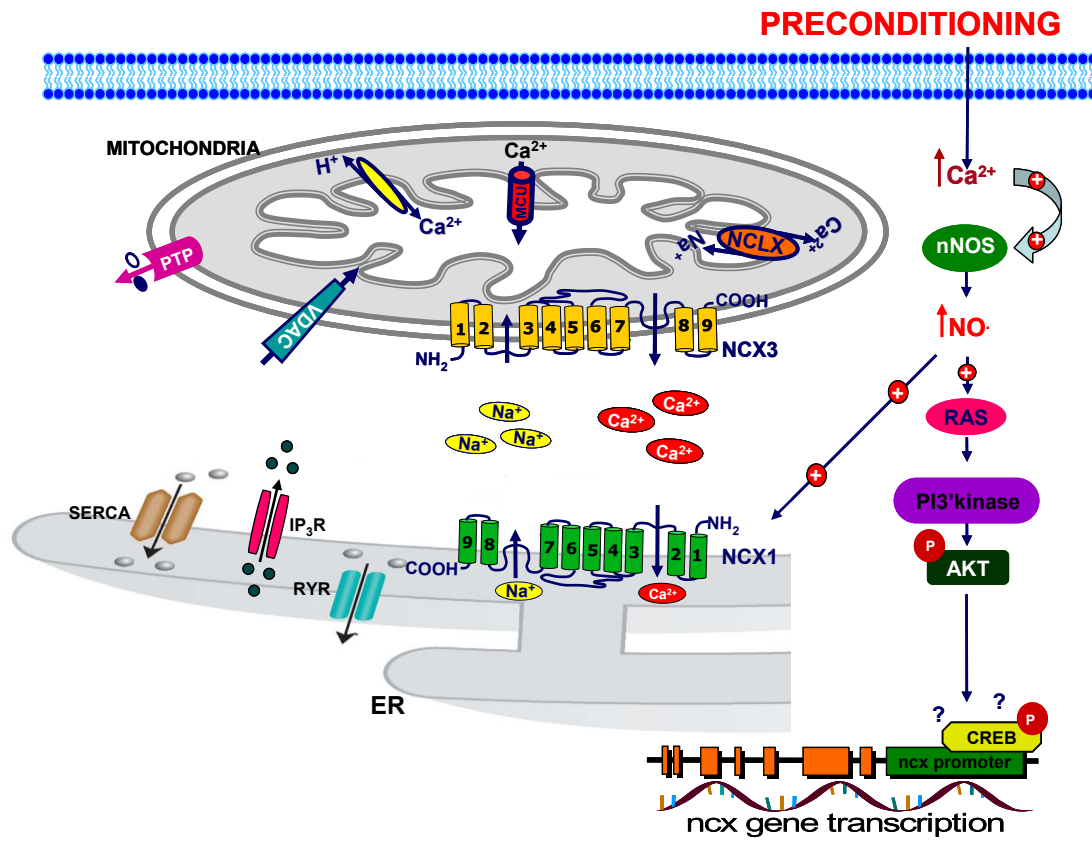


Figure 9: Proposed model

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