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The acetylation of ReIA in Lys310 dictates the NF-*κ*B-dependent response in post-ischemic injury

A Lanzillotta¹, I Sarnico¹, R Ingrassia³, F Boroni¹, C Branca¹, M Benarese¹, G Faraco⁴, F Blasi⁴, A Chiarugi⁴, P Spano^{1,2} and M Pizzi^{*,1,2}

The activation of nuclear factor kappa B (NF- κ B) p50/RelA is a key event in ischemic neuronal injury, as well as in brain ischemic tolerance. We tested whether epigenetic mechanisms affecting the acetylation state of RelA might discriminate between neuroprotective and neurotoxic activation of NF- κ B during ischemia. NF- κ B activation and RelA acetylation were investigated in cortices of mice subjected to preconditioning brain ischemia or lethal middle cerebral artery occlusion (MCAO) and primary cortical neurons exposed to preconditioning or lethal oxygen-glucose deprivation (OGD). In mice subjected to MCAO and in cortical neurons exposed to lethal OGD, activated RelA displayed a high level of Lys310 acetylation in spite of reduced total acetylation. Also, acetylated RelA on Lys310 interacted strongly with the CREB-binding protein (CBP). Conversely, RelA activated during preconditioning ischemia appeared deacetylated on Lys310. Overexpressing RelA increased Bim promoter activity and neuronal cell death both induced by lethal OGD, whereas overexpressing the acetylation-resistant RelA-K310R, carrying a mutation from Lys310 to arginine, prevented both responses. Pharmacological manipulation of Lys310 acetylation by the sirtuin 1 activator resveratrol repressed the activity of the Bim promoter and reduced the neuronal cell loss. We conclude that the acetylation of RelA in Lys310 dictates NF- κ B-dependent pro-apoptotic responses and represents a suitable target to dissect pathological from neuroprotective NF- κ B activation in brain ischemia.

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Brain ischemia is a leading cause of mortality and neurological disability. During brain ischemia, nuclear factor kappa B $(NF-\kappa B)$ rapidly activates in neurons and glial cells, where it regulates inflammatory and apoptotic events characterizing the pathophysiology of post-ischemic injury.¹⁻⁴ NF-κB activation in neurons, rather than in glial cells, has a prominent role in post-ischemic cell loss.^{4,5} NF-κB is a dimeric transcription factor created by the association of different subunits: p50. p65/ReIA, p52, ReIB and c-ReI. The most prevalent activated dimer observed after occlusion of the middle cerebral artery (MCAO) is p50/RelA.^{6,7} Accordingly, the MCAO-induced infarct size is reduced in p50 knockout mice,² as well as in mice carrying a brain-conditional deletion of RelA.7 We showed that activation of the p50/RelA dimer is associated with inhibition of c-Rel-containing dimers in ischemic brain tissues and in primary cortical neurons exposed to oxygenglucose deprivation (OGD).8 Selective targeting of c-Rel and RelA revealed that activation of c-Rel-containing dimers increases cell resistance to OGD, whereas activation of p50/ReIA contributes to the cell-death program. The effect of c-Rel dimers relies on promoter activation of the anti-apoptotic Bcl-xL gene⁸ and mediates neuroprotection induced by leptin⁹ or agonists at metabotropic glutamate receptor type 5.¹⁰ The deleterious effect of p50/RelA in brain ischemia depends on RelA-induced expression of pro-apoptotic Bim and Noxa genes.^{7,8} However, p50/ReIA is also involved in the regulation of a variety of physiological processes. Its constitutive activity is required for brain neuron survival and neurite elongation during brain development.^{11,12} In mature neurons, p50/RelA selectively localizes at the synaptic level, from which point it moves to the nucleus to transmute synaptic signals into altered gene expression, regulation of synaptic plasticity and memory formation.^{13–15} Furthermore, the activation of p50/ RelA has a role in brain tolerance, the adaptive response induced by a sublethal insult, which preserves brain health against subsequent lethal injury.¹⁶ Thus, the opposing effects elicited by NF- κ B activation in cell survival remain to be elucidated. Recent studies in tumor and peripheral cells have revealed the post-translational regulation of ReIA, including reversible phosphorylation and acetylation,¹⁷ which modulate p50/RelA transcriptional activity on target genes.¹⁸ RelA is acetylated after cell activation by tumor necrosis factor- α , phorbol myristate acetate or other stimuli. RelA acetylation may occur at multiple sites, including lysines (Lys) 122, 123,

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¹Division of Pharmacology and Experimental Therapeutics, Department of Biomedical Sciences & Biotechnologies, School of Medicine, University of Brescia and Istituto Nazionale di Neuroscienze Brescia, Brescia, Italy; ²Department of Obstetrics and Gynecology and Biomedical Technologies, University of Brescia, Brescia, Italy; ³Department of Pharmacology, University of Florence, Florence, Italy and ⁴Istituto Ricovero e Cura a Carattere Scientifico, S. Camillo Hospital, Venice, Italy *Corresponding author: M Pizzi, Division of Pharmacology, Department of Biomedical Sciences & Biotechnologies, Viale Europa, 11, Brescia 25123, Italy. Tel: + 39 030 371 7501; Fax: + 39 030 371 7529; E-mail: pizzi@med.unibs.it

Abbreviations: BCCAO, bilateral common carotid arteries occluded; CBP, p300/CREB-binding protein; DIV, days *in vitro*; FOXO3a, Forkhead transcription factors; IPC, ischemic preconditioning; JNK, c-Jun N-terminal protein kinase; LDH, lactate dehydrogenase; Lys, lysine; MCAO, occlusion of middle cerebral artery; NF-*k*B, nuclear factor kappa B; OGD, oxygen-glucose deprivation; RA, retinoic acid; SIRT1, sirtuin 1

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218, 221 and 310.17 The epigenetic regulators of histone proteins, acetyltransferases p300/CREB-binding protein (CBP) and p/CAF, appear to have a major role in the in vivo acetylation of ReIA.¹⁹ Site-specific acetylation of ReIA regulates discrete biological actions of the NF- κ B complex. For example, acetylation of Lys218 and 221 increases the DNA binding affinity of RelA for the κB enhancer and impairs RelA assembly with newly synthesized $I\kappa B\alpha$,¹⁹ whereas acetylation of ReIA at Lys122 and 123 inhibits its transcriptional activity.¹⁷ Acetylation of Lys310 does not modulate DNA binding or $I\kappa B\alpha$ assembly, but markedly enhances the NF- κB transactivation of pro-inflammatory genes. Acetylation of Lys310 is required for the full transcriptional activity of ReIA. Abolishing Lys310 acetylation by mutating Lys310 to arginine significantly inhibits the transactivation of NF-kB and the expression of inflammatory cytokines, 19,20 possibly by stabilizing Set-9 factor, which leads to the methylation of Lys314-315 and proteasomal degradation of ReIA.21 Selective deacetylation of Lys310 by sirtuin 1 (SIRT1), a class III histone deacetylase, inhibits the transcriptional activity of RelA²² and prevents the β -amyloid-induced release of neurotoxic factors from microglial cells.²³ However, the role of ReIA acetylation in NF- κ B-mediated neuronal injury during brain ischemia remains elusive.

We investigated changes in c-Rel activation and RelA acetylation in response to preconditioning or lethal ischemia and the role of these changes in ischemia-mediated gene transcription and neuronal cell death.

Results

Neuronal activation of NF-kB p50/RelA in preconditioning and lethal ischemia. It has been shown that as an early response to ischemic brain injury, NF-kB p50 and RelA rapidly activate and drive post-ischemic neuronal apoptosis. ^{4,5,7,8} We previously showed that in primary mouse cortical neurons exposed to OGD apoptosis precedes necrosis, as indicated by the early TUNEL-positivity displayed by the cells within 6 h after the OGD and the parallel release of cytochrome c in the cytosol in absence of lactate dehydrogenase (LDH) release. Subsequent necrosis causes progressive elevation of extracellular LDH level that becomes clearly detectable in the medium 24 h after the OGD exposure.⁸ Thus, we here measured delayed LDH level as a final marker of cell death. Either in primary cortical neurons or in mouse ischemic cortices, enhanced activation of p50/ReIA correlates with the diminished activation of ReIA/ c-Rel, an event that contributes to neuronal vulnerability. In order to analyze and compare the NF-kB dimers induced by preconditioning or lethal OGD, we set up an ischemic preconditioning (IPC) model by exposing the cortical neurons to 1 h OGD and, the next day, to 3 h OGD. The 1 h OGD condition did not affect cell viability per se and completely abolished the cell death induced by longer OGD applied 24 h later (Figure 1a). Enzyme-linked immunosorbent assay (ELISA) analysis of NF-kB showed similar increases in ReIA and p50 DNA-binding activity in nuclear extracts of



Figure 1 (a) Primary cortical neurons were exposed to 1 h OGD and then 24 h later to 3 h OGD. The next day, cell viability was measured by LDH assay. Sub-lethal ischemic injury totally prevented the 3 h OGD-mediated neurotoxicity. Bars are means \pm S.E.M. of three separate experiments run in triplicate; **P*<0.01 versus the corresponding control value. (b) Activation of p50 and RelA was evaluated by ELISA analysis in nuclear extracts from cortical cells exposed to 1 or 3 h OGD. Bars are means \pm S.E.M. of three separate experiments; **P*<0.05 versus the corresponding control value. (c) Representative picture of co-immunoprecipitation analysis of p50, RelA and c-Rel dimers in nuclear extracts from primary cortical neurons showed a high level of p50/RelA complex activation after 1 or 3 h OGD, whereas the activation of c-Rel dimers decreased slightly

cells exposed to 1 and 3h OGD (Figure 1b). The coimmunoprecipitation analysis of heterodimers composed of p50, RelA and c-Rel revealed superimposable increases in levels of p50/RelA under both conditions. No increase was evident in the levels of p50/c-Rel and RelA/c-Rel complexes (Figure 1c). These results indicate that p50/RelA, but not c-Rel dimers contribute to NF- κ B activation in cells exposed to preconditioning, as well as lethal OGD.

ReIA acetylation and interaction with CBP differentiate p50/RelA activation in neuronal cells during lethal as opposed to preconditioning ischemia. We checked the possibility of diverse RelA acetylation in two ischemic conditions, 1 and 3h OGD. RelA was immunoprecipitated from nuclear extracts and its acetylation was checked using a specific anti-Acetyl-RelA Lys310 antibody (RelA Ac-K310) in comparison with an anti-Acetyl-Lys antibody recognizing general RelA acetylation. We observed that acetylation at Lys310 was reduced after 1h OGD but increased significantly after 3h OGD, in spite of reduced levels of overall RelA acetvlation (Figure 2a and b). The diverse acetylation states of ReIA correlated with diverse levels of interaction with the histone acetyl transferase CBP. RelA showed higher association with CBP after lethal OGD, but not after preconditioning ODG (Figure 2a and b). Similar



Figure 2 (a) Immunoprecipitation analysis of ReIA acetylation and association with CBP in nuclear extracts of primary cortical neurons exposed to 1 or 3 h OGD. ReIA Lys310 acetylation decreased after 1 h OGD and increased after 3 h OGD. Total ReIA acetylation was not altered by 1 h OGD but was reduced by 3 h OGD. RelA association with CBP increased in nuclear extracts of cells subjected to 3 h OGD. The signal given by IgG(H) was used as a control for the quality of the immunoprecipitation. Similar results were obtained in at least four separate experiments. (b) Values from densitometric analysis of immunoblot bands are expressed as a percentage of the corresponding control value. Error bars depict means \pm S.E.M.; *P < 0.05 versus control

results were obtained in an in vivo mouse model of brain ischemia and IPC.²⁴ Mice were exposed to diverse ischemic insults: (i) 5 min bilateral common carotid arteries were occluded (BCCAO) (preconditioning ischemia); (ii) 20 min MCAO; and (iii) IPC, that is, BCCAO followed by MCAO the next day. Ischemic areas (Figure 3a) and infarct volumes (Figure 3b) were evaluated 3 days later. BCCAO induced no brain damage, but it significantly reduced the ischemic volume in mice subjected to MCAO (Figure 3b). We tested RelA acetylation in cortical nuclear extracts prepared 4h after mice were exposed to the various conditions (Figure 3c). Higher RelA activation was evident in BCCAO. MCAO and IPC extracts, but acetylation at Lys310 increased only in mice exposed to MCAO. Concomitantly with Lys310 acetylation, ReIA also displayed higher interaction with CBP in the MCAO extracts (Figure 3c and d). These results show that ReIA acetylation on Lys310 discriminates between ReIA activation after neurotoxic ischemia and that induced by preconditioning ischemia.

Acetylation of ReIA on Lys 310 is required for toxic effects elicited by NF-kB activation. The relevance of RelA acetylation on Lys310 in the NF-kB-mediated neurotoxicity after acute ischemic injury was initially investigated in neuronally differentiated SK-N-SH cells. Under exposure to retinoic acid (RA), SK-N-SH neuroblastoma cells switch from a non-neuronal to a neuronal-like phenotype²⁵ and express vulnerability to OGD by NF-kB RelA activation.⁸ Neuronal cells were exposed to OGD for 1, 2 and 4h. Western blot analysis of nuclear extracts confirmed increased acetylation of RelA on Lys310 after 4h ODG, as observed in primary cortical neurons (Figure 4a). The neuronal cultures were transiently transfected with control empty vector (pSG5) or with expression plasmids coding for human wild-type RelA or RelA carrying a mutation from Lys310 to arginine (ReIA-K310R). As arginine has the same polar side chain and charge as lysine but cannot be acetylated, this mutated form of RelA was used as a negative control to test the biological effects of specific Lys310 acetylation.¹⁹ Around 24 h after transfection, the nuclear translocation of ReIA and its acetylation were verified by immunoblot analysis in nuclear extracts of cells exposed to 4h OGD. We found that OGD promoted a marked upregulation of nuclear ReIA in cells transfected with empty vector, as well as in cells overexpressing wild-type ReIA or ReIA-K310R (Figure 4b and c). As expected, the concomitant increase in acetvlation on Lys310 after OGD was evident in control and RelAoverexpressing cells, but negligible in the ReIA-K310Rtransfected cells (Figure 4b and c). In order to test the possibility that this unique modification was specifically responsible for the regulation of neuronal injury, transfected cells were exposed to OGD for 15 h and cell viability was measured by LDH release in the medium 24 h later. Cells transfected with the empty vector were vulnerable to OGD and the overexpression of wild-type ReIA significantly increased cell death (Figure 4d), in line with previous findings.8 Instead, the RelA-induced enhancement of OGD-mediated neurotoxicity was completely abolished in cells overexpressing the ReIA-K310R mutant construct. Similar experimental conditions were used to analyze the



Figure 3 (a) Representative images of infarct areas in brain coronal sections of mice exposed to BCCAO for 5 min or to MCAO for 20 min. In preconditioning experiments (IPC), mice were subjected to BCCAO and the next day to MCAO. Ischemic lesions were evaluated 3 days later. (b) Infarct volume in mice subjected to BCCAO, MCAO or IPC. Data are reported as means \pm S.E.M. (n = 9 or 10 animals per group); *P < 0.05 versus MCAO value. (c) Representative picture of co-immunoprecipitation analysis of ReIA acetylation in nuclear extracts of mice exposed to BCCAO, MCAO or IPC (n = 3 per group). Nuclear extracts were prepared 4 h after the end of each experimental condition. Acetylation on the Lys310 residue, as well as levels of the ReIA/CBP complex, increased in mice exposed to MCAO. (d) Densitometric analysis of immunoblot bands. Values are expressed as a percentage of the control (Sham) value. Error bars depict means \pm S.E.M.; *P < 0.05 versus control

NF-*κ*B-dependent activation of mouse Bim promoterluciferase plasmid in cortical neurons during 3h OGD followed by 4h recovery, a time during which neurons express Bim and undergo apoptosis.^{7,8} We previously showed that OGD induces the activity of the Bim promoter but not the activity of BimΔ*κ*B plasmid carrying a mutation at the *κ*B site, demonstrating that Bim transcription during OGD is NF-*κ*B specific.⁸ The OGD-induced Bim promoter was significantly enhanced by ReIA overexpression, but fell to basal levels in cells overexpressing ReIA-K310R (Figure 4e). These results show that NF-*κ*B-mediated Bim transcription is completely dependent on Lys310 acetylation of ReIA. This mechanism is pivotal in driving the deleterious effects of NF*κ*B activation during lethal ischemia.

Resveratrol-mediated neuroprotection is associated with inhibition of RelA acetylation at the Lys310 residue as well as with the inhibition of Bim promoter activity. Deacetylation of Lys310 by the class III histone deacetylase SIRT1 inhibits the transcriptional activity of RelA.²² To determine whether pharmacological manipulation of RelA acetylation could repress NF- κ B-mediated pro-apoptotic transcription during OGD, we tested the SIRT1-activating molecule resveratrol. Cortical neurons were exposed to OGD for 3 h and then treated for 24 h with resveratrol (1, 3 and 30 μ M). Even when added during the post-ischemic period, resveratrol showed significant neuroprotective activity at a concentration of $30 \,\mu$ M (Figure 5a). Immunoprecipitation analysis of ReIA in nuclear extracts confirmed that after OGD, the acetylation of Lys310 increased and total acetylation decreased. Treatment with $30 \,\mu$ M resveratrol for 2 h after OGD only partially reduced ReIA activation, weakly affected general ReIA acetylation and completely deacetylated the Lys310 residue (Figure 5b). In line with this drastic effect on Lys310 acetylation, resveratrol repressed the Bim promoter activity to levels below the baseline (Figure 5c). These results represent further evidence that during ischemia, the neurotoxic activation of NF- κ B is associated with ReIA acetylation at Lys310. Drugs that deacetylate ReIA at Lys310, similar to the actions of resveratrol, can block post-ischemic transcription and neurodegeneration.

Discussion

Here, we report that although activation of NF- κ B c-Rel dimers promotes neuronal resistance to environmental noxae,^{8–10} NF- κ B activation following preconditioning OGD does not involve c-Rel-containing dimers. Moreover, the p50/RelA complex is activated in preconditioning, as well as in lethal OGD. The aberrant activation of p50/RelA dimer in neuronal cells, responsible for pro-apoptotic events in the postischemic period,^{6–8} is characterized by RelA acetylation at the Lys310 residue. It is unknown whether the acetylation of



Figure 4 (a) Immunoblot analysis of nuclear extracts from differentiated SK-N-SH cells exposed to 1–4 h OGD. RelA Lys310 acetylation significantly increased after 4 h OGD. (b) Neuronal SK-N-SH cells were transfected with wild-type RelA and RelA-K310R plasmids or with a pSG5 empty vector for 24 h and then exposed to OGD for 4 h. The OGD-induced RelA acetylation on Lys310 was significantly reduced in RelA-K310R expressing cells. (c) Data from densitometric analysis of RelA and RelA Ac-K310 immunoblots are expressed as ratios to relative nucleolin levels. Bars are the means \pm S.E.M. of three separate experiments; **P*<0.05 *versus* corresponding control value. (d) Cell survival was measured in SK-N-SH cells exposed to 15 h OGD. RelA overexpression significantly enhanced OGD toxicity, whereas RelA Ac-K310R overexpression prevented cell loss. Bars are the means \pm S.E.M. of three separate experiments; **P*<0.01 *versus* OGD in pSG5-expressing cells. (e) Primary cortical neurons were transfected with *Bim* luciferase reporter plasmid together with wild-type RelA or RelA-K310R plasmids or pSG5 empty vector for 24 h and then exposed to 3 h OGD. Luciferase activity was measured after 4 h. RelA overexpression enhanced OGD-induced *Bim* promoter activity, whereas RelA-K310R overexpression totally inhibited such activity. Bars are the means \pm S.E.M. of three experiments run in triplicate; **P*<0.01, ***P*<0.05 *versus* OGD value in pSG5-expressing neurons

other lysine residues is also involved, though the increase of Lys310 acetylation occurred in spite of decreased total RelA acetvlation. Moreover, acetvlation of RelA on Lvs310 differentiated the p50/RelA activation during lethal ischemia from that induced by preconditioning ischemia. The level of acetylation in Lys310 during OGD paralleled the increased interaction of ReIA with CBP, in line with evidence that ReIA acetylation on Lys310 is strictly dependent on the histone acetyltransferase activity of CBP/p300.¹⁹ Similar results were obtained in a model of in vivo brain ischemia and IPC. RelA is highly activated in cortices of mice exposed to either BCCAO or MCAO, as previously shown.¹⁶ Notably, the nuclear ReIA level was high also in mice exposed to 1-day IPC, that is, BCCAO followed by MCAO on the next day. This result is in contrast to what was observed in models of 3-day IPC, in which severe ischemia induced 3 days after sublethal ischemia elicited weaker NF- κ B activation as a consequence of enhanced I κ B α expression during the 3-day interval.¹⁶ Our data support the idea that brain tolerance does not impair NF- κ B translocation, but does affect ReIA acetylation and transactivation. Thus, compared with BCCAO or IPC mice, the MCAO mice displayed a higher level of Lys310 acetylation that correlated with increased interaction of ReIA with CBP.

To elucidate the relevance of Lys310 acetylation to RelA-mediated effects during ischemia, we transfected the neuronally differentiated SK-N-SH cells that were displaying RelA-dependent vulnerability to OGD⁸ with the RelA-K310R mutant construct. We found that OGD greatly enhanced Lys310 acetylation in control cells, as well as in cells transfected with RelA, whereas acetylation did not increase in cells overexpressing RelA-K310R. This result confirmed that the mutation of lysine to arginine in the RelA sequence



Figure 5 (a) Cell death was evaluated in cortical neurons exposed to 3 h OGD and then treated with 1, 3, 30 μ M resveratrol for 24 h. Resveratrol produced a significant neuroprotective effect at the 30 μ M concentration. Bars are the means ± S.E.M. of three separate experiments run in quadruplicate; **P*<0.05 *versus* the corresponding control value. (b) Representative picture of co-immunoprecipitation analysis of RelA in nuclear extracts of primary cortical neurons exposed to 3 h OGD and to 30 μ M resveratrol for 2 h during the post-OGD period. (c) Luciferase activity in cortical neurons co-transfected with Bim luciferase reporter plasmid and control vector (pSG5) or RelA-K310R expression plasmid. OGD stimulated luciferase activity in pSG5 but not in RelA-K310R-expressing cells. Resveratrol (30 μ M), added after the OGD and maintained for 4 h, repressed luciferase activity in pSG5-cells. Bars are the means ± S.E.M. of three separate experiments; **P*<0.01, ***P*<0.05 *versus* OGD value in pSG5-expressing neurons

impairs acetylation of the RelA 310 residue.¹⁹ Thus, OGDmediated cell death was enhanced in RelA-overexpressing cells,⁸ but was completely impaired in RelA-K310R-transfected cultures. The occurrence of cell death at levels even lower than those observed in cells expressing empty vector suggests that deacetylated RelA-K310R can compete with native Acetyl-RelA Lys310 to maximally reduce neuronal vulnerability, as observed in the preconditioned cells.

NF-kB fine-tunes cell survival by regulating transcription of Bcl2 family genes endowed with either anti-apoptotic activity, as the Bcl-xL, or pro-apoptotic activity, as the BH3-only members Bim and Noxa.²⁶ It has been shown that Bim and Noxa genes, being under the transcriptional control of ReIA, are significantly induced 6 h after MCAO and their upregulation is abolished in mice carrying a brain-conditional deletion of ReIA.7 Evidence that cortical neurons exposed to lethal OGD display NF-kB-dependent induction of Bim, but not Bcl-xL promoter, highlights the transcriptional specificity of p50/RelA activation during anoxic injury.8 Bim, similar to other BH3-only proteins, contributes to increase the mitochondrial permeability and activation of caspase cascade by directly activating pro-apoptotic Bax and Bak²⁷ or by releasing Bax and Bak from their complexes with pro-survival Bcl-2 homologs.^{28,29} Bim represents a converging point of diverse pro-apoptotic pathways during brain ischemia. In addition to

NF-kB, Bim can be transcriptionally induced by Forkhead transcription factors (FOXO)3a as downstream target of the PTEN-Akt-FOXO3a pathway activated in ischemic brains.³⁰ The interaction of Bim with the c-Jun N-terminal protein kinase (JNK), enhances Bim affinity to Bax and increases both proteins translocation to the mitochondria.^{31,32} Intriguingly, JNK and FOXO3a signaling can be negatively modulated by the crosstalk with NF- κ B in pro-survival pathways^{33,34} raising additional queries about the mode of Bim regulation by NF-*k*B in brain ischemia. Here, we show that Bim transcription during OGD strictly depends on Lys310 acetylation of ReIA. Bim transcription increased during OGD, it was enhanced in cortical neurons overexpressing ReIA, but was completely inhibited in ReIA-K310R-expressing cells. This result suggests that ReIA acetylation on Lys310, through the recruitment of the coactivator CBP/p300, is a mechanism evolved to regulate inducible pro-apoptotic genes during ischemia in neuronal cells. Notably, the transactivation potential of c-Rel was reported not to be influenced by the CBP/p300 interaction.³⁵ This difference may also account for diverse regulation of NF- κ B target genes by c-Rel and RelA during ischemia.8 The specific RelA acetylation on Lys310 represents a molecular target to modify gene transcription and neuronal resilience to ischemic injury.

Yeung and colleagues²² originally demonstrated that the histone deacetylase SIRT1 can interact with ReIA to inhibit

gene transcription by deacetylating ReIA at Lys310, without modifying the other lysine residue. These results suggested that acetylated Lys310 might form a platform for the binding of a bromodomain-containing protein that is required for full transcriptional activity of ReIA.³⁶ Interestingly, SIRT1 protects against neurodegeneration, and SIRT1 levels rise rapidly in various neurotoxic and neurodegenerative conditions; these results suggest SIRT1 may represent a stress sensor molecule that is important for the neuroprotective adaptation response.³⁷ We checked the effects induced in cortical neurons by resveratrol, a pharmacological activator of SIRT1.²² In line with previous evidence.^{37,38} resveratrol protected neuronal cells in a concentration-dependent manner, even if added after exposure to OGD. This effect correlated with the capability of the compound to abolish Lys310 acetylation, in spite of a modest inhibition of RelA nuclear translocation and a minor reduction in total RelA acetylation. As a consequence of specific RelA deacetylation, resveratrol repressed the acetyl-RelA Lys310-dependent transcription of Bim promoter during OGD. We cannot rule out the fact that additional targets of SIRT1 deacetylase activity, including FOXO3, p53 and peroxisome proliferator, activated receptor gamma co-activator 1α (PGC- 1α) transcription factors, ^{37,39} as well as mechanism other than SIRT1 activation³⁸ may contribute to the neuroprotection by resveratrol. Nonetheless, the present data are consistent with the hypothesis that the use of pharmacological agents modulating SIRT1 activity affects the acetylation status of RelA protein at Lys310, as well as its pro-apoptotic transactivation potential.

The issue of ReIA acetylation has been thought to explain the pro-inflammatory activity of p50/ReIA in immune cells,^{40,41} reactive astrocytes⁴² and microglial cells.²³ Here, we demonstrate that in neuronal cells does the acetylation of ReIA at Lys310 function as an intranuclear molecular switch that discriminates between the neurotoxic and the neuroprotective NF- κ B pathway in brain ischemia. The acetylation of ReIA at Lys310 may represent a new drugable target to reproduce the preconditioning-induced activation of NF- κ B.

Materials and Methods Cell culture

Primary cultures of mouse cortical neurons. Cortical neurons were prepared from 15-day-old embryonic mice, harvested with cesarean section from anaesthetized pregnant C57Bl/6 dams (Charles River, Italy) and cultured as previously described.⁸ Experiments were carried out at 11 days *in vitro* (DIV). *SK-N-SH cell culture.* The human SK-N-SH neuroblastoma cell line was purchased from American Type Culture Collection (Rockville, MD, USA). Cells were grown at 37°C in a humidified atmosphere of 5% CO₂, 95% O₂ in Dulbecco's modified Eagle's medium (DMEM) (Euroclone, Milan, Italy) supplemented with fetal calf serum, 4 mM glutamine and 100 U/ml penicillin/streptomicin. The addition of 50 μ M RA (Sigma, St. Louis, MO, USA) for 10–12 days induced mitotic arrest and differentiation into a neuronal-like phenotype.²⁵

Cerebral ischemia models

Transient middle cerebral artery occlusion (MCAO). Procedures involving animals were approved by the Institutional Animal Care Committee in compliance with the Italian guidelines for animal care (DL 116/92) and the European Communities Council Directive (86/609/EEC). C57BI/6 male mice (Harlan, Milan, Italy) were exposed to IPC and/or transient (20 min) MCAO as previously described.²⁴ For IPC, mice (n = 9) were anesthetized, and bilateral common carotid arteries were occluded (BCCAO) for 5 min with microclips. After 24 h, mice

underwent 20 min MCAO as reported.²⁴ Parallel groups were subjected to BCCAO (n = 9) or MCAO (n = 9) alone. Infarcts were measured 3 days later to rule out transient neuroprotection. Examination of infarct volume was performed in brains frozen in liquid nitrogen to avoid post-mortem changes. To prepare nuclear extracts, mice were killed by decapitation 4 h after BCCAO (n = 3), MCAO (n = 3) or BCCAO and MCAO (IPC) (n = 3).

OGD. Primary cortical neurons at 11 DIV were exposed to OGD as previously described,⁸ for 1 or 3 h. Cells recovered for 24 h in culture medium and were typically aerated in the incubator for the evaluation of cell viability. Resveratrol (1, 3 and 30 μ M) (Calbiochem, Beeston Nottingham, UK) was added in the post-ischemic period. Nuclear proteins were extracted at the end of OGD or after an additional 2 h incubation with resveratrol. SK-N-SH neuronal cells were exposed to 15 h OGD and replaced in fresh DMEM without serum for 24 h for analysis of cell viability. Neuronal injury was evaluated by measuring the amount of LDH (Promega, Madison, WI, USA) released relative to total releasable LDH.⁸

Co-immunoprecipitation and western blot analyses. Nuclear protein extracts were prepared as previously described⁸ from primary cortical neurons and from differentiated SK-N-SH cells, immediately after OGD exposure or after an additional time as indicated, or from the cortices of mice exposed to 5 min BCCAO followed by 4 h reperfusion, 20 min of MCAO followed by 4 h reperfusion, 5 min BCCAO and 24 h later, 20 min MCAO (IPC) and 4 h reperfusion.

Co-immunoprecipitation studies and immunoblot analyses were carried out as previously described.⁸ In total, 20 µg of nuclear extracts were incubated at 4°C overnight with 2 µg/ml of goat polyclonal anti-RelA antibody (sc-372G, Santa Cruz Biotechnology, Santa Cruz, CA, USA) and co-immunoprecipitated proteins were detected by immunoblotting using the following antibodies: rabbit polyclonal anti-p50 (1:500, ab7971 Abcam, Cambridge, UK), rabbit polyclonal anti-c-Rel (1:50, sc-71X Santa Cruz Biotechnology), rabbit polyclonal anti-RelA (1:200, sc-372, Santa Cruz Biotechnology), rabbit polyclonal anti-Acetyl-RelA (Lvs310) (1:500, #3045, Cell Signaling, Danvers, MA, USA), rabbit polyclonal anti-Acetyl-Lys (1:500, #06-933 Upstate-Millipore, Billerica, MA, USA) and rabbit polyclonal anti-CBP (1: 500, sc-583 Santa Cruz Biotechnology). For immunoblot analyses, nuclear proteins (25 µg proteins/sample) were resolved by 10% SDS/polyacrylamide gel. Immunodetection was performed by incubating the membrane overnight at 4°C, with the following primary antibodies: rabbit polyclonal anti-Acetyl-RelA (Lys310) (1:500, #3045 Cell Signaling), rabbit polyclonal anti-ReIA (1:200, sc-372 Santa Cruz Biotechnology), rabbit polyclonal anti-histone H3 (1:1000, #9715 Cell Signaling) and rabbit polyclonal anti-C23 nucleolin (1:300, sc-13057 Santa Cruz Biotechnology). Quantification of protein expression was performed by densitometry analysis of immunoblots, using Gel Pro.3 analysis software (MediaCybernetics, MD, USA).

NF-*κ***B** activation. The binding of mouse p50 and RelA to the NF-*κ*B-binding consensus sequence was measured in nuclear extracts using the ELISA-based Mercury TransFactor kit (BD Biosciences, San Jose, CA, USA) as previously described.¹⁰ Data are expressed as the absorbance difference observed in the presence of nuclear extracts and that observed in the absence of nuclear extracts.

Expression plasmids and transfections. The wild-type RelA plasmid⁸ was used as a template to produce the Lys-to-arginine mutant construct RelA-K310R using the Quick change site-directed method (Stratagene, La Jolla, CA, USA), and *Pfu* DNA Polymerase. The primers sequence synthesized for the mutagenesis were the following: RelA-K310R for 5'-AGGACATATGAGACCTTCAG GAGCATCATGAGAAGAGAG'; RelA-K310R REV 5'-CTCTTCTTCATGATGCTCC TGAAGGATCATGATGATCCT-3' (nucleotide substitution is in bold). The results of mutagenesis were confirmed by sequencing clones with the following internal primer, K310R 5'-GCCTGCAGGCTCCTGTGCGT-3'. Restriction map analysis was further carried out to verify construct integrity, and expression was confirmed by immunobloting with the anti-RelA antibody in differentiated SH-N-SK cells, using the following antibodies: rabbit polyclonal anti-*R*elA (1:200, sc-372, Santa Cruz Biotechnology) and mouse monoclonal anti-*β*-tubulin (1:1500, NeoMarkers, Fremont, CA, USA).

Transfection of differentiated SK-N-SH cells was carried out according to the manufacturer's instructions with Lipofectamine 2000 Reagent (LF 2000, Invitrogen Corp., Carlsbad, CA, USA), as previously described.⁸ Cells were transfected with expression plasmids encoding ReIA, ReIA-K310R or empty expression vector pSG5 as a negative control, for 24 h, before undergoing the OGD experiments.

Reporter gene assays. In order to evaluate mouse *Bim* promoter activity during OGD, cortical neurons were transfected at 10 DIV using LF 2000 Reagent with 0.2 μ g/well of the Bim-pGL3 plasmid and 0.8 μ g/well of RelA or RelA-K310R mutant construct or empty expression vector pSG5 as negative control, as previously described.⁸ To normalize the transfection efficiency, 0.02 μ g/well *Renilla* luciferase (phRLTK) control plasmid (Promega) was used. After 24 h, neurons were exposed to 3 h OGD as described above. At the end of incubation, during the 4 h recovery in Neurobasal medium with 0.4% B27 supplement (Invitrogen Corp.), the cells expressing pSG5 were treated with resveratrol (30 μ M). Cells were then harvested, and firefly and *Renilla* luciferase were measured by using Dual Luciferase Reporter Assay (Promega).

Statistics. NF-*k*B ELISA data were analyzed using one-way analysis of variance (ANOVA), followed by Dunnett's *post hoc* analysis to determine statistical significance. *P*<0.05 was considered significant. Columns represent the means \pm S.E.M. of at least four values. Data describing cell survival were analyzed by Kruskal–Wallis non-parametric ANOVA with adjustment for multiple comparisons. Data relative to densitometry analyses and luciferase reporter activity were analyzed using Student's *t*-test for independent data. *P*<0.05 was considered as significant.

Conflict of interest

The authors declare no conflict of interest.

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