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Ischemic tolerance modulates TRAIL expression and its receptors and generates a neuroprotected phenotype

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TNF-related apoptosis inducing ligand (TRAIL), a member of the TNF superfamily released by microglia, appears to be involved in the induction of apoptosis following focal brain ischemia. Indeed, brain ischemia is associated with progressive enlargement of damaged areas and prominent inflammation. As ischemic preconditioning reduces inflammatory response to brain ischemia and ameliorates brain damage, the purpose of the present study was to evaluate the role of TRAIL and its receptors in stroke and ischemic preconditioning and to propose, by modulating TRAIL pathway, a new therapeutic strategy in stroke. In order to achieve this aim a rat model of harmful focal ischemia, obtained by subjecting animals to 100 min of transient occlusion of middle cerebral artery followed by 24 h of reperfusion and a rat model of ischemic preconditioning in which the harmful ischemia was preceded by 30 mins of tMCAO, which represents the preconditioning protective stimulus, were used. Results show that the neuroprotection elicited by ischemic preconditioning occurs through both upregulation of TRAIL decoy receptors and downregulation of TRAIL itself and of its death receptors. As a counterproof, immunoneutralization of TRAIL in tMCAO animals resulted in significant restraint of tissue damage and in a marked functional recovery. Our data shed new light on the mechanisms that propagate ongoing neuronal damage after ischemia in the adult mammalian brain and provide new molecular targets for therapeutic intervention. Strategies aimed to repress the death-inducing ligands TRAIL, to antagonize the death receptors open new perspectives for the treatment of stroke.

Cell Death and Disease (2014) 5, e1331; doi:10.1038/cddis.2014.286; published online 17 July 2014

Stroke is a leading cause of death in industrialized countries¹ and the most frequent cause of disability in adults.² Although different mechanisms are involved in the pathogenesis of stroke, increasing evidence shows that ischemic injury and subsequent inflammation are responsible for damage progression,³ characterized by irreversible neuronal damage within minutes of the onset.

In the past 30 years, it has been demonstrated that the brain's resistance to ischemic injury can be transiently augmented by previous exposure to a non-injurious preconditioning (PC) stimulus.⁴ Evidence demonstrates that PC inhibits apoptosis in the penumbra region, thus preventing the spread of infarction. In addition, PC-induced neuroprotection appears related to a persistent activation of survival kinases in the penumbra.⁵ PC seems recognized by sensor molecules, such as neurotransmitters, cytokines, and others, as a sign of an event potentially more severe to come.⁶ In particular, inflammatory cytokines, beside representing PC sensor molecules, have a relevant role in acute stroke. For instance, tumor necrosis factor- α (TNF- α) and IL-1 mediate the inflammatory/immune response related to progression of the

ischemic lesion.⁷ In the brain, cytokines are expressed not only in systemic immunocytes but also in resident cells, including neurons and glia.⁸ In particular, microglia have a key role as immune-competent cells of the CNS. Recruitment of leukocytes from the blood stream and activation of microglia are thought to contribute to the extension of the infarct core into the surrounding penumbra.⁵ Substantial evidence demonstrates that ischemia-activated microglia releases several pro-inflammatory cytokines, as well as other potentially cytotoxic molecules, including NO, ROS, and eicosanoids.⁹ Such a large arsenal of cytotoxic molecules appear to be involved in the induction of neuronal death.

TNF-related apoptosis inducing ligand (TRAIL), a proapoptotic member of the TNF superfamily released by glia^{10,11} and injured neurons,¹² appears to trigger apoptosis following focal brain ischemia.¹³ TRAIL binds five receptors, death receptor-4 (DR4), DR5, decoy receptor 1 (DcR1), DcR2, and osteoprogeterin. Although DR4 and DR5 receptors contain an intracellular death domain related to the apoptotic pathway, DcR1 and DcR2 serve as decoy receptors,¹⁴ eventually buffering death receptors' binding of TRAIL, thus preventing apoptosis.^{15,16}

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Received 04.3.14; revised 26.5.14; accepted 29.5.14; Edited by D Vucic

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Abbreviations: aCSF, artificial cerebral spinal fluid; DcR1, decoy receptor 1; DR4, death receptor-4; FasL, Fas ligand; JNK, Jun-N-terminal kinase; PC, preconditioning; tMCAO, transient middle cerebral artery occlusion; TNF-α, tumor necrosis factor-α; TNFR1, tumor necrosis factor receptor 1; TRAIL, TNF-related apoptosis inducing ligand

Although recent work attempted to establish a relationship between TRAIL pathway and brain ischemia,^{17,18} only scant data are available on the role of TRAIL and its receptors in focal ischemia,¹⁹ and no data are known on the role of TRAIL in brain PC-induced neuroprotection.

In the present study, we evaluated the role of TRAIL and its receptors in stroke and ischemic PC and verified the hypothesis of a potential mode for therapeutic intervention in stroke, by administrating a specific anti-TRAIL antibody in rats subjected to transient middle cerebral artery occlusion (tMCAO).

Results

Intracerebroventricular (i.c.v.) injection of TRAIL abrogates beneficial effects of PC in rats subjected to tMCAO. In order to determine the role of TRAIL in the neuroprotection exerted by ischemic PC, TRAIL was administered i.c.v. in rats. When TRAIL was injected both at $0.2 \mu g/kg$ and at $6 \mu g/kg$ immediately after tMCAO onset, it did not induce any significant change in total ischemic volume compared with vehicle-injected rats (54 ± 4.9 and $47.8 \pm 1.9\%$ *versus* $58.6 \pm 3.5\%$, respectively). Conversely, when TRAIL was administered i.c.v. at $6 \mu g/kg$ in rats undergone the same surgical procedure preceded by PC, it was able to induce a significant increase in the ischemic volume compared with the vehicle-injected rats (54.4 ± 9.1 *versus* $22.2 \pm 2.7\%$). Ischemic volume did not change when preconditioned rats were i.c.v. treated with vehicle or with TRAIL 0.2 μ g/kg (22.2 ± 2.8 *versus* $24 \pm 4.8\%$; Figure 1a). These data were mirrored by data on neurological scores. Indeed, a worsening in neurological deficits was observed in preconditioned animals treated with TRAIL 6 μ g/kg compared with vehicle-treated animals (Figure 1b).

Expression of TRAIL and its death and decoy receptors is modulated, along with that of inflammatory molecules, by tMCAO-induced brain ischemia and by ischemic PC. Expression levels of TRAIL and both classes



Figure 1 Effect of TRAIL administration on neuroprotection elicited by ischemic preconditioning. (a) $6 \mu g/kg$ TRAIL i.c.v. administered determines the loss of the neuroprotection elicited by ischemic PC. Infarct volume in rats subjected to tMCAO + Vehicle, tMCAO + 0.2 $\mu g/kg$ TRAIL, tMCAO + 6 $\mu g/kg$ TRAIL, PC + tMCAO + Vehicle, PC + tMCAO + 0.2 $\mu g/kg$ TRAIL, and PC + tMCAO + 6 $\mu g/kg$ TRAIL. Rats were euthanized 24 h after tMCAO. **P* < 0.05 *versus* all experimental groups. Each column represents the mean ± S.E.M. (*n* = 3–7) of the percentage of the infarct volume compared with the ipsilateral hemisphere. (b) A total of 6 $\mu g/kg$ TRAIL i.c.v. administered worsens general and focal deficits in rats subjected to PC + tMCAO. **P* < 0.05 *versus* PC + vehicle-treated animals

of its receptors, death DR4 and DR5, and decoy DcR1 and DcR2 were studied by western blotting analysis in animals subjected to PC, tMCAO, or PC followed, 72 h later, by tMCAO. All animals were then allowed to 24-h reperfusion.

Expression of TRAIL increased significantly in rats subjected to tMCAO after 24 h of reperfusion. This increased

expression of TRAIL was dramatically reduced in the temporoparietal brain cortices of animals exposed to either PC alone or PC followed by tMCAO.

The increased expression of both DR4 and DR5 death receptors in the tMCAO experimental group closely paralleled that of TRAIL. In particular, DR4 and DR5 expression, after



Figure 2 Effect of tMCAO/PC followed by 24 h of reperfusion on either TRAIL and its receptor's protein expression, as well as on TNF- α , FasL, and their respective receptors. (a) Representative blots of DR4, DR5, TRAIL, DcR1, and DcR2 protein expression in sham-operated animals (SHAM) or after 100 min tMCAO (tMCAO), 30 min tMCAO (PC), and 30 min PC, followed, 72 h later, by 100 min tMCAO (PC + tMCAO) in the ipsilateral temporoparietal cortex after 24 h of reperfusion. Tubulin blots: β -tubulin (respective controls). (b) Representative blots of TNFR1, TNF- α , Fas, and FasL protein expression in sham-operated animals (SHAM) and after 100 min tMCAO (tMCAO), 30 min tMCAO (PC), and 30 min PC, followed, 72 h later, by 100 min tMCAO (PC + tMCAO) in the ipsilateral temporoparietal cortex after 24 h of reperfusion. Tubulin blots: β -tubulin (respective controls). (b) Representative blots of TNFR1, TNF- α , Fas, and FasL protein expression in sham-operated animals (SHAM) and after 100 min tMCAO (tMCAO), 30 min tMCAO (PC), and 30 min PC, followed, 72 h later, by 100 min tMCAO (PC + tMCAO) in the ipsilateral temporoparietal cortex after 24 h of reperfusion. Tubulin blots: β -tubulin (respective controls).



Figure 3 Effect of tMCAO/PC followed by 24 h of reperfusion on either caspases-8 and -3, TRAIL pathway-related kinase JNK, and the cell survival-related kinase Akt. (a) Representative blots of caspase-8 and caspase-3 protein expression in sham-operated animals (SHAM) and after 100 min tMCAO (tMCAO), 30 min tMCAO (PC), and 30 min preconditioning, followed, 72 h later, by 100 min tMCAO (PC + tMCAO) in the ipsilateral temporoparietal cortex after 24 h of reperfusion. Tubulin blots: *β*-tubulin (controls). (b) Representative blot of phospho-JNK and phospho-AKT protein expression in sham-operated animals (SHAM) or after 100 min tMCAO (tMCAO), 30 min tMCAO (PC), and 30 min PC, followed, 72 h later, by 100 min tMCAO (PC + tMCAO) in the ipsilateral temporoparietal cortex after 24 h of reperfusion. Unphosphorylated JNK and AkT are controls

PC, was comparable to normal controls and dramatically resulted in decrease after PC followed by tMCAO.

Conversely, the expression of both TRAIL-neutralizing receptors DcR1 and DcR2 was substantially reduced in animals undergone tMCAO, whereas a significant increase occurred in animals undergone PC only. The expression of both DcR1 and 2 was even more intense in animals receiving both PC and tMCAO (Figure 2a).

As TRAIL is known to significantly increase the expression of related inflammatory molecules,²⁰ the expression of TNF- α , Fas ligand (FasL) and their respective death receptors tumor necrosis factor receptor 1 (TNFR1) and Fas was also measured.

Although tMCAO significantly increased the expression of all the proteins studied, on the other hand, expression of TNF- α , FasL, TNFR1 and Fas was decreased by PC and, more substantially, when applying PC before tMCAO (Figure 2b).

Moreover, as caspase-8 and caspase-3 pathways are both activated by TRAIL and the resulting increased cell death rate involves activation of related kinase Jun-N-terminal kinase (JNK), in order to verify whether the molecular mechanisms underlying the effects of TRAIL were actually set into motion, cleavage of downstream caspase-8 and caspase-3 was first evaluated in the same lysates, along with phosphorylation of JNK.

As a consequence of tMCAO, both caspase-8 and caspase-3 were substantially cleaved (i.e., activated), whereas caspase cleavage was significantly reduced in animals undergone PC, and almost totally abrogated in rats undergone precondition procedure before tMCAO (Figure 3a).

In parallel, phosphorylation of JNK, which was significantly increased in rats undergone tMCAO, was comparable to basal levels either in rats with PC alone or in rats undergone PC before tMCAO (Figure 3b). By contrast, to correlate the increased expression of the TRAIL DcR1 and DcR2 decoy receptors to the survival kinases AKT, we evaluated the phosphorylation of the latter. Accordingly with the expression level of the two decoy receptors, phospho-AKT was substantially reduced in the tMCAO group, whereas PC and PC + tMCAO produced its robust increase (Figure 3b).

Confocal immunofluorescence imaging of TRAIL expression at different time intervals after tMCAO, PC and PC + tMCAO in the ipsilateral peri-ischemic region. TRAIL expression has been evaluated at different time intervals by immunofluorescence in tissue slices from ipsilesional temporoparietal cortex of ischemic rats subjected to 100 min tMCAO, preconditioned rats ($30 \min + tMCAO$) and ischemic preconditioned rats ($30 \min + tMCAO$), followed, 72 h later, by 100 min tMCAO).

Double fluorescence immunostaining of both TRAIL and the neuronal marker NeuN has revealed that, in ischemic rats, robust TRAIL immunoreactivity is localized mainly in the cytosol and in the nuclei of neurons. Differently, when rats were subjected to PC alone or to PC followed by tMCAO, TRAIL immunoreactivity was expressed only at the cytoplasmatic level.

With regard to TRAIL expression, immunofluorescent data confirmed those obtained with western blotting, showing an increase in TRAIL expression when rats were subjected to harmful ischemia (100 min tMCAO), and a substantial



Figure 4 Effect of 100 min of transient brain ischemia (tMCAO), ischemic PC, and PC followed 72 h later by tMCAO (PC + tMCAO) on TRAIL expression. Confocal microscopic images displaying NeuN (a–I) (green), TRAIL (b–m) (red), and Merge (c–n) (yellow) in the brain peri-ischemic region of rats after 5 h (A), 24 h (B), and 72 h (C) of reperfusion. A representative brain slice cartoon indicating the area of interest is on the left top of the figure. Scale bars in a–i: 50 μ m



reduction of TRAIL expression when the animals were exposed to PC or to PC followed by tMCAO. These data were confirmed at all the time intervals considered, 5, 24, and 72 h of reperfusion (Figures 4A–C).

Ischemic PC induces decreased expression of TRAIL death receptor DR4 and DR5 in parallel to overexpression of its decoy receptors, DcR1 and DcR2. In order to establish whether the loss of neuroprotection observed after TRAIL overexpression could be related to changes in the expression of DR4-DR5 and DcR1-DcR2, the expression of these proteins was evaluated by means of confocal microscopy analysis on coronal brain tissue slices of rats subjected to tMCAO, ischemic PC, or ischemic PC followed by 100 min tMCAO at 5, 24, and 72 h of reperfusion.

Interestingly, an inverse relationship between TRAIL death receptor expression and decoy receptor was observed.

In fact, both DR4 and DR5 protein expression increased during tMCAO and decreased after PC and after PC followed by tMCAO, thus indicating that this protein is having a detrimental role during the ischemic process (Figures 5 and 6A–C).

On the other hand, the expression of DcR1 and DcR2 was significantly reduced after harmful ischemia, whereas its expression increased after PC + tMCAO at all the considered reperfusion time intervals, suggesting that a possible buffering role of DcR1-DcR2 is set into motion to counteract TRAIL-induced neuronal death (Figures 7 and 8A–C).

Intracerebroventricular injection of an anti-TRAIL monoclonal antibody induces neuroprotection in rats subjected to tMCAO. To corroborate the hypothesis of the detrimental role of TRAIL in brain ischemia, a TRAIL-neutralizing antibody (CD253, anti-TRAIL) was injected i.c.v. in rats subjected to 100 min tMCAO, followed by 24 h reperfusion. Each rat received a total volume of 20 μ l, four times. Although anti-TRAIL at a low dose was not able to exert any significant effect (51.5 ± 4.1 *versus* 58.6 ± 3.5%), when anti-TRAIL was administered at 200 μ g/kg, it determined a significant reduction of the ischemic volume in comparison with ischemic animals treated with the anti-idiotype (30.1 ± 5.8 *versus* 58.6 ± 3.5%). No effects were observed when scrambled anti-TRAIL was administered (55.4 ± 3.8 *versus* 58.6 ± 3.5%; Figure 9a).

To assess whether the decrease in the infarct volume observed after i.c.v. administration of anti-TRAIL 200 μ g/kg in ischemic rats was accompanied by an amelioration in neurological deficits, animals were scored for general and focal neurological deficits immediately before they were killed. Results demonstrated that anti-TRAIL, injected i.c.v., besides decreasing the brain infarct volume, was also able to decrease focal and general neurological scores when evaluated at 24 h (Figure 9b).

Figure 5 Effect of 100 min of transient brain ischemia (tMCAO), ischemic PC, and PC followed 72 h later by tMCAO (PC + tMCAO) on DR4 expression. Confocal microscopic images displaying NeuN (a–l) (green), DR4 (b–m) (red), and Merge (c–n) (yellow) in the brain peri-ischemic region of rats after 5 h (A), 24 h (B), and 72 h (C) of reperfusion. A representative brain slice cartoon indicating the area of interest is on the left top of the figure. Scale bars in a–i: 50 μ m



Figure 6 Effect of 100 min of transient brain ischemia (tMCAO), ischemic PC, and PC followed 72 h later by tMCAO (PC + tMCAO) on DR5 expression. Confocal microscopic images displaying NeuN (a–I) (green), DR5 (b–m) (red), and Merge (c–n) (yellow) in the brain peri-ischemic region of rats after 5 h (**A**), 24 h (**B**), and 72 h (**C**) of reperfusion. A representative brain slice cartoon indicating the area of interest is on the left top of the figure. Scale bars in a–i: 50 μ m

Discussion

This study provides evidence that the neuroprotection elicited by ischemic PC occurs either through upregulation of the TRAIL decoy receptor DcR2 and through down-regulation of TRAIL and its death receptor DR5. In addition, here we propose an experimental strategy for inducing stroke neuroprotection by modulating TRAIL activity by using *in vivo* the selective monoclonal antibody anti-TRAIL.

Although the biological importance of TRAIL is not completely understood, it appears that this protein may have a critical role in cellular response to environmental stress.^{11,21-23} As such, TRAIL signaling also promotes an array of biological responses associated with cellular growth and survival.²⁴ In line with these findings, we report novel evidence here that TRAIL is upregulated in the ischemic brains of animals subjected to focal ischemia, whereas it appears dramatically downregulated in the brains of animals subjected to a sub-lethal ischemia, PC, or an harmful ischemia preceded by PC. Given that TRAIL represents the ligand for two death receptors and for two decov receptors, we also evaluated the expression profile of these receptors after stroke, and we observed that the increase in TRAIL would undoubtedly complement the increase of its death receptors DR4 and DR5, while its downregulation parallels the increased expression of its decoy receptors DcR1 and DcR2, resulting in a substantial increase in TRAIL signaling following stroke and a substantial decrease in TRAIL pathway after ischemic PC. Significant increase in the expression of the DcR1/DcR2 receptors, which is known to bind TRAIL not bringing about any signal transduction,¹⁵ appears to be of special interest in the ischemic context, as it represents a modality of neutralizing a part of TRAIL and its effects, which include sustained cell death rates.²⁵ Such hypothesis is supported by the parallel increase of the cell survival-related kinase AkT,⁵ indicating that PC prevents detrimental effects of TRAIL and sets into motion the cell survival machinery to rescue neurons to death.

At any rate, detrimental effects of TRAIL appeared mediated through activation of caspase-8 and -3, as well as phosphorylation of the stress kinase JNK, both key elements involved in transduction of the TRAIL-dependent cell death signaling,^{26,27} thus corroborating the hypothesis that TRAIL could represent a substantial contributor to poststroke neurodegeneration processes.

Consistent with the findings that TRAIL overexpression is deleterious, whereas TRAIL reduction is neuroprotective following stroke,¹⁷ our data provide evidence that TRAIL downregulation is, indeed, crucial to stroke recovery. This represents an important finding, as, so far, few studies have investigated the role of TRAIL in cell death induced by ischemia,^{17–19,28} and no one investigated about the possible involvement of TRAIL in the neuroprotective phenomenon known as ischemic PC.

In fact, the overexpression of TRAIL and its death receptors after hypoxia–ischemia has been underlined in some seminal papers.^{18,28} Indeed, TRAIL and DR5 are upregulated in the immature rat brain after perinatal hypoxia–ischemia¹⁸ as well as after transient global ischemia.¹⁷ In addition, it has been shown that blocking

TRAIL by soluble DR5 mitigate selective neuronal death after transient global ischemia–reperfusion in mice.¹⁷

In addition, increased TRAIL expression was paralleled by the increased levels of proinflammatory molecules, such as TNF- α and its death receptor TNFR1 or FasL and its receptor Fas. It is known that one of the biological properties displayed by cytokines, regardless of their biological role, is redundancy,²⁹ and we have shown that the increase of tissue TRAIL expression in a murine model of spinal cord injury is paralleled by an increase of both tissue TNF- α and FasL expression.²⁰ Thus it is plausible that such potent effects of TRAIL in inducing poststroke neuronal death is also related to redundancy with two cytokines, TNF- α and FasL, possessing similar properties,³⁰ therefore potentiating TRAIL-induced neuronal death, via binding to their own death receptors.

Thus we went on and proposed a feasible therapeutic strategy for stroke, based on the use of a monoclonal antibody against TRAIL as a potential compound to be used in stroke intervention. In particular, we mimicked the protective effect mediated by ischemic PC by administering anti-TRAIL to ischemic rats and proposed new insights into the putative mechanisms by which ischemic PC exerts its protective effect, with restored functional parameters.

Concerning a possible mechanism by which PC influences the expression of TRAIL and its receptor, it is possible to hypothesize that the two transcriptional factors NF κ B and HIF, involved in PC neuroprotection,^{31,32} may have a fundamental role. Interestingly, recent findings demonstrated that both these factors are tightly linked to TRAIL and its receptors. Indeed, the C-terminal element of DcR2 has signaling capacity similar to that of DR4 and DR5 with respect to NF κ B activation³³ but is unable to induce apoptosis. Thus increased expression of TRAIL decoy receptors by IPC may activate a secondary complex-linked survival kinase pathway. Studies implicating NFkB in TRAIL-induced apoptosis are few in number and conflicting. NF κ B expression is shown to be involved in protection against TRAIL,³⁴ but a functional NFkB binding site is located in the promoter region of DR4.35 NFkB has been reported to enhance TRAIL-induced apoptosis through intronic regulation of DR5.36 In addition, DcR2 expression can be induced by the other above mentioned transcription factor HIF³⁷ that is strongly upregulated by ischemic PC.³²

In conclusion, we detected upregulation of TRAIL and its death receptors, DR4 and DR5, as well as a reduction in the expression of the decoy receptors, DcR1 and DcR2, in the postischemic brain, whereas an opposite expression behavior was observed for the same proteins in the protected preconditioned brain, thus demonstrating, *in vivo*, their potency to mediate cell death or protection.

Our data shed new light on the mechanisms that propagate ongoing neuronal damage after ischemia in the adult mammalian brain and provide new molecular targets for therapeutic intervention. Strategies aimed to repress either

Figure 7 Effect of 100 min of transient brain ischemia (tMCAO), ischemic PC, and PC followed 72 h later by tMCAO (PC + tMCAO) on DcR1 expression. Confocal microscopic images displaying NeuN (A–L) (green), DcR1 (b–m) (red), and Merge (c–n) (yellow) in the brain peri-ischemic region of rats after 5 h (**A**), 24 h (**B**), and 72 h (**C**) of reperfusion. A representative brain slice cartoon indicating the area of interest is on the left top of the figure. Scale bars in a–i: 50 μ m



Α

В

С

SHAM

tMCAO 5HRS

PC 5HRS

PC+tMCAO 5HRS

SHAM

tMCAO 24HRS

PC 24HRS

PC+tMCAO

SHAM

tMCAO 72HRS

PC 72HRS

PC+tMCAO

DCR1

DCR1



Figure 8 Effect of 100 min of transient brain ischemia (tMCAO), ischemic PC, and PC followed 72 h later by tMCAO (PC + tMCAO) on DcR2 expression. Confocal microscopic images displaying NeuN (a–I) (green), DcR2 (b–m) (red), and Merge (c–n) (yellow) in the brain peri-ischemic region of rats after 5 h (A), 24 h (B), and 72 h (C) of reperfusion. A representative brain slice cartoon indicating the area of interest is on the left top of the figure. Scale bars in a–i: 50 μ m

the death-inducing ligands TRAIL, to antagonize death receptor, DR5, or to activate the decoy receptor, DcR2, open new perspectives for the treatment of stroke.

Materials and methods

Animals. One hundred and fifteen male Sprague Dawley rats (250–270 g; Charles River, Lecco, Italy) housed five per cage under diurnal lighting conditions (12 h dark–light cycle) were used in the present study. Experiments were performed in full accordance with the international guidelines for animal research and humane care. The experimental protocol was approved by the Animal Care Committee of the 'Federico II' University of Naples, Italy.

Experimental groups and surgical procedures. Transient focal ischemia was induced as previously described $^{\rm 38}$ by suture occlusion of the MCA in male rats anesthetized using 1.5% sevoflurane, 70% N₂O, and 28.5% O₂. Achievement of ischemia was confirmed by monitoring regional cerebral blood flow through laser Doppler (PF5001; Perimed, Jarfalla, Sweden). Animals not showing a cerebral blood flow reduction of at least 70% were excluded from the study. Before treatment, rats were randomly divided into three main experimental groups: (1) preconditioned. (2) ischemic, and (3) preconditioned ischemic rats (undergone PC+tMCAO). The sham-operated animals underwent the same experimental conditions except that the occluding filament was not introduced in the vessel lumen; in the ischemic group, the MCA was occluded for 100 min; in the preconditioned ischemic group, rats were subjected to 30 min of tMCAO 72 h before 100 min of tMCAO. All animals were killed by decapitation 24 h after 100 min of tMCAO to quantify the infarct volume. Rectal temperature was maintained at 37 ± 0.5 °C. Blood gas analysis was carried out in all animals, and no differences in pH, pCO₂ and PO₂ was detected among all the experimental groups.

Intracerebroventricular (i.c.v.) administration. In rats positioned on a stereotaxic frame, a 23-g stainless steel guide cannula was implanted into the right lateral ventricle using the stereotaxic coordinates from the bregma: 0.4 mm caudal, 2 mm lateral and 2 mm below the dura.³⁹ TRAIL at doses ranging from 0.2 to 6 μ g/kg was administered in a final volume of 5 μ l, either immediately after tMCAO induction or after PC + ischemia. Conversely, anti-TRAIL (20 and 200 μ g/kg, respectively) was injected four times, dissolved in artificial cerebral spinal fluid (aCSF) at a final volume of 20 µl, at 24 and 12 h before ischemia induction, as well as immediately after ischemia onset and 6 h after. Vehicle-injected rats undergoing tMCAO or PC+tMCAO received i.c.v. aCSF in the same volume and on the same time schedule as TRAIL- or anti-TRAIL-treated animals. Animals were randomly allocated to each experimental group. The number of animals excluded because they died during the surgery procedures or because the reduction in the CBF was <70% was the following: tMCAO vehicle-treated group: 2, 3; tMCAO 20 µg/kg anti-TRAIL: 1, 2; tMCAO 200 µg/kg anti-TRAIL: 0, 3; tMCAO 200 µg/kg scrambled anti-TRAIL: 1, 2; tMCAO 0.2 µg/kg TRAIL: 0, 3; tMCAO 6 µg/kg TRAIL: 1, 2; PC followed by tMCAO vehicle-treated group: 0, 1; PC followed by tMCAO 0.2 µg/kg TRAIL: 1, 3; and PC followed by tMCAO 6 µg/kg TRAIL: 2, 4.

Evaluation of the infarct volume and of neurological deficit scores. Animals were killed by decapitation 24 h after 100 min. Brains were quickly removed, sectioned coronally at 1-mm intervals, and stained by immersion in the vital dye (2%) 2,3,5-triphenyltetrazolium hydrochloride. The infarct volume was calculated by summing the infarction areas of all sections and by multiplying the total by slice thickness. To avoid that edema could affect the infarct volume value, it has been chosen to express the infarct volume as a percentage of the infarct, calculated by dividing the infarct volume by the total ipsilateral hemispheric volume.^{40,41}

Neurological scores were evaluated 24 h after reperfusion according to two scales: a general neurological scale and a focal neurological scale. In the general score, the following six general deficits were measured: (1) hair conditions, (2) position of ears, (3) eye conditions, (4) posture, (5) spontaneous activity, and (6) epileptic behavior. For each of the six general deficits measured, animals received a score ranging between 0 and 12 depending on the severity of signs. The scores of investigated items were then summed to provide a total general score. In the focal score, the following seven areas were assessed: (1) body symmetry, (2) gait, (3) climbing, (4) circling behavior, (5) front limb symmetry, (6) compulsory circling, and (7) whisker response. For each of these items, animals were rated between 0 and 4 depending on severity. The seven items were then summed to give a total focal score. Infarct volumes, neurological scores, and animal survival were evaluated in a blinded manner.

Western blotting analysis. In the western blotting experiments, lysates were obtained from rats subjected to tMCAO, ischemic PC, or ischemic PC followed by 100 min tMCAO 24 h of reperfusion (n = 3 per group; 2 were excluded because of death and 3 excluded because CBF did not reach a reduction of at least 70%). The protein concentration of the supernatant was determined by the

npg



Figure 9 Effect of anti-TRAIL administration on ischemic damage elicited by 100 min of transient middle cerebral occlusion followed by 24 hours of reperfusion. (a) In all, 200 μ g/kg antiTRAIL, i.c.v. injected, induces neuroprotection in rats subjected to tMCAO followed by 24 h reperfusion. Infarct volume in rats subjected to tMCAO + Vehicle, tMCAO + 20 μ g/kg anti-TRAIL, tMCAO + 200 μ g/kg anti-TRAIL, and tMCAO + 200 μ g/kg scrambled anti-TRAIL. **P*<0.05 versus all experimental groups. Each column represents the mean ± S.E.M. (*n*=5–7) of the percentage of the infarct volume compared with the ipsilateral hemisphere. (b) In all, 200 μ g/kg anti-TRAIL injected i.c.v. ameliorates general and focal deficits in rats subjected to PC + tMCAO. **P*<0.05 versus vehicle-treated animals

Bradford method.⁴² Equal amounts of protein (50 µg) were subjected to sodium dodecyl sulfate-polyacrylamide gel electrophoresis on 8 and 12% gels and then transferred onto Hybond ECL nitrocellulose membranes (Amersham Life Science, Buckinghamshire, UK). The membranes were blocked with 5% milk in PBST and then incubated overnight at 4 °C with a rabbit anti-DR4 polyclonal antibody (ProSci Inc., Poway, CA, USA) or a rabbit anti-DR5 polyclonal antibody (Abcam, Cambridge, UK) or a rabbit anti-TRAIL polyclonal antibody (Abcam) or a rabbit anti-DcR1 (ProSci) polyclonal antibody or a rabbit anti-DcR2 polyclonal antibody (Abcam) or a rabbit anti-TNFRI polyclonal antibody (Abcam) or a rabbit anti-TNF alpha polyclonal antibody (Novus Biologicals, Littleton, CO, USA) or a mouse anti-Fas monoclonal antibody (BD Biosciences, San Jose, CA, USA) or a rabbit anti-FasL polyclonal antibody (Abcam) or a mouse anti-Caspase-8 monoclonal antibody (Cell Signaling Technology, Danvers, MA, USA) or a rabbit anti-Caspase-3 monoclonal antibody (Cell Signaling Technology) or a mouse anti-Phospho-JNK monoclonal antibody (Santa Cruz Biotechnology, Santa Cruz, CA, USA) or a rabbit anti-JNK polyclonal antibody (Santa Cruz Biotechnology) or a rabbit anti-Phospho-Akt monoclonal antibody (Cell Signaling Technology) or a rabbit anti-Akt monoclonal antibody (Cell Signaling Technology) and horseradish peroxidaseconjugated anti-rabbit or anti-mouse IgG secondary antibody (Amersham Life Science). *β*-Tubulin (Santa Cruz Biotechnology) was used as an internal control to validate the right amount of protein loaded in the gels. Detection was performed by means of ECL chemiluminescence assay (Amersham Life Science).

Immunohistochemistry. In immunofluorescence experiments, rats were subjected to tMCAO, ischemic PC, or ischemic PC followed by 100 min tMCAO at 5, 24, and 72 h of reperfusion (n=3 per group; 1 excluded because of death)and 2 excluded because CBF did not reach a reduction of at least 70%). Immunostaining and confocal immunofluorescence procedures were performed as previously described.⁴³ Rats were anesthetized with chloral hydrate (300 mg/kg, intraperitoneally) and perfused transcardially with 4% paraformaldehyde and 15% picric acid in phosphate buffer. The brains were sectioned coronally at 40 mm on a vibratome. After blocking, sections were incubated with the following primary antisera: mouse monoclonal anti-NeuN (1:1000, Millipore, Milan, Italy), rabbit polyclonal anti-TRAIL (1:200; Abcam), rabbit polyclonal anti-DcR2 (1:200; Abcam), rabbit polyclonal anti-DcR1 (1:200; ProSci), rabbit polyclonal anti-DR4 (1:200; ProSci), and rabbit polyclonal anti-DR5 (1:200; Abcam). The sections were incubated with the corresponding fluorescent-labeled secondary antibodies (Alexa 488/Alexa 594-conjugated antimouse/antirabbit IgGs). Images were observed using a Zeiss LSM510 META/laser scanning confocal microscope. Single images were taken with an optical thickness of 0.7 m and a resolution of 1024×1024 .

In double-labeled sections, the pattern of immunoreactivity for both antigens was identical to that seen in single-stained material.

Controls of the methods in the double immunofluorescence experiments included replacement of the primary antisera with normal serum. To control for a possible cross-reactivity between IgGs in double immunolabeling experiments, some sections were processed through the same immunocytochemical sequence except that primary antisera were replaced with normal serum or only one primary antibody was applied, but the full complement of secondary antibodies was maintained. In addition, the secondary antibodies utilized were highly preadsorbed to the IgGs of

numerous species. Tissue labeling without primary antibodies was also tested to exclude autofluorescence. No specific staining was observed under these control conditions, thus confirming the specificity of the immunosignals.

Statistical analysis. For the statistic analysis of the effect of TRAIL and anti-TRAIL on the infarct volume, data were expressed as mean \pm S.E., and statistical analysis was performed with two-way ANOVA followed by Newman–Keuls test.

The data related to focal and general neurological deficits, being non-parametric data, were analyzed using the non-parametric Kruskal–Wallis test, followed by the Nemenyi test for the non-parametric multiple comparison. Statistical significance was accepted at the 95% confidence level (P < 0.05).

Conflict of Interest

The authors declare no conflict of interest.

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