Interference of apoptosis in the pathophysiology of subarachnoid hemorrhage

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ABSTRACT
Programmed cell death is crucial for the correct development of the organism and the clearance of harmful cells like tumor cells or autoreactive immune cells. Apoptosis is initiated by the activation of cell death receptors and in most cases it is associated with the activation of the cysteine proteases, which lead to apoptotic cell death. Cells shrink, chromatin clumps and forms a large, sharply demarcated, crescent-shaped or round mass; the nucleus condenses, apoptotic bodies are formed and eventually dead cells are engulfed by a neighboring cell or cleared by phagocytosis. The authors have summarized the most important data concerning apoptosis in subarachnoid hemorrhage that have been issued in the medical literature in the last 20 years.

Key words: Apoptosis, cell death receptors, programmed cell death, subarachnoid hemorrhage

Introduction
Subarachnoid hemorrhage [Figure 1] is associated with high mortality as 14% of patients die before reaching the hospital.[¹]

These deaths occur mostly as a result of the initial hemorrhage, and no effective treatment is available for brain injury after the hemorrhage.[²] For survivors, early brain injury caused by the initial hemorrhage and delayed ischemic neurologic deficits due to cerebral vasospasm [Figure 2] are major causes of the subsequent morbidity and mortality.[³]

Although cerebral vasospasm has been studied and treated using a wide array of drugs during the past several decades, the outcome is not improved by the reversal of vasospasm.[⁴] Early brain injury is considered a prime target for future research and may be also an important factor in preventing symptomatic vasospasm. In this respect, early brain injury may predispose the brain to ischemic injury due to vasospasm. Recent studies showed that apoptosis is involved in the pathogenesis of early brain injury after experimental subarachnoid hemorrhage (SAH) or in a clinical setting.[⁵,⁶] Therefore, it is thought that an antiapoptotic treatment can be a therapeutic candidate for early brain injury after SAH.

Pathophysiology of Early Brain Injury
Most available information about early brain injury after SAH comes from endovascular filament perforation animal models, which show high mortality and acute metabolic changes similar to the clinical settings.[⁷,⁸]

Intracranial pressure in this model was increased to 40 mmHg immediately after SAH and then decreased to plateau (15-25 mmHg), whereas cerebral perfusion pressure was decreased to 35-40 mmHg from 70 mmHg, cerebral blood flow was decreased with 20-30% beneath the baseline after SAH induction, and then each of the values were gradually recovered.[⁹] Interestingly, the mortality rate was 100% when cerebral blood flow was reduced to less than 40% underneath the baseline for 60 min after SAH, while a less augmented cerebral blood flow reduction resulted in a 19% mortality.[¹⁰]

Many factors, such as global ischemia,[¹¹] microcirculatory disturbance,[¹⁰] and subarachnoid blood toxicity[¹²] are involved in apoptosis-related mechanisms in early brain injury after SAH, whereas distribution of apoptotic cell death is controversial.[¹¹,¹³] Although apoptotic cell death was seen in both the cortex and subcortex, neuronal cell death in the

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Palade, et al.: A apoptosis and the mechanisms of subarachnoid hemorrhage

hippocampus, which is related to global ischemia, may depend on intracranial pressure.\(^6,13\)

Blood immediately spreads in the subarachnoid space after SAH, and then the cerebral cortex is covered with a thick blood clot. Hemoglobin is metabolized by neurons and microglia,\(^14\) and the released iron induces apoptosis via lipid peroxidation. Thus, subarachnoid blood clotting, which has been linked to cell injury and oxidative stress,\(^12\) may cause greater apoptotic cell death in the cerebral cortex compared with the subcortex. Apoptotic cell death has been reported to occur in neurons\(^12,15,16\) and endothelial cells\(^17,18\) in early brain injury after SAH. Both these situations may be correlated with brain edema.\(^19\) In this article, we focus on neuronal cell apoptosis, which consists of the intrinsic and extrinsic pathways.\(^20\)

Apoptosis represents the most well-characterized type of programmed cell death. Morphologically, cells typically round up, form blebs, undergo chromatin condensation and nuclear fragmentation. These morphological changes are largely the result of the activation of a set of cell-suicide cysteine proteases referred to as caspases.\(^21\)

The biochemical activation of apoptosis occurs through two general pathways: The intrinsic pathway, which is mediated by the mitochondrial release of cytochrome C and resultant activation of caspase-9; and the extrinsic pathway, originating from the activation of cell surface death receptors such as Fas, resulting in the activation of caspase-8 or -10 (Salvesen and Dixit, 1997). A third general pathway, which is essentially a second intrinsic pathway, originates from the endoplasmic reticulum and also results in the activation of caspase-9.\(^22\) Both extrinsic and intrinsic apoptotic pathways are synthesized in Figure 3.

**Intrinsic Mechanisms of Apoptosis and SAH**

**Caspase-dependent pathway**

The intrinsic pathway (mitochondrial pathway), which is mediated by the Bcl-2 family, begins with the increase in outer mitochondrial membrane permeability. This alteration of membrane permeability leads to the leakage of cytochrome C. Cytochrome C is translocated from mitochondria to the cytosolic compartment and interacts with apoptotic proteases, activating factor-1 and forming the apoptosome while leading to caspase-9 activation.

Caspase-9 activates caspase-3, and results in DNA damage.\(^23\) Caspase-3 is well known as one of the effectors of apoptosis, and cleaved caspase-3 is upregulated in the hippocampus and cortex after SAH.\(^11, 24, 25\)

Some reports showed that some protein kinases might directly interact with mitochondrial proteins in cerebral ischemia.

Their role mainly concentrates on the phosphorylation of pro- and anti-apoptotic proteins (Bad, Bax, Bcl-2, Bcl-xl).\(^26\) Akt (protein kinase B) and mitogen-activated protein kinase (MAPK) were the best studied in early brain injury after SAH. Akt, which is a serine/threonine kinase, is a key anti-apoptotic signaling enzyme positioned downstream of phosphoinositide 3-kinase (PI3K) in a growth factor mediated signaling cascade.

Stimulation of receptor tyrosine kinases or GTP-binding protein-coupled receptors activates Akt via PI3K. Activated Akt modulates many substrates, including Bax, Bad, glycogen synthase kinase-3, apoptosis signal-regulating kinase 1, and caspase-9, which inhibit apoptosis.\(^27\) Akt has also been shown to promote cyclic AMP response element-binding (CREB) protein phosphorylation and lead to Bcl-2 induction.\(^28\)

A decrease in Akt activity is responsible for ischemic neuronal cell death. Last but not least, Akt activation is a principal factor in the prevention of apoptosis via the caspase-dependent pathway in cerebral ischemia.\(^29,30\)

Some studies suggested that Akt might be involved in the
mechanism for early brain injury after SAH. This conclusion was drawn using a PI3K inhibitor, which prevented phosphorylation of Akt and increased DNA damage. Akt activation by overexpression of copper/zinc superoxide-dismutase (SOD1), which is one of the antioxidant enzymes, attenuated early brain injury caused by SAH. The timing of Akt phosphorylation after SAH depended on the damaged brain regions; Akt was rapidly phosphorylated in the cortex, but it took 24 h to phosphorylate Akt in the hippocampus. Because early brain injury after SAH is the most severe in the cortex, it is suggested that Akt phosphorylation depends on the severity of brain injury.

The roles of MAPKs are very important in early brain injury after SAH. MAPK, including extracellular signal-regulated kinase (ERK), c-Jun N-terminal kinase (JNK) and p38, is involved in the apoptotic responses in cell death during cerebral ischemia. These kinases are activated by various stimuli, e.g. vascular endothelial growth factor (VEGF), oxidative stress, and inflammatory cytokines.

After SAH in a perforation model, these kinases were phosphorylated and induced brain edema, continuous high intracranial pressure, and high mortality. Since ERK is activated in response to growth and differentiation factors and might be part of the survival pathway, whether activation of ERK is protective or detrimental to neurons in cerebral ischemia is controversial. In contrast, JNK and p38 are activated in response to inflammatory cytokines and cellular stress, which were highly elevated in the cerebrospinal fluid and in cerebral arteries after SAH. JNK phosphorylates c-Jun, which upregulates apoptotic cascades by inducing expression of the proapoptotic member of Bcl-2 family Hrk/DP5, Bim, and Fas.

Phosphorylated JNK and expression of c-Jun were increased after SAH induction and c-Jun mRNA were upregulated in the rat cerebral cortex and hippocampus after SAH. p38 activation by TNF-a and IL-1b was associated with neuronal death, and suppression of p38 activation by Bcl-2 suggested that p38 might be involved in apoptosis.

**Caspase-independent pathway**

The caspase-independent component of the intrinsic pathway is carried out by the mitochondria-released apoptosis inducing factor (AIF), endonuclease G and Bcl-2/adenovirus EIB 19 kDa-interacting protein (BNIP3). Apoptosis inducing factor, which is the best studied among them, is normally in the mitochondrial intermembrane space and is translocated to the nucleus by some stimulations, inducing large-scale DNA fragmentation and cell apoptosis, which is independent of caspase activity. Nuclear AIF upregulation was reported in cerebral ischemia, and the translocation might be triggered by poly (ADP-ribose) polymerase activity. There has not been much reported about AIF expression in early brain injury after SAH and it is not clear which compartment of AIF expression increases.

**Oxidative stress and early brain injury**

It is important to hold the balance between reactive oxygen species (ROS) and antioxidants, which control oxidative stress.
ROS such as superoxide anion, hydrogen peroxide (H₂O₂), and hydroxyl radical (OH⁻) are generated at low levels and play important roles in signaling pathways.[31]

Under normal conditions, they are regulated by endogenous antioxidants including SOD, glutathione peroxidase, glutathione, and catalase.[52] Overproduction of ROS and/or inactivation of antioxidants cause tissue injury from oxidative damage.[31] Oxidative stress can play important roles in the pathogenesis of early brain injury after SAH.[31] Mitochondria disruption, the production of hydroxyl radicals from extravasated hemoglobin, and disruption of the intrinsic antioxidant systems have all been reported in either experimental or human SAH.[53,54] O₂⁻ production was observed 1 h after SAH, and overexpression of SOD1 inhibited the production and reduced apoptotic cell injury after SAH.[31] The reduction in oxidative stress by SOD1 overexpression attenuated early brain injury after SAH via activation of Akt.[31]

**DNA damage**

p53 is a tumor suppressor gene involved in the regulation of apoptosis.[50] Responding to cell damage, p53 upregulates pro-apoptotic molecules including Bax, p53-upregulated modulator of apoptosis, and Bid, and downregulates antiapoptotic molecules Bcl-2 and Survivin.[55] p53 is upregulated after an ischemia insult and induces mitochondrial damage and activation of caspases.[56] It was reported that in SAH, p53 is one of the key factors in neuronal cell death. p53 was upregulated both at 24 and 72 h after SAH, and p53 inhibitor decreased brain edema and neuronal cell death.[54,57,58]

**Extrinsic pathway of apoptosis**

The death receptors, which are located on the cell surface, are involved in the extrinsic apoptosis pathway.[50] The receptor ligands expression, including Fas and tumor necrosis factor (TNF), are upregulated after cerebral ischemia.[59,60] The death receptors can activate caspase-8 or -10, which then directly activate caspase-3 or cause Bid/Bax activation, inducing cytochrome C release.[61] Moreover, forkhead transcriptional factors were activated after cerebral ischemia and then expression of Fas ligand increased, resulting in neuronal cell death.[62]

However, little is known regarding the relationship between early brain injury and death receptors or their ligands, whereas TNF-a was upregulated after SAH.[63]

**Therapeutic modalities**

For evaluating neuronal apoptosis in early brain injury after SAH, neurological examination should be performed to examine the outcome of neuronal cell injury. These molecular apoptotic pathways in neurons may induce brain edema, neurological deficit, and higher mortality. Previous studies showed that apoptotic-related pathway modulation by treatment could improve the outcome in early brain injury after SAH.

**Conclusions**

Recent studies have demonstrated the apoptosis mechanism in cerebral ischemia, whereas relatively few have studied the relationship between apoptosis and SAH, especially in early brain injury. It would be very helpful to study the relationship between SAH and another apoptotic mechanism, including autophagy and endoplasmic reticulum stress, which may lead to novel therapies in early brain injury.

Studies regarding early brain injury after SAH are limited, and further studies are needed for the clarification of the exact mechanism. For example, MAPKs, including ERK, JNK, and p38, were reported to induce apoptosis in the brain and cerebral artery after SAH,[62] whereas it has reported that ERK phosphorylation induced a beneficial effect on cerebral vasospasm.[64]

It is suggested that elevated ERK phosphorylation blocks apoptosis by enhancing the antiapoptotic protein Bcl-2 via CREB activation in cerebral ischemia.[60] The opposite effects may depend on the localization in the brain including neurons, glia, and endothelial cells.

In conclusion, apoptosis may play an important role in early brain injury after SAH. Further studies regarding apoptosis may lead to the development of new therapies and the improvement of outcome of SAH patients.

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Palade, et al.: A apoptosis and the mechanisms of subarachnoid hemorrhage

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