MINI-SYMPOSIUM: Role of the Inflammasome in Brain Pathogenesis: a Potential Therapeutic Target?

A brain in flame; do inflammasomes and pyroptosis influence stroke pathology?

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Abstract

Stroke is one of the leading causes of death and disability worldwide. Inflammation plays a key role across the time course of stroke, from onset to the post-injury reparative phase days to months later. Several regulatory molecules are implicated in inflammation, but the most established inflammatory mediator of acute brain injury is the cytokine interleukin-1. Interleukin-1 is regulated by large, macromolecular complexes called inflammasomes, which play a central role in cytokine release and cell death. In this review we highlight recent advances in inflammasome research and propose key roles for inflammasome components in the progression of stroke damage.

INTRODUCTION

Stroke is the second leading cause of death worldwide, affecting approximately 17 million people per annum (26). Of those affected by stroke, around two thirds either die or are left disabled (17). Two major subtypes of stroke exist and are defined by their etiology. Ischemic stroke occurs through embolic or thrombotic occlusion of a cerebral vessel, resulting in insufficient oxygen and glucose delivery required for cellular respiration. Hemorrhagic stroke occurs when a diseased or abnormal vessel ruptures, leading to the development of a neurotoxic hematoma within the brain. Ischemic strokes are most prevalent, accounting for 87% of all cases, with spontaneous non-traumatic intracerebral hemorrhage (ICH) being the second most common subtype of stroke, accounting for around 12% of all stroke cases worldwide (25). Roughly 40% of ICH cases prove fatal and, of those that survive, less than 40% live independently at 12 months (4). It is becoming increasingly clear that inflammatory processes are inherent across the time course of stroke, from endothelial activation shortly after stroke onset (in both ischemic and hemorrhagic stroke) through to the post-injury reparative phase days to months after stroke (41, 71). During the acute phase of stroke, sterile inflammation (ie, inflammation processes that occurs in the absence of pathogens) leads to the production of pro-inflammatory cytokines, such as interleukin-1 (IL-1), which, in turn, propagate the neuroinflammatory response through activation of resident cells (ie, microglia) and recruitment of inflammatory cells. Neutrophils are the first cells to migrate into the brain after stroke, followed by macrophages and lymphocytes a few days after injury (34, 46). Moreover, inflammatory cell activation is not contained within the necrotic core of ischemic brain tissue but, instead, is observed within the peri-infarct region. The size of this "inflammatory penumbra" positively correlates with the extent of brain damage and, consequently, is a key target for therapeutic intervention (33).

Minutes to hours after stroke, cell death leads to the induction of an inflammatory response through the release of danger signals termed damage associated molecular patterns (DAMPs). DAMPS are endogenous components which, under normal conditions, are either inert or sequestered intracellularly and thus not detected by immune cells. Examples of DAMPs released from dying cells include the chromatin-associated protein HMGB1, purine metabolites such as ATP, heat shock proteins and uric acid. In addition to DAMPs, some intracellular stores of pro-inflammatory cytokines can be released by necrotic cells such as IL-1 α (14). DAMPs trigger inflammatory responses through the activation of pattern recognition receptors (PRRs) present on or within innate immune cells. The activation of some cytosolic PRRs leads to the formation of large multiprotein complexes called inflammasomes. Inflammasomes are a type of intracellular PRR that combine pathogen associated molecular patterns (PAMPs) or DAMPs recognition to the activation of caspase-1. The presence of oligomerization domains allows the inflammasome sensor molecules to seed large quaternary structures, which ultimately concentrate caspase-1 peptides leading to auto-cleavage and activation (87). The majority of inflammasomes contain an N-terminal pyrin domain, which recruits the adaptor molecule apoptosis-associated

Table 1. Evidence for inflammasomes in stroke pathology.

Inflammasome	Evidence of involvement in stroke	Cell type	Stroke model
NLRP3	Increased mRNA and/or protein expression in stroke brains	Neurons (58, 97), microglia (54, 66)	Clinical ischemic brain tissue (97) Rat and mouse tMCAo (23, 54, 58, 61, 66, 91, 97, 100, 101) Mouse eMCAo (43) Mouse autologous blood injection (70, 96, 98) Rat collagenase injection (27) Rat and mouse endovascular perforation (15, 20, 62, 88)
	Smaller infarcts, edema, BBB breakdown, and neurological deficits in NLRP3 ^{-/-} mice	Microglial and endothelial cells	Mouse tMCAo (95)
	Infiltrating leukocytes express high levels of NLRP3 and inhibition of the inflammasome lessens brain injury	Infiltrating leukocytes	Mouse tMCAo (44)
NLRP1	Increased mRNA and/or protein expression in stroke brains	Neurons	Rat and mouse tMCAo (23, 58)
	The NLRP1 inflammasome complex was shown to form in neurons, whilst a neutralizing antibody reduces infarct size	Neurons, Astrocytes, Microglia	Mouse thromboembolic stroke (1)
AIM2	mRNA upregulation in the peri-infarct region	NA	Rat tMCAo (58)
	Smaller infarcts, less neurological deficits and dampened neuroinflammation in AIM2 ^{-/-} mice	NA	Mouse tMCAo (19)
NLRC4	mRNA upregulation in the peri-infarct region	NA	Rat tMCAo (58)
	Smaller infarcts, less neurological deficits and dampened neuroinflammation in NLRC4 ^{-/-} mice	NA	Mouse tMCAo (19)

NOD- LRR- PYD-containing (NLRP), Absent in melanoma 2 (AIM2), NOD- LRR- CARD-containing 4 (NLRC4), transient (t), embolic (e), permanent (p) middle cerebral artery occlusion (MCAo), blood brain barrier (BBB).

speck-like protein containing a CARD (ASC) through homotypic pyrin domain interactions (87). In turn, ASC forms a large speck structure consisting of multimeric ASC dimers and recruits caspase-1 to the assembling inflammasome complex via a C-terminal CARD (28). Caspase-1 then processes pro forms of the cytokines IL-1 β and IL-1 β , and facilitates their release through a caspase-1 dependent form of cell death termed pyroptosis (18). Each inflammasome senses PAMPs or DAMPs through unique, direct or indirect, interactions, allowing a multitude of molecular patterns to be sensed and acted on, some of which are implicated in chronic and acute brain injury (59, 90).

As a role for IL-1 in stroke damage is discussed at length elsewhere (12), here we focus on the role of inflammasomes in the inflammatory and cell death processes seen in stroke.

EVIDENCE FOR THE INVOLVEMENT OF INFLAMMASOMES IN STROKE

Of the five known inflammasome forming proteins, to date, four have been cited in stroke literature, namely NOD-, LRR- and PYD-containing 1 (NLRP1), NLRP3, absent in melanoma 2 (AIM2), and NOD-, LRR- and CARD-containing 4 (NLRC4). It is important to note NLRP3 is unique amongst inflammasomes in that, at

basal levels, its expression is insufficient to respond to activating stimuli (6). Thus, similar to IL-1 β the NLRP3 inflammasome has a two-step processing system and must receive an initial priming signal which instigates NF κ B signaling. The second signal is provided by stimuli that are unique to each inflammasome (discussed later) and cause the assembly of the inflammasome structure. Therefore, when critically analyzing inflammasome activity within tissues it is essential to have a true readout of inflammasome complex formation.

There are a number of studies highlighting upregulation of inflammasome components in various animal models of stroke (see Table 1). At the mRNA and protein level NLRP3 is significantly upregulated 12 hours after transient middle cerebral artery occlusion (tMCAo) in mice, peaking at 24 hours and remains elevated beyond 48 hours (95). Using the same model of stroke another group blotted for protein NLRP3, NLRP1 and the adaptor ASC, and found all to become elevated within one hour and remain above sham levels for 24 hours (97). Within the ischemic penumbra of rats subjected to tMCAo there is a steady rise in NLRP3, NLRP1, NLRC4, AIM2 and ASC mRNA over time, which remained at higher levels than shams up to 72 hours post-surgery (58). In the permanent embolic MCAo (eMCAo) mouse model of stroke NLRP3 is expressed within the ischemic penumbra within

24 hours (43) and in a thromboembolic occlusion mouse model, ASC protein levels are significantly raised in cortical lysates 7 days post-stroke (1). Data from both ICH and subarachnoid hemorrhage (SAH) stroke models indicate that the NLRP3 protein is upregulated at 12 hours and remains elevated beyond 24 hours (15, 20, 27, 70). Clearly, NLRP3 is the most cited inflammasome sensor molecule in pre-clinical stroke research and has also been shown in the brain tissue of stroke patients (97). Whilst the expression of inflammasomes within the stroke brain does not indicate causative effects, it does highlight the presence of sensor molecules in a pathology heavily reliant on IL-1 signalling. Some studies have taken inflammasome biology further by identifying the cellular distribution of inflammasomes in the brain post-stroke.

The NLRP1 molecule was the first inflammasome sensor to be imaged within murine ischemic brain tissue, where it was found to be ubiquitously expressed in neurons and astrocytes, but specifically upregulated in microglia within 6 hours of thromboembolic stroke (1). In mice, NLRP3 has been localized to microglia and the endothelium of ischemic and hemorrhagic brain tissue (70, 95), whilst in rat tMCAo models, it was found to be upregulated in neurons (58). The most recent study concerning inflammasomes in stroke used flow cytometry to identify cells expressing NLRP3 (44). In this study the authors used fluorescence activated cell sorting and found that NLRP3 mRNA was specifically expressed within infiltrating myeloid cells. It is well established that inflammasomes are functional in many cell types relevant to stroke—see review (60)—though functional readouts for inflammasome activation in the brain are currently lacking.

Using size exclusion chromatography of ischemic brain lysates, an NLRP1 inflammasome complex was found in fractions above 600 kDa, along with the interacting molecules ASC, XIAP and caspase 1 (1). NLRP1 was the first inflammasome to be characterized and is unique in its double death domain structure. The most established activator of NLRP1 is the Bacillus anthracis virulence factor and metalloproteinase, anthrax lethal toxin, which acts by cleaving the N-terminal domain, causing oligomerization and caspase-1 activation. It is suggested NLRP1 is present in neurons in a pre-assembled state that is sensitized to neuronal stress, leading to activation under metabolic disruption (90). Interestingly, glucose starvation and hypoxia (both conditions of paramount importance to ischemic stroke pathology) have been shown to activate NLRP1 through intracellular ATP depletion (64). In a thromboembolic stroke model, the application of an anti-NLRP1 antibody inhibited the formation of the NLRP1 inflammasome and reduced brain lesion volume by 18% (1). Since this finding in 2009 no further studies have addressed NLRP1 activation in stroke and focus has turned toward characterizing stroke pathology in transgenic animals.

NLRP3^{-/-} mice subjected to tMCAo are protected from ischemic damage, with better functional outcomes and reduced inflammatory pathology (95). Lesion volumes and neurological deficit scores were reduced by around 40% in NLRP3 deficient mice, correlating with less endothelial damage and blood brain barrier (BBB) breakdown. This is contrary to results from Denes *et al* (19) which suggest NLRP3 does not contribute to MCAo injury in mice. The discrepancy between these results could be explained by differences in occlusion time causing modifications of the inflammatory response. Bruton's tyrosine kinase (BTK) physically associates with NLRP3 and ASC, and these interactions are essential for

inflammasome activation in response to multiple stimuli (44). Both BTK and NLRP3 are expressed within infiltrating myeloid cells present in the ischemic penumbra 1 day post-stroke. Moreover, pharmacological inhibition of BTK activity reduces lesion volumes in an IL-1B specific manner. This finding suggests that infiltrating leukocytes express NLRP3 and that this directly contributes to ischemic brain injury. Many of the NLRP3 activating DAMPs are present within the ischemic brain. Oxygen depletion leads to a metabolic switch toward glycolysis, causing lactic acid and proton build up and subsequent extracellular acidosis. The NLRP3 inflammasome is able to sense acidic conditions through an uncharacterized mechanism leading to K+ efflux and formation of the inflammasome complex (79). Similarly, ATP is released by dying cells and is a well-established activator of the NLRP3 inflammasome through P2X7R activation. The latter mechanism has been shown to be an important facet of hemorrhagic stroke. The P2X7R antagonist brilliant blue G is protective in both SAH and ICH models, resulting in smaller lesions, reduced neuronal death, and fewer infiltrating neutrophils (15, 27). One of the most injurious aspects of hemorrhagic stroke is the toxicity of parenchymal blood components (3). Interestingly, the red blood cell breakdown product haem activates the NLRP3 inflammasome through the production of mitochondrial ROS (21). In ICH, genetic deletion of the haem scavenging protein hemopexin accentuates brain injury, whereas, mitochondrial ROS scavengers reduce brain injury through NLRP3 downregulation (69, 70). A major predictor of mortality in ischemic and hemorrhagic stroke patients is the development of cerebral edema. Cytotoxic edema occurs foremost through accumulation of intracellular ions causing influx of water molecules. This cell swelling may also be a mechanism by which NLRP3 becomes active within the brain, as hypotonic environments are well established activators of the inflammasome (16). Common amongst studies investigating NLRP3 in preclinical stroke models is the lack of preexisting stroke co-morbidities. The co-morbidities commonly linked with stroke, such as obesity and atherosclerosis, have significant NLRP3-dependent inflammatory changes (12). Whether these inflammatory changes systemically prime the NLRP3 inflammasome and affect stroke outcomes is unknown. However, it is known that hypertensive rats have a higher number of cells expressing NLRP3 in the brain compared with normotensive counterparts (66).

Our group have shown that both absent in melanoma 2 (AIM2) and NLRC4 contribute to damage after ischemic stroke, although the mechanisms leading to their involvement remain elusive (19). When subjected to MCAo, infarct volumes were significantly smaller in NLRC4^{-/-} and AIM2^{-/-} mice compared with wild type (WT) C57BL/6 mice. NLRC4^{-/-} and AIM2^{-/-} mice showed a dampened immune response to stroke, with fewer active microglia and leukocytes observed within the ipsilateral hemisphere. AIM2 is a cytosolic double-stranded DNA (dsDNA) sensor that can react to both pathogenic and endogenous dsDNA. The AIM2 sensor molecule is composed of a pyrin domain and a DNAbinding HIN domain, and thus recruitment of caspase-1 to the inflammasome complex requires ASC (29, 40). How AIM2 becomes active during stroke is unclear. However, AIM2 contributes to autoinflammatory disease through recognition of endogenous dsDNA. Moreover, similar to NLRP3, the AIM2 inflammasome has recently been described as a sensor of glycolysis and could contribute to stroke damage through this pathway (93).

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NLRC4, on the other hand, possesses a CARD domain and can directly interact with procaspase-1 to cause cell death, although the presence of ASC is required for cytokine processing (72). Bacterial PAMPs such as flagellin, and rod and needle units of bacterial type III secretion systems are the most cited activators of the NLRC4 inflammasome (30). NLRC4 responds to these exogenous molecules indirectly through interacting proteins termed NLR-family, apoptosis inhibitory proteins (NAIPs). It is currently unclear whether NAIPs can respond to endogenous molecules, although NLRC4 is known to respond to hyperosmotic conditions (42), which may be present in the post-stroke brain. Interestingly, NLRC4 does not respond to necrotic cell death (45) but its role in sterile inflammation is becoming increasingly apparent, with reported links to acute coronary syndrome (47), breast cancer progression and diabetic nephropathy (53, 99).

Caspase-1 activation after stroke

Studies evaluating caspase 1 activity in the stroke affected brain also give some insight into inflammasome activity in the disease. Following a permanent model of stroke, the proteolytic activity of caspase-1 is increased 30 minutes after occlusion with a second wave of activation 12 hours later (7). The increase of caspase-1 expression has been described in neurons and astrocytes after thromboembolic stroke and observed later in microglia (24 h post stroke) (1). Other studies have demonstrated a role of caspase-1 after stroke by using transgenic mice. Caspase-1 knockout mice and mice expressing a dominant-negative caspase-1 exhibit a reduction of brain damage compared with wild type after experimental stroke (32, 86). Moreover, intra-cerebroventricular administration of Ac-YVAD-cmk or VRT-018858, both caspase-1 inhibitors, provides protective effects in experimental stroke models

(38, 78, 83). Beneficial effects of caspase-1 inhibition are also observed in a model of oxygen and glucose deprivation in rat organotypic hippocampal slices (81). Altogether, these studies demonstrate a key role of caspase-1 during stroke.

Caspase 11 and non-canonical inflammasome after stroke

In addition to the canonical mode of inflammasome activation, caspase-4 and -5 in humans and caspase-11 in mice can participate in IL-1/IL-18 processing and cell death, through the "noncanonical inflammasome" (36, 80). Initial studies showed that activation of caspase-11 promoted subsequent activation of caspase-1 and -3 and that caspase-11 knockout mice exhibited markedly reduced apoptosis following a permanent model of MCAO (49). In more recent studies, caspase-11 has been described to detect intracellular LPS and promote cell death and the release of IL-1 α (37, 52). Caspase-11 can also lead to caspase-1 dependent IL-1 β release and caspase-1 independent pyroptosis via direct cleavage of gasdermin D (GSDMD—see below) by caspase-11 (51). However, the precise role of caspase-11 in the context of sterile inflammation is not well understood and further studies are required in order to fully delineate the role of the non-canonical inflammasome during stroke.

IL-1 INDEPENDENT ROLES OF INFLAMMASOMES IN STROKE

Pyroptosis

As shown in Figure 1, one of the defining features of inflammasome activation is the induction of a pro-inflammatory programmed

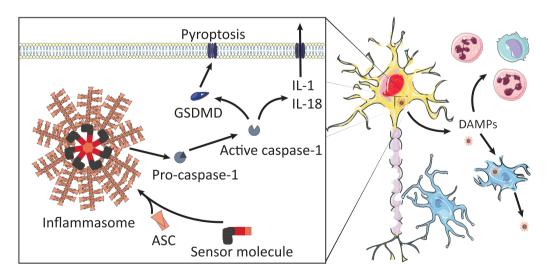


Figure 1. The detection of pro-inflammatory stimuli leads to the formation of macromolecular inflammasome complexes. Sensor molecules react to an array of damage associated molecular patterns (DAMPs) by recruiting a scaffold protein termed apoptosis-associated speck forming protein containing a CARD (ASC). ASC recruits the zymogen caspase-1 causing auto-cleavage and activation. Active caspase-1 imparts a pro-inflammatory lytic cell death termed pyroptosis through cleavage of gasdermin D (GSDMD). The N-terminus of

GSDMD forms pores in the plasma membrane which allows passage of proinflammatory cytokines, such as interleukin (IL-) 1 and 18 that are also processed by inflammasomes. During stroke, inflammasomes contribute to brain injury through the spread of inflammation, leading to microglial activation and ingress of circulating immune cells. Inflammasomes themselves can be released during pyroptosis and propagate an inflammasome activation cascade within tissue.

cell death termed pyro- (fire/fever) ptosis (falling). The two hallmark features of pyroptosis are DNA damage and membrane rupture, the former akin to apoptosis whilst the latter in complete contrast (8). Another distinction from apoptosis is the initiation of pyroptosis through the inflammatory caspases, namely caspase 1 or caspase 4/5/11, and the subsequent release of pro-inflammatory cytosolic components. The process of pyroptosis is essential for the clearance of pathogens *in vivo* and has recently been shown to retain bacteria within intracellular traps during the lytic cell death process (48). With pyroptosis strongly linked with IL-1 signaling, it may well have emerged as a vehicle to disseminate the now termed canonical DAMPs (73). We know these canonical DAMPs play implicit roles during sterile inflammation, yet it is still unclear how pyroptosis is involved in stroke.

The caspase 1/11 dependent cell death process may contribute enormously to damage post-stroke independently of IL-1 signaling. Caspase $1/11^{-/-}$ mice have reduced infarcts after MCAo, yet IL-1β deficient mice are not protected (10, 86). Pyroptosis may be the principal cell death mechanism of neurons within the ischemic core during stroke. Neurons are the earliest source of caspase 1 activation, where it has been identified to precede pro-apoptotic caspase 3 activity (7). Moreover, the expression of a neuronal dominant negative caspase 1 is known to lessen stroke brain injury (32). The pyroptosome forming ASC protein has also been shown to substantially contribute to acute brain injury, with knockout mice exhibiting reduced infarct volumes and improved neurological outcome independently of IL-1B levels (19). However, pyroptosis can occur irrespective of ASC pyroptosome assembly (13), thus the true extent of pyroptotic cell death cannot be inferred solely through the use of ASC-deficient mice. Despite strong evidence implicating multiple inflammasomes in CNS disease (90), there is no conclusive evidence describing pyroptosis within the brain. More studies are required to evaluate the activity of caspase 11 and GSDMD during acute brain injury.

In 2015 multiple articles were released which described the previously uncharacterized GSDMD protein as an effector of pyroptosis (39, 50, 89). Cells lacking GSDMD were unable to undergo pyroptotic cell death or secrete canonically processed IL-1β. Kayagaki et al (2015) found that, in the absence of GSDMD, IL-1B failed to be processed when LPS was delivered intracellularly, suggesting GSDMD may also initiate non-canonical inflammasome formation (50). Cleavage of GSDMD releases an N-terminal fragment which has recently been shown to associate with the plasma membrane and form 13 nm pores (2, 67, 85). Interestingly, Shi et al (2015) showed GSDMD to be the effector of pyroptosis for all of the previously mentioned stroke-related inflammasomes bar NLRP1 (89), suggesting that inflammasomes formed within the brain induce pyroptosis through GSDMD. However, it is currently unclear if pyroptosis occurs within brain cells at all and whether it would follow the same route as described for myeloid cells.

Inflammasome dependent protein release

On top of the processing of IL-1 β , IL-18 and GSDMD, caspase 1 activation also facilitates the release of other DAMPs. In fact, the principal output of caspase 1 activation may indeed be the induction of pyroptosis and the subsequent release of DAMPs. Genetic knockout of IL-1 β in mice still renders the species susceptible to endotoxemia (24). Whereas caspase $1/11^{-/-}$ mice given an 800 μ g

dose of LPS do not enter septic shock and have a far greater survival rate than their WT counterparts (63). More recent evidence suggests caspase 11 is crucial for LPS induced sepsis and this may occur through HMGB1 signalling (51, 57, 92). HMGB1 released during pyroptosis is hyperacetylated at its nuclear localization sequence (68), which has previously been shown to cause cytoplasmic redistribution and facilitation of its secretion (9). Interestingly, HMGB1 levels are elevated in stroke patient serum and neutralizing antibodies are protective in different experimental models of stroke (65, 75).

Unlike IL-1 β , IL-1 α is biologically active in its pro-form and its release is heavily dependent of cell death (22, 35). Despite lacking a caspase-1 cleavage site, IL-1α secretion can be triggered through inflammasome activation (35). In contrast to IL-1β, inflammasome dependent IL-1α release does not require membrane permeabilization (35, 74), suggesting that GSDMD cleavage is dispensable for its secretion. Despite this, IL- 1α secretion is proposed to heavily rely on caspase-1 dependent processes when; (i) the stimulus does not cause intracellular calcium levels to rise, (ii) IL- 1α is sequestered by the cytosolic IL-1RII (35, 77). Studies have shown there to be a substantial reduction in the amount of IL-1 α released from caspase1/11 -/- macrophages and monocytes stimulated with LPS (56, 63), and this reduction in IL-1 α release is likely to be caspase 11 dependent (51). IL-1α expression precedes IL-1β in ischemic brain tissue and is known to be a critical mediator of acute brain inflammation, which is extensively reviewed elsewhere (11, 76). HMGB1 and IL-1a posses the potential to cause a positive NLRP3 inflammasome signalling loop within the parenchyma post-stroke, with the former associated with inflammasome activation and the latter in priming (94).

Self-propagating inflammasomes

ASC speck formation may have pleiotropic roles in the propagation of inflammation, the best characterized being the aforementioned processing of zymogens and the induction of pyroptosis. There is some evidence to suggest the supramolecular ASC assembly is able to stably interact with hydrophobic molecules and may act to sequester bacterial PAMPs or endogenous DAMPs and deliver these pro-inflammatory signals to neighboring phagocytes, which in the brain would be microglial cells (84).

During pyroptosis the inflammasome complex is released into the extracellular milieu where the platform can potentiate tissue inflammation (5, 31). Using in vitro assembled ASC specks, two groups have shown that specks are extremely stable and can activate extracellular caspase 1. Within the necrotic regions of the damaged stroke brain this would enable activated inflammasomes to capture any unprocessed zymogens released during necrosis, maximizing the inflammatory response in the process. ASC specks were described as having "prionoid" activity by Franklin et al (31) as they were shown to nucleate inflammasome formation when ingested by phagocytic cells. As previously discussed, there is evidence for inflammasome activation in stroke and ASC specks have been quantified within the ischemic penumbra (55). This data begs the question; what role do extracellular inflammasome complexes play in the stroked brain? Interestingly, there is evidence that inflammasome targeting antibodies lessen neuronal damage after stroke and traumatic brain injury in mice (1, 82). As antibodies cannot enter the cytosol of live cells, these neutralizing antibodies

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may be limiting the acute spread of inflammation via effects on extracellular inflammasomes. Both inflammasome complexes and anti-ASC antibodies have been identified in the serum of patients suffering with inflammatory illnesses (5, 31), suggesting ASC specks can have long-lasting physiological effects. As ASC is an implicit mediator of stroke damage, it may also propagate an acute injury into chronic inflammation, possibly contributing toward inflammatory-driven post-stroke sequelae such as depression and cognitive decline.

CONCLUSION

Sterile inflammation contributes enormously to brain damage during stroke. Inflammasomes, which are central effectors of sterile inflammation, are expressed and activated after stroke leading to IL-1 processing and cell death through pyroptosis. The role of IL-1 in the pathogenesis of stroke is well described, however inflammasome-induced cell death in acute brain injury is less understood. Moreover, release of inflammasome complexes could represent an inflammasome dependent inflammatory drive after stroke. Further understanding of the molecular mechanisms of inflammasome activation within the brain after stroke is required in order to elucidate therapeutic targets. The development of therapeutic strategies to target inflammasomes will, one day, provide new opportunities in the management of stroke.

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