Cyclooxygenase-2 inhibitor blocks the production of West Nile virus-induced neuroinflammatory markers in astrocytes

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Inflammatory immune responses triggered initially to clear West Nile virus (WNV) infection later become detrimental and contribute to the pathological processes such as blood-brain barrier (BBB) disruption and neuronal death, thus complicating WNV-associated encephalitis (WNVE). It has been demonstrated previously that WNV infection in astrocytes results in induction of multiple matrix metalloproteinases (MMPs), which mediate BBB disruption. Cyclooxygenase (COX) enzymes and their product, prostaglandin E2 (PGE2), modulate neuroinflammation and regulate the production of multiple inflammatory molecules including MMPs. Therefore, this study determined and characterized the pathophysiological consequences of the expression of COX enzymes in human brain cortical astrocytes (HBCAs) following WNV infection. Whilst COX-1 mRNA expression did not change, WNV infection significantly induced RNA and protein expression of COX-2 in HBCAs. Similarly, PGE2 production was also enhanced significantly in infected HBCAs and was blocked in the presence of the COX-2-specific inhibitor NS-398, thus suggesting that COX-2, and not COX-1, was the source of the increased PGE2. Treatment of infected HBCAs with NS-398 attenuated the expression of MMP-1, -3 and -9 in a dosedependent manner. Similarly, expression of interleukin-1 β , -6 and -8, which were markedly elevated in infected HBCAs, exhibited a significant reduction in their levels in the presence of NS-398. These results provide direct evidence that WNV-induced COX-2/PGE2 is involved in modulating the expression of multiple neuroinflammatory mediators, thereby directly linking COX-2 with WNV disease pathogenesis. The ability of COX-2 inhibitors to modulate WNV-induced COX-2 and PGE2 signalling warrants further investigation in an animal model as a potential approach for clinical management of neuroinflammation associated with WNVE.

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INTRODUCTION

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West Nile virus (WNV), a mosquito-borne flavivirus that causes lethal encephalitis, has emerged as a significant cause of viral encephalitis in the USA (Brinton, 2002). In a small subset of cases, WNV targets the central nervous system (CNS), clinically progressing to meningitis, encephalitis or acute flaccid paralysis syndrome, leading to mortality in 10 % of hospitalized patients and complicated neurological sequelae in some who survive (Sejvar *et al.*, 2003). WNV-associated encephalitis (WNVE) in the mouse model is characterized by disruption of the blood–brain barrier (BBB), neuroinflammation, microglial activation and loss of neurons (van Marle *et al.*, 2007; Wang *et al.*, 2004, 2008). Inflammation in the CNS is a major hallmark of WNVE in mice and is associated with a dramatic

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increase in several pro-inflammatory cytokines such as tumour necrosis factor alpha (TNF- α) and interleukin (IL)-1 β and -6 (Garcia-Tapia *et al.*, 2007; Wang *et al.*, 2004) and chemokines such as CCL2 and CXCL10, which regulate leukocyte trafficking into the brain (Glass *et al.*, 2006; Klein *et al.*, 2005; Lim *et al.*, 2006). Although many of these virus-induced cytokines and chemokines play a critical role in the recruitment of virus-specific T cells and virus clearance in the mouse brain, increased production of pro-inflammatory cytokines also contributes to the overall disease pathogenesis.

In the CNS, neurons are the prime target for WNV infection; however, infection of non-neuronal CNS cells such as astrocytes and BBB endothelial cells has been documented (Cheeran *et al.*, 2005; Shrestha *et al.*, 2003; van Marle *et al.*, 2007; Verma *et al.*, 2009). Activation of glial cells along with loss of neurons is considered a key pathogenic feature in WNV infection in humans (Kelley *et al.*, 2003). Although virus infection in human glial cells is

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not as robust as in neurons, they secrete much higher levels of immune mediators such as chemokines (CXCL10, CCL2 and CCL5) and cytokines (Cheeran *et al.*, 2005; Glass *et al.*, 2005). Furthermore, cytotoxic factors secreted from WNVinfected astrocytes can induce bystander death of naïve neurons (van Marle *et al.*, 2007). Our previous data also demonstrated that WNV infection of human astrocytes results in the induction of multiple matrix metalloproteinases (MMPs), which are capable of degrading the tight junction proteins of human brain microvascular endothelial cells, thereby compromising the integrity of the BBB model (Verma *et al.*, 2010). Thus, although astrocytes are one of the key players in WNV-induced neuroinflammatory responses, the upstream events modulating these inflammatory responses are not well understood.

Prostaglandin E2 (PGE2), the most abundant prostaglandin in the brain, is considered to play an essential role as a local regulator of pathogenic processes in several neurodegenerative diseases (Bazan et al., 2002; Candelario-Jalil & Fiebich, 2008; Hickey et al., 2007). Cyclooxygenase enzymes (COX-1 and -2) catalyse the committed step in the conversion of arachidonic acid to PGE2. COX-1 is expressed ubiquitously and is considered to be an isoform responsible for homeostatic prostaglandin synthesis. By contrast, COX-2 is rapidly induced in many cell types including astrocytes in response to inflammatory stimuli (Bazan, 2001; Bazan et al., 2002; Tzeng et al., 2005). However, in the CNS, COX-2 is also expressed constitutively in the hippocampal neurons (Yang & Chen, 2008). Accumulating evidence suggests that of the two COX isoforms, COX-2, but not COX-1, plays a crucial role in inflammation and disease pathogenesis. PGE2 activates several downstream inflammatory pathways via autocrine or paracrine mechanisms, resulting in the induction of pro-inflammatory mediators (Ferri & Ferguson, 2005; Kyrkanides et al., 2002). Multiple downstream effects of COX-2/PGE2 include induction of chemotactic cytokines, mediators of BBB disruption such as MMPs and plasmin/ urokinase plasminogen activator (uPA), apoptotic death and activation of microglia (Bazan, 2001; Bazan et al., 2002; Im et al., 2006). Both MMPs and the plasmin/uPA system belong to the family of multi-domain zinc-containing serine proteases, and their increase in glial cells has been associated with BBB disruption (Conant et al., 2004). Treatment with COX-2 inhibitors such as NS-398 in both in vitro and in vivo model systems can attenuate secretion of cytokines, cell-adhesion molecules, MMP-9 and uPA, and improve overall pathology associated with several neurodegenerative diseases (Im et al., 2006; Iwamoto et al., 2008; Ottino & Bazan, 2001; Pompl et al., 2003; Thomas & Kuhn, 2005). CNS infection of other neurotropic viruses such as human immunodeficiency virus (HIV) and Japanese encephalitis virus (JEV) also results in increased COX-2 and PGE2 production (Flora et al., 2006; Ghoshal et al., 2007).

In WNV infection, once the virus enters the brain and triggers inflammation, there is very little that can be done

to prevent disease progression. Therefore, understanding the key molecular mechanisms by which WNV modulates inflammation is of utmost importance in the design of effective treatments. The extent of the host innate immune response is critical in determining the end point of WNV infection, i.e. virus clearance versus morbidity. As the COX/PGE2 pathway has been demonstrated to modulate the host innate and inflammatory response in viral infections, a better understanding of this pathway in WNV neuropathogenesis is warranted. Therefore, the purpose of this study was to investigate the role of COX enzymes and their product, PGE2, in WNV infection using primary human brain cortical astrocytes (HBCAs) and to determine the mechanisms by which they contribute to neuroinflammation. Our results indicated that COX-2 and its product, PGE2, are significantly induced in WNVinfected HBCAs and that increased COX-2/PGE2 levels correlate with a peak in the virus titres. Using the COX-2specific inhibitor NS-398, we further demonstrated that the expression of MMPs and key inflammatory cytokines are tightly regulated by COX-2/PGE2 levels in WNV-infected astrocytes.

RESULTS

WNV infection induces COX-2 expression in HBCAs in a dose- and time-dependent manner

Profound inflammatory reactions including induction of MMPs, cytokines, chemokines and cell-adhesion molecules have been demonstrated in WNV-infected brain cells (Arjona et al., 2007; Glass et al., 2005; Wang et al., 2008). COX-2, an important mediator of inflammation, can modulate inflammatory reactions by inducing cytokines and MMPs. In the brain, astrocytes are an important source of COX-2 and inflammatory mediators in several neurodegenerative diseases. Our previous study demonstrated that WNV infection of primary HBCAs peaked at day 2 post-infection (p.i.) and remained high up to day 4 p.i., following which there was a decline in the virus titre (Verma et al., 2010). Therefore, in this study we analysed the mRNA and protein expression of COX enzymes in HBCAs at the same time points. The global response of HBCAs infected with WNV at an m.o.i. of 5 was determined at days 1 and 3 p.i. by cDNA microarray analysis. As observed in Table 1, WNV infection did not alter the expression profile of the COX-1 gene in HBCAs at either time point, whereas increased expression of the COX-2 gene was observed at day 1 and peaked further at day 3 p.i. An increase in the expression of COX-2 was further validated by qRT-PCR from day 1 to day 4 p.i. in HBCAs infected with WNV at an m.o.i. of 1 and 5. As observed in Table 1, WNV infection at an m.o.i. of 1 induced a time-dependent increase in the mRNA expression of COX-2 from day 1 to day 4 p.i. This increase was first detected at day 1, increased sharply at day 2 and remained high up to day 4 p.i. In concurrence with the Table 1. mRNA fold changes in COX-1 and -2 levels in WNV-infected HBCAs compared with uninfected controls as determined by cDNA microarray analysis and qRT-PCR

Days p.i.	COX-1	COX-2					
	cDNA microarray	cDNA microarray	Real-time qRT-PCR				
			m.o.i. 1	m.o.i. 5	UV-WNV		
Day 1	1.01	55	27	42	1.27		
Day 2	ND	ND	55	87	2.1		
Day 3	0.47	102	83	93	1.1		
Day 4	ND	ND	76	97	1.6		

UV-WNV, UV-inactivated WNV; ND, not done.

microarray data, in HBCAs infected with WNV at m.o.i. of 5, a gradual increase in COX-2 gene expression was observed from day 1 to day 3 p.i. Moreover, infection with UV-inactivated WMV did not induce COX-2 expression (Table 1), suggesting that increased COX-2 expression is a result of active virus replication. Expression of COX-2 protein was characterized by Western blotting. As demonstrated in Fig. 1, although COX-2 protein expression in control HBCAs was minimal, a dramatic increase was observed in infected HBCAs at days 2 and 3 p.i., which coincided with high mRNA transcripts and a peak in virus replication.

We next analysed the increase in COX-2 protein expression by immunocytochemical analysis. Whilst mock-infected HBCAs did not exhibit a COX-2 fluorescence signal (Fig. 2a), a strong signal for COX-2 expression was observed at day 2 p.i. in WNV-infected HBCAs (Fig. 2b). Fig. 2(c) demonstrates WNV antigen staining in HBCAs, and, as depicted in Fig. 2(d), increased COX-2 immunostaining was observed mostly in WNV-infected cells. This was further confirmed by counting the number of COX-2- and WNV-positive cells. Based on the counting of approximately 1000 cells from a total of four coverslips, it was found that 79% of COX-2-positive cells were also positive for WNV antigen staining.



Fig. 1. WNV induces COX-2 protein expression in HBCAs. Cell lysates extracted from WNV-infected (m.o.i. 5) (I) and mock-infected control (C) HBCAs at days 2 and 3 p.i. were separated by PAGE, transferred onto PVDF membranes and immunoblotted with antibodies specific to COX-2. Equal loading was confirmed by reprobing with anti- β -actin antibody and the bands were detected by an ECL method.

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Treatment with COX-2 inhibitor prevents WNV-induced PGE2 production in HBCAs

As PGE2 is the main product of COX-2 and is released extracellularly, it was important to determine its release in the supernatant by ELISA. We observed a four- to fivefold increase in PGE2 levels in the supernatant of WNVinfected HBCAs at day 3 p.i. (P<0.01, Fig. 3). To further confirm the contribution of COX-2 as the main source of PGE2, infected HBCAs were treated with 25 and 50 uM concentrations of the COX-2-specific inhibitor NS-398, and PGE2 levels were assaved at the same time point. As seen in Fig. 3, the level of PGE2 released by infected HBCAs was significantly reduced in the presence of both doses of NS-398 (P<0.01, compared with infected HBCAs without NS-398 treatment). The inhibition was greater in cells treated with 50 µM NS-398, suggesting a dosedependent effect. HBCAs infected in the presence of vehicle control (0.001 % DMSO) did not exhibit any change in expression levels of COX-2 enzyme (data not shown), thus validating the specificity of NS-398 in blocking the COX-2 enzyme. This observation is similar to previous studies where the vehicle control for NS-398 had no effect on the COX-2 enzyme (Baek et al., 2007; Lu et al., 2008). These findings suggested that the increased levels of PGE2 observed in WNV-infected cells were indeed dependent on COX-2 activation and were not a result of cellular COX-1 activity, which is not inhibited by NS-398. Furthermore, NS-398 treatment did not affect the virus titres in infected HBCAs as measured by plaque assay (data not shown). Taken together, these results demonstrated that WNV infection induces COX-2 and PGE2 production in astrocytes.

WNV-induced COX-2/PGE2 regulates the expression of uPA and MMPs in HBCAs

We have demonstrated previously that WNV infection induces MMP-1, -3 and -9 in HBCAs (Verma *et al.*, 2010). As the COX-2/PGE2 signalling pathway is associated with the induction of MMPs, we assessed the ability of WNVinduced COX-2 and PGE2 to modulate MMP expression in HBCAs. The mRNA expression of MMP-1, -3 and -9 increased in HBCAs at day 3 p.i. (m.o.i. 5), thereby



Fig. 2. Localization of COX-2 in WNV-infected HBCAs using immunofluorescence. Confluent primary HBCAs were infected with WNV at an m.o.i. of 5 and stained with anti-COX-2 (red) or anti-WNV envelope (green) antibody and the nuclei counterstained with DAPI (blue). (a) Mock-infected HBCAs demonstrated faint and diffused COX-2 immunoreactivity. (b) A marked increase in COX-2 immunostaining was observed in infected HBCAs at day 2 p.i. (c) WNV-infected HBCAs stained for WNV antigen. (d) Strong COX-2 staining was observed mostly in WNV-infected HBCAs. (e) HBCAs incubated with secondary antibody only did not demonstrate any immunostaining. The images depicted are representative results of three independent experiments in duplicate. Bars, 10 μm.

corroborating our previous results. However, treatment of infected HBCAs with NS-398 drastically reduced the expression of MMPs in a dose-dependent manner (Fig. 4a). Compared with WNV-infected cells, treatment of infected HBCAs with NS-398 at a concentration of 50 uM decreased the mRNA expression of MMP-3 by 55 % and of MMP-1 and -9 by 70-75%. Furthermore, a similar inhibitory effect of NS-398 was also observed in the amount of MMP-3 and -9 secreted into the supernatant of infected HBCAs as assayed by ELISA. As observed in Fig. 4(b), whilst the levels of MMP-3 and -9 increased by fiveto sixfold (P<0.005) in the supernatant of infected HBCAs at day 3 p.i., the presence of NS-398 at concentrations of 25 and 50 µM decreased their levels significantly, thereby supporting the qRT-PCR data. These results collectively indicated that COX-2/PGE2 modulates the expression of MMPs in WNV-infected HBCAs. As the activation of uPA/



Fig. 3. HBCA were infected at an m.o.i. of 5 in the presence or absence of NS-398 and the supernatants collected at day 3 p.i. were assayed by ELISA for PGE2 production. The presence of NS-398 significantly reduced the levels of PGE2 released into the supernatant. The data expressed are the mean concentration \pm sD (pg ml⁻¹) of the amount of PGE2 secreted into the cell supernatant from at least four independent experiments in duplicate. *, *P*<0.01 compared with uninfected control; **, *P*<0.05 compared with infected cells.

uPA receptor (uPAR) system, which is associated with matrix degradation, is also governed by COX-2, we analysed the response of uPA gene expression to WNV infection. As depicted in Fig. 4(a), the expression of uPA increased by four- to fivefold in WNV-infected HBCAs compared with control cells and treatment of infected HBCAs with NS-398 almost completely blocked any increase in uPA mRNA.

Expression of inflammatory cytokines is governed by COX-2-derived PGE2

Increased production of inflammatory cytokines is another major neuropathological event associated with WNV infection. Therefore, we next investigated the possibility of COX-2 as one of the cellular factors responsible for modulating cytokine production in WNV-infected HBCAs. Increases in pro-inflammatory cytokines such as IL-1 β and -6 has been documented recently in WNV-infected astrocytes (van Marle et al., 2007). Here, we measured both mRNA and protein levels of key inflammatory cytokines such as IL-1 β , -6 and -8 induced by WNV in HBCAs in the presence or absence of NS-398. At an m.o.i. of 5, the levels of mRNA expression of IL-1 β , -6 and -8 increased by 5-11-fold at 24 h p.i. (data not shown), followed by a dramatic increase at day 2, which peaked further at day 3 p.i. (Table 2). This sharp increase in IL-1 β , -6 and -8 occurred at days 2 and 3 p.i. when the expression of COX-2 was also high (Fig. 1) and coincided with the peak in virus titres. However, as depicted in Table 2, treatment of HBCAs with NS-398 led to a 60-90 % reduction in mRNA expression of these cytokines at both time points. This decrease was more prominent in HBCAs treated with 50 µM NS-398 compared with treatment with 25 µM NS-398, suggesting that the reduction in cytokine expression is dose dependent. As cytokines are secretory proteins, their release into the supernatant of infected HBCAs treated with or without NS-398 was confirmed by ELISA. A significant increase in the levels of IL-1 β , -6 and -8 was observed in the supernatant from WNV-infected HBCAs at day 3 p.i.



Fig. 4. COX-2 inhibitor attenuates the expression of MMPs and uPA. (a) gRT-PCR was conducted on RNA extracted from HBCAs treated with or without NS-398 at day 3 p.i. (m.o.i. 5) to determine the fold change in MMP-1, -3 and -9 gene expression. Changes in the levels of each gene were first normalized to the glyceraldehyde-3-phosphate dehydrogenase (GAPDH) gene and the fold change in infected cells compared with corresponding control cells was calculated. Data represent the means \pm SD of at least four independent experiments conducted in duplicate. (b) The supernatant medium from WNV- and mock-infected HBCAs with or without NS-398 treatment was analysed for total MMP-3 and -9 levels by ELISA at day 3 p.i. WNV-induced MMP-3 and -9 were significantly reduced in the presence of 50 µM NS-398. Results are expressed as means \pm SD (pg ml⁻¹) of the amount of MMP-3 and -9 secreted into the cell supernatant in duplicate from at least three independent experiments. Mean comparisons were based on extrapolated confidence intervals for at least four data points. *, P<0.005 compared with uninfected control; **, P<0.05 compared with infected cells.

 Table 2. mRNA fold changes in cytokine levels as determined

 by qRT-PCR in WNV-infected HBCAs in the presence or

 absence of the COX-2-specific inhibitor NS-398

Gene	Day 2			Day 3		
	WNV	WNV + NS-398		WNV	WNV+NS-398	
		25 µM	50 µM		25 µM	50 µM
IL-1 β	81	18	9	187	68	32
IL-6	76	15	8	108	37	21
IL-8	48	8	3	137	22	11

(P<0.005), which decreased significantly in the presence of NS-398 (P<0.05, Fig. 5). These data suggested that COX-2 has a significant regulatory effect on the induction of inflammatory cytokines in WNV-infected cells and that blocking of COX-2 alone was sufficient to reduce the levels of key inflammatory cytokines released from the infected cells.

DISCUSSION

Neuroinflammation and disruption of the BBB are the common pathological features of WNVE (Sejvar *et al.*, 2003; Wang *et al.*, 2004, 2008), but the upstream signalling pathways modulating neuroinflammation are unclear. In the present study, we investigated the upstream mechanisms by which WNV modulates the expression of MMPs as well as key inflammatory cytokines. Our results not only highlight the fact that astrocytes respond to WNV infection



Fig. 5. COX-2 inhibitor limits the expression of cytokines in WNV-infected HBCAs. Supernatants collected from HBCAs infected in the presence or absence of NS-398 at day 3 p.i. were analysed by ELISA for levels of IL-1 β , -6 and -8. The data are expressed as mean concentration ± sD (pg ml⁻¹) of the amount of protein secreted into the supernatant from at least three independent experiments in duplicate. *, *P*<0.005 compared with uninfected control; **, *P*<0.05 compared with infected cells.

by significantly increasing COX-2 expression and production of its substrate, PGE2, but also link its increase to downstream pathogenic events such as induction of matrix-degrading enzymes and inflammatory cytokines.

The COX-2 enzyme is specifically implicated as being important in host responses to infection, and the ability of PGE2 to modulate inflammation and the immune response is well documented (Bazan, 2001; Candelario-Jalil & Fiebich, 2008). COX-2 enzymic processes not only generate PGE2 but also promote the production and release of reactive oxygen intermediates such as oxygen radicals and nitric oxide (Gebicke-Haerter, 2001) and are a major target of widely prescribed non-steroidal anti-inflammatory drugs and newer COX-2-specific inhibitors such as celecoxib. We have demonstrated for the first time that the COX-2 enzyme is induced at both the mRNA and protein level in human astrocytes (Fig. 1). This increase coincided with a peak in virus titres, and infection with UV-inactivated WNV failed to induce COX-2 in HBCAs, suggesting that induction of COX-2 is a result of active virus replication. The increase in COX-derived PGE2 as observed in Fig. 3 also correlated with COX-2 induction, and its dramatic decrease in the presence of the COX-2specific inhibitor NS-398 clearly indicated that COX-2, rather than COX-1, was the source of PGE2 in HBCAs. It is likely that COX-2-derived PGE2 acts as an autocrine or paracrine molecule, thereby mediating cellular events involved in WNV-induced neuroinflammatory responses. Although our in vitro data identified astrocytes as one of the sources of COX-2 in infected brain, it is possible that in vivo other cell types such as microglia and infiltrating monocytes may also contribute to the increased COX-2/ PGE2 levels. A recent study by Alvarez et al. (2008) demonstrated that COX-2 induced by HIV-2 in human astrocytes modulated their activation status, such as expression of glial fibrillary acidic protein, and that this effect was blocked by NS-398 treatment. Increased COX-2/ PGE2 has been linked to pathogenesis associated with other viruses such as HIV (Flora et al., 2006), hepatitis C virus (Lu et al., 2008), Epstein-Barr virus (Murono et al., 2001), Theiler's murine encephalomyelitis virus (Molina-Holgado et al., 2002), vesicular stomatitis virus (VSV) (Chen et al., 2002) and respiratory syncytial virus (Radi et al., 2010). Recent findings also report COX-2 induction and microglial activation following infection with JEV, another flavivirus (Ghoshal et al., 2007).

The integrity of the BBB in neuroinflammation is highly dependent on endothelial cells, production of cytokines and matrix-degrading enzymes, and recruitment of circulating inflammatory cells such as monocytes. An increase in the levels of MMPs, key matrix-degrading enzymes, has been linked to a compromised BBB and tight junction protein degradation in infections with neurotropic viruses such as HIV, herpes simplex virus and WNV (Keogh *et al.*, 2003; Sellner *et al.*, 2006; Wang *et al.*, 2008). Our previous results also demonstrated that increased MMPs produced by WNV-infected HBCAs are capable of digesting tight

Here, we further demonstrated that uPA is induced by WNV in HBCAs. Increased uPA observed in neurodegenerative diseases and infection with pathogens (Conant et al., 2004; East et al., 2005; Paul et al., 2005) is believed to activate plasmin, which can degrade most components of basal lamina either directly or through activation of procollagenases such as MMP-3 (Iwamoto et al., 2008; Zhao et al., 2008). These data suggest that several matrixdegrading components such as MMPs and uPA are induced in the astrocytes in response to WNV infection, which collectively might contribute to the BBB disruption in vivo. Upstream signalling mechanisms that govern expression of these matrix-degrading enzymes include COX-2 and cytokines such as IL-1 β and TNF- α (Aid *et al.*, 2010; Johnatty et al., 1997). Inhibition of COX-2 limits lipopolysaccharide-induced MMP-3 and -9 levels in both in vitro and in vivo mouse models (Aid et al., 2010; Candelario-Jalil et al., 2007). Similarly, in infections with pathogens such as Helicobacter pylori, the COX-2/PGE2 pathway is involved in the induction of the uPA/uPAR signalling pathway (Iwamoto et al., 2008). Our results demonstrating a significant reduction in the levels of MMPs and uPA in the presence of NS-398, as depicted in Fig. 4, suggest that WNV-induced COX-2 and its product, PGE2, are one of the key upstream signals that govern the activity of these matrixdegrading enzymes. However, these studies do not address the question of whether these MMPs are induced directly by PGE2 or by PGE2-modulated cytokines such as IL-1 β . Treatment with TNF- α results in the induction of MMPs and BBB disruption in mice, and the use of COX-2-specific inhibitors can eliminate TNF-*a*-induced MMP activity and reverse BBB disruption, thereby supporting the role of COX-2 in governing MMP activities (Candelario-Jalil et al., 2007). Similarly, post-ischaemic treatment with the COX-2 inhibitor nimesulide reduces BBB disruption and leukocyte infiltration following transient focal cerebral ischaemia in rats (Candelario-Jalil et al., 2005).

junction proteins of endothelial cells (Verma et al., 2010).

Pro-inflammatory cytokines such as TNF- α and IL-1 have been reported as potent inducers of neuronal injury in several neurodegenerative diseases such as cerebral ischaemia, spinal cord injury, multiple sclerosis and viral infections including HIV and JEV (Brabers & Nottet, 2006; Ghoshal et al., 2007; McColl et al., 2008). WNV infection also results in dramatic increases of several pro-inflammatory cytokines and chemokines such as IL-1 β and CXCL10 in both human brain cells and mouse brain (Garcia-Tapia et al., 2007; Kumar et al., 2010). Our data demonstrating an increase in IL-1 β in infected astrocytes is consistent with a previous observation reported by van Marle et al. (2007) where they not only documented increased production of IL-1 β by WNV-infected astrocytes but also established the role of astrocyte-derived neurotoxic mediators in inducing the death of naïve neurons (van Marle et al., 2007). Expression of IL-6 and -8, which are endogenous pyrogens exerting multiple downstream inflammatory signalling pathways, is elevated in many CNS

disorders including infection with JEV. The upstream mechanisms that control the production of different cytokines are complex and include COX-2-derived PGE2, intracellular potassium levels and reactive oxygen species (Aktas et al., 2007; Flora et al., 2006; Salminen et al., 2008). Cytokines such as TNF- α and IL-1 β can stimulate the expression of several secondary cytokines and COX-2. In contrast, it is also well established that COX-2-derived PGE2 can amplify the signalling cascade and production of cytokines, MMPs and cell-adhesion molecules (Kyrkanides et al., 2002; Mark et al., 2001; Tzeng et al., 2005). Similarly, PGE2 mediates the cellular effects of IL-1 β on parvocellular neurons in the hypothalamus, and the presence of NS-398 can abolish that effect (Ferri & Ferguson, 2005). Virus infections can induce multiple signalling pathways, which may cross-talk with each other or converge on common downstream effectors. In WNV infection, the initial stimulus for cytokine production in the brain is not clear. In the present study, our data demonstrating the ability of NS-398 to significantly block the production and release of IL-1 β , -6 and -8 (Fig. 5) indicate that COX-2/PGE2 is one of the pathways through which expression of key inflammatory cytokines is modulated in WNV-infected HBCAs. This finding is particularly important because it suggests that targeting COX-2 alone is likely to reduce multiple neuroinflammatory downstream molecules that contribute to complex pathological events such as chemoattraction of peripheral immune cells into the CNS, disruption of the BBB and neuronal injury.

NS-398 administration to modulate inflammation has been employed successfully in several neurodegenerative disease models and significantly attenuates levels of several inflammatory mediators (Bazan et al., 2002; Candelario-Jalil & Fiebich, 2008), thus establishing a complex feedback regulation. Furthermore, it is COX-2 activity, but not COX-1 activity, that contributes to the progression of focal ischaemic brain injury, and the beneficial effects observed with non-selective COX inhibitors are associated with COX-2 rather than COX-1 inhibition (Candelario-Jalil et al., 2003). Although the data are limited, a few studies have characterized the relative effects of COX-1 and COX-2 in modulating the host immune response (Aid et al., 2010; Candelario-Jalil et al., 2003). In influenza virus infection, important but opposite effects of the COX-1 and COX-2 enzymes have been documented, wherein COX-1 deficiency is detrimental to the host whilst COX-2 deficiency is beneficial (Carey et al., 2005). Similarly, selective inhibition of COX-2 also suppresses replication of VSV (Chen et al., 2000, 2002).

It has been suggested that it is the overwhelming inflammatory response that contributes to the severity of WNV disease, blocking of which not only improves disease parameters but also reduces virus titres in the brain (Wang *et al.*, 2004, 2008). Several of these cytokines act synergistically, resulting in cell death; therefore, it is important to identify upstream mechanism(s) by which the production of inflammatory molecules is modulated in WNV infection. Extrapolation of our results to animal

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models and testing the effect of COX-2 inhibitors on WNV titres, BBB disruption and CNS inflammation ultimately resulting in improved WNV disease outcome are warranted. In conclusion, our findings establish a critical role for COX-2 and PGE2 in modulating the inflammatory response in the CNS to WNV infection.

METHODS

Cells and virus. Primary HBCAs were obtained from the Applied Cell Biology Research Institute (Kirkland, WA, USA) and propagated as described previously (Verma *et al.*, 2010). All experiments were performed with cells between passage 6 and 10. Infections with WNV (lineage I, WNV strain NY99 originally isolated from crow brain and passaged once in Vero cells) and UV-inactivated WNV were conducted as described previously (Verma *et al.*, 2008, 2009) and cells were harvested at various time points p.i.

Treatment of HBCAs with COX-2 inhibitor. The toxicity of the COX-2-specific inhibitor NS-398 (Cayman Chemical) was tested by cultivating HBCAs in medium in the presence of NS-398 in a concentration range of 10–80 μ M using a CellTiter 96 AQ_{ueous} One Solution Cell Proliferation Assay kit (Promega) (Verma *et al.*, 2008). None of these concentrations of NS-398, or use of vehicle only (0.001 % DMSO), was found to be toxic to HBCAs. For all inhibitor experiments, HBCAs were treated with 25 or 50 μ M NS-398 at 12 h p.i. with a final concentration of DMSO of 0.001 %.

qRT-PCR. cDNA prepared from total RNA extracted from HBCAs was used for qRT-PCR as described previously (Verma *et al.*, 2010). The primer sequences and annealing temperatures used for the amplification of GAPDH and MMPs have been described previously (Verma *et al.*, 2009), and the primers for COX and inflammatory cytokines are described in Supplementary Table S1 (available in JGV Online). For all *in vitro* fold change analyses, the housekeeping gene GAPDH was used as an internal baseline reference to normalize each gene of interest, and the mRNA fold change of the gene of interest was determined at each time point compared with its corresponding infected control. All experiments were performed on samples from at least three independent infections in duplicate.

Western immunoblot analysis of COX-2. Total protein from HBCAs was extracted as described previously (Verma *et al.*, 2008, 2009, 2010). The nitrocellulose membranes were incubated with rabbit polyclonal anti-COX-2 antibody (1 mg ml⁻¹; Santa Cruz Biotechnologies) followed by HRP-conjugated secondary antibody and developed using an ECL detection kit (Pierce Technologies).

Immunocytochemical analysis. Fixed HBCAs were immunostained using anti-3.67G WNV envelope mAb (diluted 1:800; a generous gift from Dr D. Gubler, Duke-NUS Graduate Medical School, Singapore; originally produced by Dr R. H. Hall, University of Queensland, St Lucia, Australia) or anti-COX-2 (diluted 1:100) mAb followed by secondary antibodies conjugated to Alexa Fluor 488 or 546 (Invitrogen) and examined using a Zeiss confocal Pascal microscope equipped with a Zeiss Axiovert 200, as described previously (Verma *et al.*, 2009).

Analysis of PGE2, MMPs and cytokines by ELISA. At day 4 p.i., the supernatant medium from WNV- and mock-infected HBCAs in the presence or absence of NS-398 was collected to assay the levels of PGE2, MMPs and cytokines. All ELISAs were performed using undiluted supernatant medium in Biosafety Level 3 laboratories without any UV treatment. PGE2 was measured using a PGE2 Parameter kit (R&D Systems) according to the manufacturer's

instructions. ELISAs for total MMP-3 and -9 in the cell supernatant were conducted using Quantikine human MMP-3 and -9 ELISA kits, respectively (R&D systems), as described previously (Verma *et al.*, 2010). IL-6 and -8 levels were measured using Quantikine ELISA kits, whilst the IL-1 β level was measured using a Quantikine High Sensitivity kit according to the manufacturer's instructions. The ELISA plates were analysed using a Victor 3 microtitre reader equipped with Workout version 1.5 software (Perkin Elmer).

Statistical analysis. All *in vitro* data are reported as the mean \pm SD of at least three independent experiments in duplicate. An unpaired Student *t*-test using GraphPad Prism software version 5.0 was used to compare the levels of cytokines and MMPs assayed by ELISA. Differences where *P*<0.05 were considered statistically significant.

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