Loss of interneurons and disruption of perineuronal nets in the cerebral cortex following hypoxia-ischaemia in near-term fetal sheep

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Hypoxia-ischaemia (HI) in term infants is a common cause of brain injury and neurodevelopmental impairment. Development of gamma-aminobutyric acid (GABA)ergic circuitry in the cerebral cortex is a critical event in perinatal brain development. Perineuronal nets (PNNs) are specialised extracellular matrix structures that surround GABAergic interneurons, and are important for their function. Herein, we hypothesised that HI would reduce survival of cortical interneurons and disrupt PNNs in a near-term fetal sheep model of global cerebral ischaemia. Fetal sheep (0.85 gestation) received sham occlusion (n = 5) or 30 min of reversible cerebral ischaemia (HI group; n = 5), and were recovered for 7 days. Expression of interneurons (glutamate decarboxylase [GAD]1+; parvalbumin [PV]1+) and PNNs (Wisteria floribunda agglutinin, WFA) was assessed in the parasagittal cortex by immunohistochemistry. HI was associated with marked loss of both GAD1+ and PV1+ cortical interneurons (all layers of the parasagittal cortex and layer 6) and PNNs (layer 6). The expression and integrity of PNNs was also reduced on surviving GAD1+ interneurons. There was a trend towards a linear correlation of the proportion of GAD1+ neurons that were WFA1+ with seizure burden (r2 = 0.76, p = 0.0534). Overall, these data indicate that HI may cause deficits in the cortical GABAergic system involving loss of interneurons and disruption of PNNs, which may contribute to the range of adverse neurological outcomes following perinatal brain injury.

Perinatal hypoxic-ischemia (HI) is a common trigger of brain injury in term infants, being responsible for approximately 1–3 cases of moderate to severe encephalopathy per 1,000 live births, and is associated with a high risk of death or disability1. A major neuropathology observed following HI at term involves parasagittal watershed zone injury with neuronal loss, and damage to the underlying subcortical white matter2–9. This pattern of injury is strongly associated with adverse neurological outcomes, including cerebral palsy, cognitive delay, and epilepsy10–14.

During late prenatal and early postnatal development, the brain undergoes a period of marked growth and wiring, resulting in the establishment of highly ordered and complex functional networks15. A major aspect of this circuit development involves the integration of gamma-aminobutyric acid (GABA)ergic interneurons into the cerebral cortex16,17. GABAergic interneurons are the major population of cortical inhibitory neurons18, representing approximately 20% of all cortical neurons19. GABAergic circuits provide a critical source of inhibition required for the regulation of neuronal signalling, and thus maintain the balance between excitatory and inhibitory activity20,21. Importantly, a major period of GABAergic network development in the cerebral cortex occurs during the late prenatal period22,23. Thus, injury to the brain during this time has the potential to affect GABAergic circuitry and cortical function. Indeed, there is limited evidence for loss and dysfunction of interneurons in the cerebral cortex in human post-mortem brain tissue following perinatal brain injury24 and in experimental rodent models of neonatal HI25–27.
Perineuronal nets (PNNs) are specialised structures, formed by dense mesh-like aggregates of multiple extra-cellular matrix (ECM) molecules, which enwrap interneurons in the cerebral cortex and other brain regions\(^{28,29}\). PNNs are commonly detected using lectins such as Wisteria floribunda agglutinin (WFA) that bind to the N-acetylgalactosamine residues of chondroitin sulphate proteoglycans\(^{30,31}\). PNNs are important for normal GABAergic neuronal function\(^{32-34}\), including the regulation of cortical synaptic formation and stabilisation\(^{35-37}\), and have key roles in neuronal plasticity throughout development (reviewed in\(^{16}\)). Alterations in PNN formation and integrity are also associated with various human neurological diseases including schizophrenia\(^{39,40}\), epilepsy\(^{41,42}\), autism and Rett Syndrome\(^{43}\); PNN disruption was proposed to contribute to the associated deficits in neuronal signalling and cognition/behaviour in these disorders. Further, loss of cortical PNNs was reported in adult sheep and rodents following focal ischaemia\(^{44,45}\).

The effects of perinatal HI on cortical interneurons and PNNs remain unclear. Thus, in the present study we tested the hypothesis that cortical PNNs are associated with interneurons in the near-term fetal sheep, and that global cerebral ischaemia is associated with loss of both cortical interneurons and PNNs.

**Methods**

**Experimental protocols.** Fetuses were randomised to cerebral ischaemia (HI group; \(n = 5\)) or sham occlusion (control group; \(n = 5\)). At 128 ± 1 d gestation, ischaemia was induced by reversible inflation of the carotid occluder cuffs with sterile saline for 30 min. Successful occlusion was confirmed by the onset of an isoelectric EEG signal within 30 s of inflation. The carotid occluder cuffs were not inflated in sham control experiments. Fetal blood samples were drawn just before the occlusion, and at 2, 4, and 6 h after occlusion, followed by daily sampling for the remainder of the experiment. Animals were euthanized at 7 d recovery with an overdose of sodium pentobarbitone (9 g i.v. to ewe; Pentobarb 300; Chemstock International, Christchurch, New Zealand).

**Immunohistochemistry.** All antibodies/markers used in this study are summarised in Table 1.
Diaminobenzidine labelling of glutamate decarboxylase. At post-mortem, fetal sheep brains were perfusion fixed in situ in 10% phosphate-buffered formalin. Brains were then embedded in paraffin, cut into 10 μm thick sections using a microtome (Leica Jung RM2035; Leica Microsystems, Albany, New Zealand), and mounted onto glass slides. Regions of the forebrain used for analysis included sections taken at the level of the mid striatum, 26 mm anterior to stereotaxic zero according to the fetal sheep stereological atlas, with two adjacent sections selected from each animal (levels were matched between animals) for each analysis.

Sections were dewaxed for 2 × 15 min in 100% xylene, and rehydrated by immersion in decreasing concentrations of ethanol for 5 min each (100%, 90%, and 75%). Following 3 × 5 min washes in 0.1 M phosphate buffered saline (PBS), antigen retrieval was performed in 10 mM citrate buffer at >120 °C using the pressure cooker method (Antigen 200 Retriever; Electron Microscopy Sciences, Emigrad, Australia), and sections were again washed for 3 × 5 min in PBS. Sections were then incubated in 1% hydrogen peroxide in methanol to block endogenous peroxidase activity. Sections were washed for 3 × 5 min in PBS, then blocked in 5% normal goat serum (NGS)/PBS for 1 h at room temperature, followed by incubation with rabbit anti-glutamate decarboxylase (GAD) 65/67 (1:200; #ab49832; Abcam, Cambridge, MA, USA) overnight at 4 °C. After 3 × 5 min washes in PBS, sections were incubated in biotin-conjugated goat anti-rabbit IgG secondary antibody (1:200) in 3% NGS/ PBS for 3 h at room temperature, washed for 3 × 5 min in PBS, and then incubated in ExtrAvidin®-Peroxidase (1:200; #E2886; Sigma-Aldrich, St. Louis, MO, USA) in PBS for 2 h at room temperature. Sections were washed for 3 × 5 min in PBS, and then antibodies visualised by incubation in 3,3’-diaminobenzidine tetrahydrochloride hydrate (DAB; #D4293; Sigma-Aldrich). The reaction was stopped by washing in distilled water, and sections were dehydrated in increasing concentrations of ethanol (75%, 90%, 100%, 5 min each), followed by 2 × 10 min incubations in xylene, and then coverslipped with DPX mounting media (Sigma-Aldrich).

Fluorescent labelling. For multiple-labelling experiments, mid-striatal brain sections were selected, and sections were dewaxed, rehydrated, and antigen retrieved as described above. For detection of PNN expression (biotinylated WFA: 1:400; #L1516; Sigma-Aldrich), sections were incubated for 15 min in 0.1% avidin/PBS, and then 15 min in 0.1% biotin/PBS, for blocking endogenous biotin. For all other primary antibodies, including rabbit anti-GAD65/67 (1:200; #ab49832; Abcam), rabbit anti-parvalbumin (PV; 1:50; #ab11427; Abcam), mouse anti-NeuN (1:20; #MAB377; Merck Millipore, Billerica, MA, USA), and chicken anti-myelin basic protein (MBP; 1:500; #MBP; Aves Labs, Tilgard, OR, USA), sections were incubated in 0.1 M glycine for 20 min to reduce tissue autofluorescence, washed for 3 × 5 min in PBS, and blocked in 5% NGS. Sections were then incubated with WFA and primary antibodies in PBS/3% NGS for 3 nights at 4 °C, washed for 3 × 5 min in PBS, followed by appropriate secondary antibodies (Thermo Fisher Scientific, Waltham, MA, USA), including goat anti-rabbit Alexa Fluor 488 or 594 (1:200), goat anti-mouse Alexa Fluor 647 IgG, (1:100; #A-21240; used to detect NeuN primary antibody), streptavidin-conjugated Alexa Fluor 594 (1:200) or 680 (1:200), goat anti-chicken fluorescein conjugate (#F-1005; Aves Labs), and Hoechst 33342 (1:10,000; Thermo Fisher Scientific), for 2.5 h at room temperature. Slides were washed again for 3 × 5 min in PBS, and coverslipped using Vectashield mounting medium (Vector Laboratories, Burlingame, CA, USA).

For calmodulin-dependent protein kinase IIα (CaMKIIα) immunolabelling, paraffin-fixed sections were processed as above until the NGS blocking step, except the glycine step was omitted. Sections were then incubated in rabbit anti-CaMKIIα (1:200; #C6974; Sigma-Aldrich) for 3 nights at 4 °C, and then washed for 3 × 5 min in PBS. To amplify the CaMKII signal, biotin-conjugated goat anti-rabbit antibody (1:200; Vector Laboratories) was added for 3 h at room temperature, followed by washing and incubation in streptavidin-conjugated Alexa Fluor 488 (1:200) for 2.5 h at room temperature. Sections were washed and a second avidin–biotin block was performed, and sections were then incubated in WFA (1:400) overnight at 4 °C. Finally, sections were washed, streptavidin–conjugated Alexa Fluor 594 and Hoechst 33342 were added for 2.5 h, followed by washing and coverslipping as above.

For immunolabelling with cortical layer markers, brain sections were immersion-fixed in 4% paraformaldehyde, cut at 50 μm using a freezing microtome, and stored in cryoprotectant at −20 °C. Free-floating sections were washed for 3 × 5 min in PBS, incubated in 10 mM citrate buffer at 85 °C for 5 min for antigen retrieval, cooled for 20 min at room temperature, and then washed for 3 × 5 min in PBS. Sections were blocked for 1 h in 5% normal donkey serum (NDS; Thermo Fisher Scientific)/PBS at room temperature, and then double-labelled with rat anti-CTIP2 (1:500; #ab18465; Abcam) and biotinylated WFA (1:400) in 3% NDS/PBS for 3 nights at 4 °C.

### Table 1. Antibodies and markers used for immunolabelling studies.

| Antibody | Concentration | Specificity | Supplier details |
|----------|---------------|-------------|------------------|
| Calmodulin-dependent protein kinase IIα (CaMKIIα) | 1:200 | Excitatory neurons | #C6974; Sigma-Aldrich |
| CTIP2 | 1:500 | Layer 5 cortical neurons; weaker expression in layer 6 neurons | #ab18465; Abcam |
| Gamma-aminobutyric acid (GAD)65/67 | 1:200 | GABAergic interneurons | #ab49832; Abcam |
| Parvalbumin (PV) | 1:50 | PV interneuron subpopulation | #ab11427; Abcam |
| Myelin basic protein (MBP) | 1:500 | Myelin | #MBP; Aves Labs |
| NeuN | 1:20 | All post-mitotic neurons | #MAB377; Merck Millipore |
| Biotinylated Wisteria floribunda agglutinin (WFA) | 1:400 | Perineuronal nets | #L1516; Sigma-Aldrich |
were washed for 3 × 5 min, and incubated in donkey anti-rat Alexa Fluor 594 (1:500) and streptavidin-conjugated Alexa Fluor 680 (1:250) secondary antibodies, with Hoechst 33258 (1:10,000), in 3% NDS/PBS for 2.5 h at room temperature. After 3 × 5 min PBS washes, sections were mounted and coverslipped.

Imaging and analysis. Quantification of WFA, GAD, and PV expression in fluorescent sections. Using imaging software (Stereo Investigator; MBF Bioscience, Williston, VT, USA) driving a motorised stage (MAC 6000, MBF Bioscience) connected to a microscope (Zeiss AxioImager M2; Carl Zeiss Microscopy, LLC, Thornwood, NY, USA), the first and second parasagittal gyri (PG1 and PG2, respectively) of the right hemisphere (the left hemisphere was used in cases where there was damage to the right hemisphere) of each brain section were traced (2.5× objective), using the common sulcus as a guide for dividing the two gyri (Fig. 1). The dense WFA immunoreactive layer was then traced (5× objective) for each gyrus using WFA, NeuN, and Hoechst labelling as guides. The inner boundary of this layer was adjacent to the border of the white matter, defined using Hoechst and NeuN. The numbers of WFA+ , GAD+ , PV+ , and CaMKII/WFA+ neurons in this layer were counted (40× objective; counting frame size: 100 µm × 100 µm; ~25 sites per gyrus) for each gyrus, beginning at the intersection between PG1 and PG2. Positive cells were selected based on the following criteria: (1) a nucleus size within two standard deviations of the mean control nuclear size (based on data from ~20 control neurons per animal), and (2) a staining pattern and morphology similar to control neurons. Cell somata touching either of the two inclusion lines of the counting frame were included, while somata touching either of the two exclusion lines were not counted. Counting frames with greater than one-third falling outside of the traced boundary were excluded, as were regions with evidence of marked tissue loss. The densities of WFA+ , GAD+ , and PV+ cells (per mm²) were calculated for each gyrus, and two slides per animal were averaged to obtain final data.

Quantification of GAD+ cells in the parasagittal cortex in DAB labelled sections. The cortices of PG1 and PG2 were traced and counted as described above. The numbers of GAD+ neurons in all layers of the parasagittal cortex were counted using the fractionator probe (grid size: 500 µm × 500 µm; counting frame size: 150 µm × 150 µm; ~25 sites per gyrus) for each gyrus. The density of GAD+ cells (per mm²) was calculated for each gyrus, and two slides per animal were averaged to obtain final data.

Data analysis. Off-line physiological data analysis was performed using LabVIEW based customised programs. Seizures were identified visually and defined as sudden repetitive and evolving waveforms in the EEG signal lasting more than 10 s and of an amplitude greater than 20 µV. All biochemical, neurophysiological, cardiovascular, and fetal growth data from this cohort of animals were previously reported. For neuronal count data, a two-way ANOVA was performed to test for overall differences in cell densities in PG1 and PG2 of control and HI animals. Where an overall effect was found, a Fisher’s least squared derivative test was performed to assess for differences between the groups. Linear regression analysis was used to compare the relationship between loss of GAD+ neurons or PNNs in cortical layer 6, and seizure burden following HI. All statistical analyses were performed with statistical software (GraphPad Prism; GraphPad software, La Jolla, CA, USA). Statistical significance was accepted at p < 0.05. Data are presented as mean ± standard deviation. Histology and neuronal counts were performed by an assessor (T.M.F.) who was blinded to the treatment groups.

Figure 1. Sampling regions in the parasagittal cortex of the near-term fetal sheep brain. (A) Sheep brain atlas showing the first (PG1) and second (PG2) parasagittal gyri at the mid-striatal level. (B) Representative tracing and sampling sites of the Wisteria floribunda agglutinin+ (WFA+) layer for PG1 and PG2. The WFA+ layer is traced in pink (PG1) or blue (PG2), while the sampling sites are marked with circles (PG1) or diamonds (PG2). (A) was adapted with permission from http://www.brains.rad.msu.edu, supported by the US National Science Foundation and the National Institutes of Health. Scale bar: 5 mm.
Results

PNN and interneuron localisation in uninjured sheep brain. We first determined the pattern of WFA labelling in control near-term fetal sheep brains. Intense WFA staining was observed in the cerebral cortex (Fig. 2A), particularly in the parasagittal cortex (including both PG1 and PG2), and largely confined to the infragranular layers. Within these regions, WFA staining included a pattern of dense pericellular labelling around a subpopulation of somata (e.g., Fig. 3A,B [arrowheads]), as well as a more diffuse extracellular component (e.g., Fig. 3A,B [layer 6 region defined within dotted lines]). Co-labelling with the cortical layer marker CTIP2, which is most strongly expressed in cortical layer 5\(^56,57\), revealed predominant WFA labelling in cortical layer 6 (i.e., directly below the layer of highest CTIP2 expression). Immunohistochemistry for NeuN (post-mitotic neuron marker) and MBP (myelin marker) confirmed that the WFA layer was adjacent to the white matter (Fig. 2C,D). We also assessed GAD expression in the parasagittal cortex, and found that GAD\(^+\) neurons were distributed throughout all cortical layers, including the WFA-rich layer 6 (Fig. 2C,D).

Figure 2. Localisation of WFA staining and interneurons in the uninjured near-term sheep cortex. (A) WFA reactivity in the whole brain (red) co-labelled with Hoechst 33258 (blue). Representative high magnification examples of co-labelling of WFA (red) and CTIP2 (green) (B), WFA (white), glutamic acid decarboxylase (GAD, red), and NeuN (green) (C), and myelin basic protein (MBP, white), GAD (red), and NeuN (green) (D) taken from the boxed area in PG1 (A). CTX, cortex. Scale bar: (A), 5 mm; (B–E), 200 µm.
As expected, all WFA+ cells co-expressed the neuronal marker NeuN (Fig. 3A), and exhibited a pattern resembling immature PNNs, with dense immunoreactivity surrounding the soma and occasionally proximal processes (by contrast, mature PNNs surround proximal processes more frequently, and often extend further along processes)28,30,58. Double labelling revealed that 20.1 ± 8.8% (PG1) and 15.2 ± 4.5% (PG2) of total PNNs were expressed on GAD+ interneurons in layer 6 (e.g., Fig. 3B), while 6.6 ± 7.2% (PG1) and 4.3 ± 3.4 (PG2) of total PNNs were expressed on PV+ interneurons in layer 6 (e.g., Fig. 3C). Note that the levels of neuronal GAD and PV expression varied from weak to strong. Finally, 7.2 ± 6.2% (PG1) and 20.3 ± 15.5% (PG2) of total PNNs were expressed on CaMKIIα+ excitatory neurons in layer 6 (e.g., Fig. 3D).

**Figure 3.** PNN expression on various neuronal subtypes in the parasagittal cortex. Dense pericellular WFA reactivity (arrows) was observed around subsets of NeuN+ neurons (A; arrowheads), GAD+ interneurons (B; arrowheads), and CaMKIIα+ excitatory neurons (D; arrowheads), in a pattern resembling immature PNNs. By contrast, very few PNNs (arrows) were localised to parvalbumin (PV)+ interneurons (C). Note that the diffuse extracellular component of WFA staining is observed within layer 6 (dotted lines). Scale bar: 50µm.

Loss of total interneurons in the parasagittal cortex following term HI. To quantify the effect of HI on cortical interneuron survival, we assessed the density of GAD+ neurons in all layers of the parasagittal cortex in PG1 and PG2 (see tracing boundaries in Fig. 1B) at 7 d recovery after HI in near-term fetal sheep. There was a
significant reduction in the density of GAD\(^+\) neurons in both PG1 (Control: 151 \(\pm\) 38 vs. HI: 101 \(\pm\) 25 cells/mm\(^2\); \(p = 0.0391\)) and PG2 (Control: 134 \(\pm\) 45 vs. HI: 86 \(\pm\) 27, respectively; \(p = 0.0443\)) in HI animals (Fig. 4). There were no significant differences in GAD\(^+\) cell densities between PG1 and PG2 for the control or the HI groups.

**Loss of PNNs in layer 6 of the parasagittal cortex following term HI.** Next, we assessed PNN expression in layer 6 of the parasagittal cortex following HI. There was a marked overall decrease in WFA staining intensity in layer 6 of PG1 and PG2 after HI (Fig. 5B) compared with control animals (Fig. 5A), with complete loss in some regions (e.g., Fig. 5C,D). Areas with reduced but visible WFA reactivity were predominantly restricted to the superior surface of PG1 and PG2. Interestingly, the remaining PNNs in layer 6 of the injured PG1 and PG2 cortices were often less sharply defined (i.e., more diffuse), with reduced pericellular staining and lower WFA intensity compared with controls (Fig. 5E–H).

We then assessed the effect of HI on the number of PNNs in cortical layer 6. In HI animals there was a significant reduction in the density of total WFA\(^+\) neurons compared with controls in both PG1 (Control: 223 \(\pm\) 75 vs. HI: 16 \(\pm\) 17 cells/mm\(^2\), \(p < 0.0001\)) and PG2 (Control: 144 \(\pm\) 29 vs. HI: 42 \(\pm\) 30 cells/mm\(^2\), \(p = 0.0019\)) (Fig. 6A). Next, we examined whether this decrease in WFA\(^+\) cell density was because of interneuron death or loss of PNNs on surviving interneurons. HI was associated with a significant reduction in the density of GAD\(^-\) neurons (all GABAergic interneurons) and PV\(^+\) neurons within layer 6 in PG1 (GAD\(^-\): 369 \(\pm\) 119 vs. 74 \(\pm\) 39 cells/mm\(^2\), \(p < 0.0001\); PV\(^+\): 79 \(\pm\) 47 vs. 7 \(\pm\) 11 cells/mm\(^2\), \(p = 0.0014\)) and PG2 (GAD\(^-\): 343 \(\pm\) 117 vs. 61 \(\pm\) 28 cells/mm\(^2\), \(p < 0.0001\); PV\(^+\): 65 \(\pm\) 31 vs. 13 \(\pm\) 15 cells/mm\(^2\), \(p = 0.0139\)) compared with controls (Fig. 6B,E). Double-labelling for GAD/WFA\(^+\) and PV/WFA\(^+\) neurons in control animals showed that approximately 11\% (PG1) and 7\% (PG2) of the total GAD\(^+\) cell population expressed PNNs (Fig. 6C), and approximately 18\% (PG1) and 10\% (PG2) of the total PV population expressed PNNs (Fig. 6C). By contrast, HI was associated with reduced WFA colocalisation with GAD\(^+\) (to 1.7\% for PG1 and 1.0\% for PG2, \(p \leq 0.0001\) for both) and PV\(^+\) (to 0\% for both PG1 [\(p = 0.0217\)] and PG2, \(p = 0.2011\)]) cells, indicating that PNNs were also lost on surviving interneurons.

**Loss of cortical interneurons and PNNs is not significantly associated with increased electrographic seizure burden.** Finally, we examined the relationship between the density of GAD\(^+\) neurons or PNNs in cortical layer 6, and seizure burden in the HI animals. There was no significant linear correlation of GAD\(^+\) cell density (\(p = 0.6638\); Fig. 7A), WFA\(^+\) cell density (\(p = 0.2188\); Fig. 7B), or the proportion of GAD\(^+\) neurons that were WFA\(^+\) (\(p = 0.0534\); Fig. 7C) with seizure burden.

**Discussion**

Disrupted GABAergic signalling in the cerebral cortex is thought to contribute to the neurophysiological and cognitive impairments observed in numerous neurodevelopmental disorders, and there is limited preclinical and human evidence of damage to cortical GABAergic interneurons after perinatal brain injury\(^{16-27}\). Herein, we show that HI in the near-term fetal sheep (equivalent to the term human with respect to brain development)\(^{47,48}\) results in marked loss of GABAergic interneurons throughout the parasagittal cortex. We also provide new evidence for formation of PNNs on cortical interneurons during the last third of gestation in the fetal sheep, localised mainly to cortical layer 6. This population of cortical interneurons was particularly vulnerable to injury following HI, with widespread interneuron and PNN loss in layer 6, along with disruption of PNNs on surviving neurons. Overall, these data suggest that perinatal HI may cause deficits in the cortical GABAergic system that involve, at least in part, disruption of PNNs that are important for cortical inhibitory network function and regulation of CNS plasticity.

Injury to the parasagittal cortex is a common pattern of brain damage observed following HI in term infants. This pattern of injury is reproduced in near-term animal models of cerebral HI, including the 0.85 gestation fetal sheep\(^{24-27}\). Our findings demonstrate that HI near term causes marked loss of GABAergic interneurons (GAD\(^+\): by \(\approx 34\%\)) throughout the parasagittal cortex at 1 week of recovery. While these changes may relate to cortical layer 6. This population of cortical interneurons was particularly vulnerable to injury following HI, with widespread interneuron and PNN loss in layer 6, along with disruption of PNNs on surviving neurons.
to downregulation of interneuron markers on surviving neurons rather than cell death, this is unlikely as the degree of interneuron loss in the parasagittal cortex following HI in the present study was similar to the total cortical neuronal loss we previously reported for this cohort. Interneuron loss was even more pronounced in cortical layer 6 (GAD⁺: by ~88%; PV⁺: by ~86%), suggesting a particular vulnerability of this cell layer. Previous

Figure 5. Representative examples of PNNs in the parasagittal cortex of control and HI animals at 7 d recovery. Control (left column), HI (right column). WFA (A,B: grey, C–H: red), Hoechst 33258 (blue). Note that because of the marked reduction in WFA staining in HI animals, the image intensity of injured tissues (D,F,H) was manually adjusted (four-fold increase relative to control images) to allow visualisation of staining patterns. Representative examples of actual WFA staining levels between control and HI animals are shown in panels A and B. Images (C–H) were taken from PG1. Images (E–H) were taken from layer 6. Arrowheads indicate PNNs. The diffuse extracellular WFA staining is observed within layer 6 indicated by dotted lines. Scale bars: (A,B) 5 mm, (C, D) 500µm, (E, F) 100µm, (G, H) 20µm.
reports showed that unilateral HI caused acute death of GAD+ interneurons in the cerebral cortex in postnatal day (PND)9 mice 26, while asphyxia in newborn rats (PND0) resulted in long-term (PND56) loss of cortical calbindin+ interneurons (~20%) in the frontal cortex 27. Further, maternal hypoxia at embryonic day 17 in mice caused an acute, but transient, reduction of GAD protein expression in the cerebral cortex, and a reduced density of cortical calbindin+ interneurons in both superficial and deep layers by PND14 25. Despite no equivalent reports in term humans, post-mortem studies in preterm born infants with white matter lesions (25–32 weeks gestation) showed a reduced density of cortical calretinin+ interneurons and reduced numbers of cortical cells with GABA receptor expression24. Overall, these studies in combination with our observations indicate that injury to cortical interneurons may form an important component of the neuropathology observed following perinatal HI.

In the present study, HI resulted in loss of PNNs in layer 6 (by ~85%), which, at least in part, reflects the loss of cortical interneurons. Nevertheless, there was also a marked reduction in the expression of PNNs on surviving GAD+ (by ~85%) and PV+ (by 100%) interneurons, while remaining PNNs showed overall weaker and less defined staining, indicative of reduced PNN integrity. Similarly, expression of brevican, a major PNN component, was reported to be reduced in the cortex after HI in neonatal rats 26. Loss of cortical PNNs was also shown

Figure 6. Density of PNNs, and GAD+ and PV+ interneurons, in the parasagittal cortex of control and HI animals at 7 d recovery. Density of (A) WFA+, (B) GAD+, and (C) PV+ neurons in cortical layer 6. (D) Proportion of total GAD+ neurons that were WFA+. (E) Proportion of total PV+ neurons that were WFA+. n = 5 animals/group. *p < 0.05, **p < 0.01, ***p < 0.001. Data are mean ± standard deviation.

Figure 7. Relationship between loss of GAD+ neurons or PNNs in layer 6 of the parasagittal cortex with seizure activity in HI animals at 7 d recovery. Linear correlations of (A) GAD+ cell density, (B) WFA+ density, and (C) the proportion of GAD+ neurons with PNNs with seizure burden are shown.
after stroke in the adult rat45, which was attributed to degradation of PNN components. In support, expression of matrix metalloproteinase 9 (MMP9), which is particularly important for PNN degradation, was increased after HI in the neonatal mouse cortex46, and after focal ischaemia in the adult rat cortex47–49. ADAMTS (a disintegrin and metalloproteinase with thrombospondin motifs) and hyaluronidase (Hyal) enzymes also degrade PNNs, and are upregulated in the brain of adult humans and rats after focal ischaemia44–46. In these studies, upregulation of MMP, Hyal, and ADAMTSs occurred rapidly (2–24 h) after injury, and often persisted for days. Thus, the reduction in cortical PNNs with HI in the present study may reflect an increase in enzymatic PNN degradation. Further studies examining the exact timing of PNN production during fetal life and expression of PNN-degrading enzymes after HI will aid in determining the mechanisms underlying these changes.

Numerous studies have reported key roles of GABAergic neurons in normal cortical circuit development37,21, and of PNNs in GABAergic network function32–34. Deficits in cortical GABAergic signalling cause an imbalance in neural excitatory-inhibitory activity50, and can produce phenotypes similar to those observed in children following perinatal brain injury, including altered cortical plasticity and excitability, epilepsy, cognitive delay, and behavioural dysfunction68–74. Even though we did not observe a relationship between loss of interneurons or PNNs and seizure burden following HI in the present study, others showed that degradation of PNNs in hippocampal cultures in vitro caused increased interneuron excitability and seizure-like activity32,75. Further, loss of PNNs may increase interneuron vulnerability to oxidative stress and death76–78. Thus, we suggest that changes in cortical interneuron and PNN expression may still contribute to the cortical hyperexcitability, seizures, and delayed neuronal loss previously reported in this model76,79. Greater animal numbers are required to confirm the relationship between loss of interneurons or PNNs and seizure burden. Loss of GABAergic interneurons and PNN disruption around term age is also likely to have longer-term implications for cortical function. For example, GABAergic circuit establishment is the trigger for the beginning of critical periods of cortical development (periods during which synaptic circuitry is most easily modified by experience), while PNNs are required for restriction of plasticity and critical period closure79,80. Further studies are required to examine the longer-term effects of interneuron and PNN injury following perinatal HI.

PNNs are known to exhibit developmental and regional changes in their cortical expression patterns. Although there are no previous reports of fetal PNN expression, in humans, immature PNNs were first observed in the medial prefrontal cortex at 2 months of postnatal age, reaching mature levels by 8 years41. In rats, immature PNNs first develop in layer 6 of the parietal cortex at PND7, an age of brain maturation equivalent to the late gestation human82,83, followed by more widespread PNN expression in cortical layers 2–6 by PND14, and then adult-like patterns by PND3584. A similar timing of PNN formation was shown in the mouse visual cortex, but PNNs were highest in layers 4 and 5, and lower in layers 2/3 and 6 at all ages (PND10–PND70)85. In adult sheep, PNNs are predominantly located in layers 3 and 5, with less in layers 4 and 686, while in adult humans and monkeys, PNNs are expressed in cortical layers 3 and 4, with less in layers 2, 5, and 689,90. The initial appearance of PNNs in cortical layer 6 in the present study likely reflects the earlier formation of this layer compared with other cortical lamina, as during cortical development, the deeper layers (5 and 6) form first, followed by superficial layers (2–4)91. In sheep, neurogenesis and formation of cortical layers 5 and 6 occur from approximately 30 d gestation, and are largely complete by 60 d44. We found that only a low proportion of PNNs in layer 6 were localised to inhibitory GABAergic neurons identified using GAD (~18%) or PV (~6%), which contrasts with adult human and animal studies showing a majority of cortical PNNs (up to 87%) on interneurons89,90. These adult studies also imply that the maximum proportion of PNNs localised to excitatory neurons is ~13%, which is similar to that observed on excitatory neurons identified using CaMKIIα (~14%) in the present study. Thus, in the near-term fetal sheep brain, there remain numerous PNNs expressed on neurons that do not express common inhibitory markers. In support, initial appearance of PNNs and PV+ interneurons was reported to occur earlier than their colocalisation during postnatal development in the human prefrontal cortex42. These findings may relate to the ongoing maturation of cortical inhibitory circuitry during fetal and postnatal periods. Cortical interneuron fate, including the interneuron subtype, is specified during fetal life, prior to cell migration from the ganglionic eminences, while the differentiation of these neurons (including expression of various interneuron markers) then occurs once the cells reach their cortical destination51,92. In humans, the migration of GABAergic interneurons into the cerebral cortex continues into early postnatal life (~6 months of age)52. Further, in many species, including rodents, cats, and humans, the expression of GABA, GAD, and PV in the cerebral cortex does not reach adult levels until around adolescence (e.g., postnatal day 15–21 in the rodent, 5 weeks postnatally in the kitten, and 12–20 years in the human)53–55; human cortical GAD expression during late gestation is only <10% of adult levels, and then progressively increases postnatally and into adulthood53. Thus, the total proportion of PNNs on interneurons during early development may be higher than that observed by GAD and PV labelling; i.e., there may be neuronal populations with mRNA expression of GABAergic interneuron markers that do not yet express the protein products (CaMKIIα or Hyal enzymes) or degrade PNNs in the cortical GABAergic system involving loss of interneurons and disruption of PNNs. Given the importance of GABAergic networks in CNS maturation and function, these deficits may contribute to the wide range of adverse neurological outcomes associated with cerebral HI.

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**Author Contributions**

T.M.F., J.M.D., and R.G. conceptualised and designed the study. R.G., J.O.D., and G.W. performed sheep surgeries, experiments, and care. T.M.F. performed immunohistochemistry, cell counts, imaging, and analysis. R.G. undertook immunohistochemistry and seizure analysis. T.M.F. and J.M.D. wrote the main manuscript text. A.J.G. and L.B. provided overall oversight of the research. R.N.K. and J.D.P. provided critical review of the manuscript and laboratory support. All authors critically reviewed the manuscript and approved the final manuscript as submitted and agree to be accountable for all aspects of the work.

**Additional Information**

**Competing Interests:** The authors declare no competing interests.

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