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Role of microglia in a mouse model of paediatric traumatic brain injury

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ABSTRACT

The cognitive and behavioural deficits caused by traumatic brain injury (TBI) to the immature brain are more severe and persistent than TBI in the mature brain. Understanding this developmental sensitivity is critical as children under four years of age sustain TBI more frequently than any other age group. Microglia (MG), resident immune cells of the brain that mediate neuroinflammation, are activated following TBI in the immature brain. However, the type and temporal profile of this activation and the consequences of altering it are still largely unknown.

In a mouse model of closed head weight drop paediatric brain trauma, we characterized i) the temporal course of total cortical neuroinflammation and the phenotype of *ex vivo* isolated CD11B-positive microglia/macrophage (MG/MΦ) using a battery of 32 markers, and ii) neuropathological outcome 1 and 5 days post-injury. We also assessed the effects of targeting MG/MΦ activation directly, using minocycline a prototypical microglial activation antagonist, on these processes and outcome.

TBI induced a moderate increase in both pro- and anti-inflammatory cytokines/chemokines in the ipsilateral hemisphere. Isolated cortical MG/MΦ expressed increased levels of markers of endogenous reparatory/regenerative and immunomodulatory phenotypes compared with shams. Blocking MG/MΦ activation with minocycline at the time of injury and 1 and 2 days post-injury had only transient protective effects, reducing ventricular dilatation and cell death 1 day post-injury but having no effect on injury severity at 5 days.

This study demonstrates that, unlike in adults, the role of MG/M Φ in injury mechanisms following TBI in the immature brain may not be negative. An improved understanding of MG/M Φ function in paediatric TBI could support translational efforts to design therapeutic interventions.

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1. Introduction

Traumatic brain injury (TBI) is the most common injury leading to significant lifelong disability that occurs in children (Stanley et al., 2012). Unfortunately, the cognitive and behavioural deficits caused by traumatic brain injury (TBI) to the immature brain are more severe and persistent than those observed following comparable injuries to the mature (adult) brain (Anderson et al., 2005; Ewing-Cobbs et al., 2006; Hessen et al., 2007; Rivara et al., 2012) (reviewed in (Giza et al., 2007)) with injury in an experimental setting progressing into a chronic brain disorder (Ajao et al., 2012; Kamper et al., 2013). This is in contrast to Kennard's Principle that the immature brain has superior potential for repair (Bennet et al., 2013). This is of particular concern as children under the age of

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four years sustain TBI more frequently than any other age group (Koepsell et al., 2011) and in children under the age of 2 years, the rates of TBI serious enough to require intensive care support are as high as 50 per 100,000 (Keenan et al., 2003). A developmental sensitivity to TBI as seen in humans is also observed in a rodent model of TBI, where within the first 30 days of life, injury is maximal when TBI is caused at postnatal day 7 (P7) (Bittigau et al., 1999). In addition, during the first three postnatal weeks, rodents display a heightened sensitivity to excitotoxicity (Ikonomidou et al., 1999). In mouse and humans this period is when developmental processes such as maximal brain growth, synaptogenesis and myelination occur.

In the paediatric population, TBI is caused by injuries and insults, which include acceleration/deceleration injuries (shaken baby syndrome) and contusion injuries (direct skull impact) (Pinto et al., 2012). Contusion injuries are the prevailing form of non-inflicted injuries and also represent a large proportion of inflicted injuries (Pinto et al., 2012). The primary injury process in TBI is mechanical damage (i.e. shear forces inducing vascular damage and bleeding), followed immediately by mast cell degranulation (Stokely and Orr, 2008), and secondary pathological processes. including excitotoxicity, ischemia, mitochondrial dysfunction, activation of matrix metalloproteinases (MMPs) and activation of caspases leading to apoptosis (Xiong et al., 2013). These secondary injury processes induce neuroinflammation, which itself has the potential to be neurotoxic (Hagberg et al., 2012), but which is poorly understood in the immature brain following TBI.

Microglia (MG) are the central regulators of neuroinflammation, involved in the pathological processes of the majority of acute and chronic brain injuries, such as stroke, Alzheimer's disease and multiple sclerosis (for review see (Prinz et al., 2011)). Thus MG are logical candidates to mediate neuropathological changes following TBI in the immature brain. MG possess enormous functional plasticity that allows them to participate in both injury and repair, as reviewed in (Colton and Wilcock, 2010; Ransohoff and Perry, 2009). The nomenclature of these functional activation states (phenotypes) of MG has been simplified to facilitate their description and a common nomenclature includes classic pro-inflammatory or cytotoxic, anti-inflammatory or reparatory/regenerative and immunomodulatory phenotypes.

There are specific differences in the immune and inflammatory responses to injury between neonatal and adult humans and experimental animals (Copland et al., 2004; Giza et al., 2007; Schultz et al., 2004; Zhu et al., 2005), including in microglia responsiveness (Butovsky et al., 2014). Studies of neuroinflammatory profile and MG activation states have recently been published in adult models of TBI (Bye et al., 2007; Kumar et al., 2015) but it is unknown how microglia would respond to a similar injury to the developing brain. As such, this study investigates for the first time the characteristics of MG- driven neuroinflammation in a mouse model of paediatric TBI. Furthermore, as a proof-of-concept, we aimed to assess the effects of modulating MG activity on injury severity using the immunomodulatory tetracycline minocycline. Minocycline reportedly has strongly anti-inflammatory actions and has been used to reduce MG activation and injury with success in numerous pathological models (see Table 1 and review, (Garrido-Mesa et al., 2013)).

2. Materials and methods

2.1. Animals

Study ethics were approved by the Bichat and Robert Debré Hospital ethics committee (No 2011-14/676-0050) and adhered to the European Union Guidelines for the Care and Use of Animals. Procedures were typically carried out between 10am and 1 pm (light phase: 7am-7 pm daily), all animals were monitored daily during experimentation. A single animal represents an experimental unit with groups spread between and across litters where possible and each litter had an approximate 50–50% spread of males-females. Specifically, data in Fig. 2 are derived from 6 litters; Fig. 3 derived from 24 litters; Figs. 4, 5 and 7 derived from 6 litters each; Fig. 6 derived from 6 litters. Animals were housed in Plexiglas cages (30x18x15 cm) together with littermates and their dam for the whole of the experiment. Animals had access to standard chow and water *ad libitum* and bedding was wood-chips with shredded paper for nesting (Pharmaserv, France).

2.2. Traumatic brain injury model and experimental procedure

Postnatal day 7 (P7: weight 4–5 g) OF1 mice (Charles River. L'Arbresle, France) of both sexes were randomly (alternating animals) allocated to TBI, control or TBI+ treatment (phosphate buffered saline [PBS] or minocycline) groups. The study protocol is detailed in Fig. 1. A dose of 45 mg/kg of minocycline was chosen based on its prior use in models of adult TBI, stroke and paediatric excitotoxic lesion, see Table 1. In a separate experimental workspace within the animal facility, mice were anesthetized with isoflurane (8% induction) and subjected to a closed head weightdrop head trauma at P7 in a model as described previously (Kaindl et al., 2007). In brief in a process lasting no more than 3 min, the skull was fixed into a stereotaxic frame, the skull surface exposed with a skin incision and the impact device was oriented parallel to the parietal bone with the centre of the foot plate (2 mm diameter) positioned 2 mm anterior and 1 mm lateral to lambda on the parietal bone. The foot-plate was first allowed to touch the skull and was then further depressed by 0.5 mm. The impact device consisted of a hollow stainless-steel cylinder 20 cm in length, perforated at 1 cm intervals to prevent air compression, and guiding a 10 g weight falling from a height of 10 cm onto the foot-plate (2.0 mm in diameter). The contusion impact was delivered unilaterally to the left side of the skull, the same operator conducted all experiments and cortical contusions were of comparable severity in all animals. Body temperature was kept constant via the use of a heating pad maintained at 37 °C until pups were returned to their dams at approximately 15 min post-TBI. Sham animals were anesthetized and an incision made in their scalp, this was then sutured and animals were recovered after 3 min in line with the time taken for the TBI procedure. Minocycline (45 mg/kg in PBS: Sigma, Lyon, France) (Cai et al., 2006; Dommergues et al., 2003) or PBS alone was injected intraperitoneal immediately following TBI, and at 24 and 48 h post-TBI, depending on the protocol. A group of sham minocycline was not included in this study as the specific aim was to investigate the effects of modulating the microglial activation state associated with TBI. Furthermore, minocycline has been widely reported to have no effect on microglial gene expression in a basal state (Kobayashi et al., 2013; Scholz et al., 2015).

2.3. Tissue preparation, and histology

One or five days after TBI, animals were euthanatized via an overdose of pentobarbital and decapitation and brains were immersion fixed (formol 4% for 5 days), embedded in paraffin and coronally sectioned (16 μ m) from the frontal pole to the occipital lobes. Ventricular area was determined as described previously (Kaindl et al., 2007; Moretti et al., 2016) on cresyl-violet-stained sections. In short, the border of each lateral ventricle from three serial sections spanning the hippocampus and midstriatum was outlined, then the cross-sectional ventricular areas were

Table 1

Summary of selected studies investigating the neurotherapeutic effects of minocycline.

Study	Animal	Injury	Dose	Regime	Cell death / Lesion Volume	MG number	Outcome	
Dommergues et al. (2003)	P5 mouse	Excitotoxic	45 mg/kg	Twice daily from P5-P7	Decreased Cleaved Caspase-3 at +1 day and decreased lesion volume at +5 days	Decreased numbers of Lectin+ MG	Decreased lesion volume at +5 days	Solution
Fox et al. (2005)	P7 rat	MCAO	45 mg/kg	+2 h & +2 h, or +8 h & +18 h	Decreased lesion volume at +1 day	No change in ED1+ MG numbers	No improvement in lesion volume at +7 days	Ţ
Yang et al. (2015)	Adult rat	MCAO	5 mg/kg	+5 min	Decreased infarct on MRI	60% decreased (Increased anti- inflammatory type MG)	Improvements on MRI at 4 week	A
Cai et al. (2006)	P4 rat	HI	45 mg/kg	12 h before, immediately after & daily for 3 days	Decreased pyknosis at +4 days	50% decrease in numbers of lectin positive MG	Decreased loss of mature oligodendrocytes and myelin at +2 weeks	A
Lechpammer et al. (2008)	P6 rat	НІ	50 mg/kg	Immediately following HI	Decreased white matter injury at +3 days	Decreased numbers of CD68+ & MHCII+ cells at +3 days		A
Arvin et al. (2002)	P7 rat	ні	22.5–45 mg/kg	Immediately before or +3 h	Decreased lesion volume at +7 days	-	Decreased lesion volume at +7 days	E)
Tsuji et al. (2004)	P7 mouse	HI	22–135 mg/kg	 (1) Twice in first 24 h (45 mg/kg) & twice in the next 24 h (23 mg/kg), or (2) Twice in first 24 h (135 mg/kg) & twice in the next 24 h (68 mg/kg), or (3) Single dose 12 h before HI (45 mg/kg) 	Exacerbated total injury score for all treatments (1–3) at +7 days	-	<i>Mouse:</i> Exacerbated total injury score for all treatments <i>(1–3)</i> at +7 days	Ţ
	P7 rat		45 mg/kg	(4) Immediately before HI, or(5)12 h before HI	Decreased total injury score for both treatments (4–5) at +7 days	-	<i>Rut:</i> Decreased total injury score for both treatments $(4-5)$ at +7 days	E)
Hanlon et al. (2016)	P11 rat	Repeated TBI (CCI)	45 mg/kg	Once immediately after the third and final TBI	No change in fluro-jade B+ cell number at +3, +7 & +21 days	No change	Exacerbated defects in retention tasks. No improvements in tissue loss or spatial memory defects at +21 days.	P
Bye et al. (2007)	Adult mouse	TBI (CCI)	45 mg/kg	+30 min & every 12 h for 3 days	Decreased at +1 day, no change at +4 days	Decreased amoeboid ED1+ MG	No improvement in motor function at +1 week	P
Homsi et al. (2010)	Adult mouse	TBI (CCI)	90-45 mg/kg	+5 min (90 mg/kg), +3 h & +9 h (45 mg/kg)	50% decrease in cortical tissue loss	50% Decrease in CD11b+ MG/M Φ	Improvement in locomotor hyperactivity at +8 weeks	
Current study	P7 mouse	TBI (WD)	45 mg/kg	Immediately after & at +24 h & +48 h	Decreased cleaved caspase-3+ cell numbers, decreased ventricular volume at +1 day	15% decrease in numbers of Iba1+ MG (minimal change in activation by gene expression)	No improvements in neuropathology at +5 days	P

HI, hypoxic/ischemic. CCI, controlled cortical impact. WD, weight drop.



Fig. 1. Schematic representation of the experimental procedures including administration of drugs and tissue collection, and injury distribution. Injury is indicated by the expression of CCasp3 at 1 day (+1d) and 5 days (+5d) following TBL Slightly increased areas of labelling shown by orange stars, moderate increases by red stars and intense changes shown as blocks of red. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

determined using ImageJ software (version 1.43; National Institute of Health, Bethesda, Md., USA) and the ratio between the left (ipsilateral) and right (contralateral) ventricular areas determined. All tissue processing and analyses were carried out by investigators blind to the treatment group due to coding of the brains and covering of the codes during analysis. There were no differences in the ventricular area of the contralateral hemisphere between sham and TBI mice. Immunohistochemistry (IHC) was performed as previously described (Fleiss et al., 2012), and the antibodies used included: rabbit monoclonal anti-ionized calcium binding adaptor molecule-1 (Iba-1; 1:1000, Wako Chemicals USA, 019-19741), Rabbit monoclonal anti-cleaved caspase 3 (CCasp3; 1:200, Cell Signalling, 9661), mouse monoclonal anti-myelin basic protein (MBP; 1:500, Millipore, MAB382) and mouse monoclonal antimicrotubule-associated protein 2 (MAP2; 1:2000, Sigma, M4403). After overnight incubation with primary antibodies and washing, sections were incubated with appropriate secondary antibodies (1:200; Vectorlabs, California, USA).

2.4. Analysis of neuropathology

Ventricular volume was assessed by measuring the area of the ipsilateral and contralateral ventricles and expressed as percentage of change compared to the contralateral values. Iba-1-positive and CCasp3-positive cells were counted in the parietal cortex, hippocampal CA1 region and striatum of the traumatized hemisphere, on two images captured using a Leica DM6000 B microscope (Leica Microsystems Ltd.) and a 10X objective at the level of maximum lesion (approximately -1.50 mm from bregma). Counts were carried out using Image J and cell numbers within a given region expressed as cells/mm². The area of MAP2 and MBP immunolabeling was measured at 4–6 levels per brain (one 16 µm-thick serial section every 576 µm) as previously described (Fleiss et al., 2012). Volumes of MAP2 and MBP immunolabeling were calculated from area measurements according to Cavalieri's principle using the following formula: $V = SA \times P \times T$, where V is total volume, SA is the sum of the areas measured, P is the inverse of the sampling fraction and T is the section thickness. Volume loss was estimated by the difference in calculated volumes between the contralateral (right) and the ipsilateral hemispheres (left).

2.5. CD11B-antibody-coupled magnetic cell isolation

At different time-points following TBI (2, 6, 14, 24 h and 5 days), cells positive for CD11B (cluster of differentiation 11 beta, a marker for M Φ and MG), were extracted using the antibody-coupled magnetic bead system (MACS) following the manufacturer's recommendations (Miltenyi Biotec, Bergisch Gladbach, Germany) and as previously reported (Schang et al., 2014). In brief, the olfactory bulbs and cerebella were removed and the hemispheres mechanically and enzymatically digested using the Neural Tissue Dissociation Kit (Miltenyi Biotec, Germany). Three or four hemispheres were pooled for each sample to ensure sufficient RNA quantities. In a preliminary analysis, comparisons of MG/M Φ activation and cytokine gene expression between left and right sham hemisphere did not show any differences and samples were pooled. Homogenized and digested tissue was incubated with magnetic coupled anti-CD11B antibodies and CD11B-positive cells were separated in a magnetic field before being counted and frozen at -80 °C. The purity of separated cells was assessed using quantitative real-time polymerase chain reaction (gRT-PCR) for glial fibrillary acidic protein (GFAP; astrocytes), MBP (oligodendrocytes), neuronal nuclear antigen (NeuN; neurons) and CD11B (MG/M Φ), and showed levels of contamination less than 5%. We have described the CD11B-positive population extracted from the brain as MG/M Φ as we cannot exclude a contribution of macrophages to the cell population (Hsieh et al., 2013).

2.6. RNA extraction and quantification of gene expression by real-time qPCR

MG/M Φ qRT-PCR, primer design, and PCR setups were similar to that previously described (Chhor et al., 2013; Husson et al., 2005; Schang et al., 2014). In brief, RNA was extracted using Qiagen RNA extraction columns as per the manufacturers instructions, including initial homogenisation in Trizol (Invitrogen). RNA purity was verified using a nanodrop. Reverse transcription was performed using an iScript RT kit (Biorad) as per manufacturers instructions. PCR reactions were setup on a loading robot in 384 well plates with Sybr green from Biorad as per recommended protocol. Primer sequences are given in Table 2. Gapdh (glyceraldehyde-3-phosphate dehydrogenase) was used to normalize the quantitative experiments based on prior referencegene suitability testing and we verified for each experiment that the raw Gapdh values were not significantly different between groups. The relative quantities are expressed as the specific ratio between the gene of interest and the reference gene. Genes were classified as cytotoxic, repair/regeneration or immunomodulatory based on the literature (Colton and Wilcock, 2010; Ransohoff and Perry, 2009) and previous characterization in our lab (Chhor et al., 2013).

2.7. Protein extraction procedure and multiplex cytokine/chemokine assay

Frozen cortices from 6, 14 and 24 h post-TBI were homogenized in 0.1 M PBS, and extracts sonicated in ice-cold homogenization buffer (3 mM ethylenediaminetetraacetic acid [EDTA] and 1% protease inhibitor cocktail, [P8340, Sigma] in 0.1 M PBS) and centrifuged (800xg for 10 min). The supernatant was collected and stored at -80 °C. Protein concentrations were determined via a bicinchoninic acid (BCA) assay. After thawing on ice, supernatants were centrifuged briefly to remove particulates (300g for 10 min). Cytokine and chemokine levels were measured using a 96-well magnetic plate assay on a Bio-Plex 200 according to the manufacturer's instructions (BioRad laboratories, Marnes la Coquette, France). Cytokines and chemokines measured included interleukins

Table 2			
Primer sequences	and	NCBI	references.

GapdhGGC CTT CCG TGT TCC TACTGT CAT CAT ATC TGG CAG GTTNM_008084.2iNosCCC TTC AAT GGT TGG TAC ATG GACA TTG ATC TCC GTG ACA GCCNM_010187.2CD32CTG GAA GAA GCT GCC CAA AACCCA ATTG CCA TGC GAG ACT AANM_010187.2CD86GAG CGG GAT ACT AAC GCT GAGGC TTT CCA TGC CTT CAC TCNM_01188.3Ptgs2TCA TTC ACC AGA CAG ATT GCTAAG GCT TTG CGT TCG GTT GANM_008625.2Arg1GTG AAG AAC CCA CGG TCT GTTGC AAG AGC GCT TCG GTT CAA CTGNM_008625.2Jgg13GAT GAC AAT CAT GGG GCA CAGATT GAA GAC GC GGT CAA CTGNM_0016705.3Jgg14TGG ATG CTC TT GAT TG GGGCA ACA CTC ATC CAC AAT GCNM_010705.3Jgg15TGC AAG CCC GT GT GT GGGCA ACA CTC ATC CAA CTG CT NM_010705.3Jgg14TGC AGA GCC CCG AGG CCGCA CAC GCG GT GAA GTNM_00172475.1JltmTGC AGC CAA GCC GT GT GGGCA CAC GT GT GCA GT GAT GAMM_001707.3JltmTGC AGC GA GCC CGA AGC CCACT CTG GAG AGA CTT GGT GGNM_0018700.3Socs3GGT TCAA AGC AGC CCT TCA GAGCC CCT TCT GG GCC GAG AAG ATNM_0011584.2Ll-16GAA CCA CA CA GCA GT CT TCA GAGCC CTT GG GCC CTT GG GCCNM_0011584.2Ll-12bATC CAG CCC AAG GAA GAA TCTCT TCT GGT GT GC CTT GG GCCNM_0011593.3Ll-12bATC CAG CCC AAG GAA GAAATA ACG CAT CT GG GCC TT MM_001593.3Ll-12bATC CAG CCC AAG CAA CAAATA CCG AGC CT TCC GGT CCNM_001393.3CxC11GCA CCC AAG CAA GCA GCTT CAG CT GG GCC TT CC CT CM MM_001393.3<	Gene	Sense	Antisense	NCBI Reference
iNosCCC TTC AAT GGT TGG TAC ATC GACA TTG ATC TCC GTG ACA GCCNM_010927.3CD32CTG GAA GAT GCC AAA ACCCA ATG CCA AGG CACT AANM_010187.2CD86GAG CCG GAT AGT AAC GCT GAGGC TCT CAC TCC CTT CAC TCNM_011988.3Prgs2TCA TTC ACC ACA CAG ATT GCTAAG GCT TTG CGG TAC TA TTNM_008625.2Arg1GTG CAG AAC CCA CGG TTC GTGCC CAG AGA TGCT TCC ACA TGNM_008625.2Igals3GAT CAC AAT CAT GGG CAC AGATT GA AGG GGC GTT GANM_0010705.3Igf1TGG ATG CTC TTC AGG CAC AGATT GA CGG GGG GTT AAA GTNM_0010712.4Sphk1TCC AGA AAC CCC TGT GTA GCCAC ACA CT CAC AAT GCNM_001072.4IlmTTG TGC CAA GT CGG GAG CACT CTG GAG ACA GT TGG CAG AGA GT TGC GA GAG GTNM_00100870.3Socs3CGT TGA CAG AC CCG AGA GCACT CTG GGA AGA GT TGGNM_00100870.3IL1bGGG CCT CAA AGG CAA GAC CCGCC TT CT TTG GGT ATT GG TCGNM_0011548.2IL10CTC CC GT GGA AAT AAG AGCGCC TT CT TG TG GA CT TGC GT CCNM_0011548.2IL-12aTCA CAA CCA AGA GA CCTCT CTA GAG CCT CT CT GG CT CM M_001548.2IL-12aIL-12bATC CAG CCAA AG CAG CACTCT CTG CG CG CG GT CA CTCNM_0011593.3Cxcl10GGG TAA AGC GAG GAG CAG CCTAT CAG CTC GG CG CT CA GAG CTNM_001593.3Cxcl11GCA CCA ACC GA GA CAG CCTAT CAG CCA CAG CT CCNM_001393.3Cxcl10GGG TAA AGC GAG CG CGA GAG GG GCT TAT TGA CGC GGG GT CA NM_001333.3Cxcl10GGG CAA AGC AGA CCTTG GG CT CA AGC GGG GT CA AGG GGT CGA AGA CGCMM_001337.2 <t< td=""><td>Gapdh</td><td>GGC CTT CCG TGT TCC TAC</td><td>TGT CAT CAT ATC TGG CAG GTT</td><td>NM_008084.2</td></t<>	Gapdh	GGC CTT CCG TGT TCC TAC	TGT CAT CAT ATC TGG CAG GTT	NM_008084.2
CD32CTG GAA GAA GCT GCC AAA ACCCA ATG CCA AGG CAG ACT AANM_010187.2CD86GAG CGG GAT AAC ACC GCT GAGGC TCT CAC TGC CTT CAC TTCNM_019388.3Ptgs2TCA TTC ACC AGA CGA TT GCTAAG CGT TTG CG TAC TCA TTNM_011198.3CD206CTT CGG CCC TTT GGA ATA ATTAG AAG AGC CCT TGG GTT GANM_008625.2Arg1GTG AAG CAC CGG TCT GG CACGCC AGG GAT GCT TCA AC TGNM_001705.3Igd1TGG ATG CTC TTC AGT TGG TG CGCA AGA CTC ATC CAC AGA GTNM_0101512.4Sphk1TCG AGA CCC TCT GTG AGCCA CGA CTC ATC CAC AAT GCNM_0011272.5.1IllmTTG TGC CAA GTC TGG AGA TGTTC CAC GGC GAT GA AG GTNM_0011087.0.3Socs3CGT TGG ACA CCC CTT CTA GCACT CTG GA GAG GT GT GT GGNM_0010870.3IL1bGGG CTC CAA AGG AAA GAA TCTCT TCT GT GG GCC GAG AGA GAT NM_010187.2NM_001165.5IL1bGGG CCT CAA AGG AAA GAA TCTCT TCT GT GG GC GGC GAG AGA GAT NM_0010870.3IL1bGGG CCT CAA AGG AAA GAA TCTCT TCT GT GT GG CC CAG MG ATNM_001159.4IL10CTC CCC TGT GAA AGA GACGCC ATT CT TCT GT GG TCCNM_001159.42.1IL12aTCA CAA CCA AGG CAT CATACT AGG GCC TCA TT CA GC TCCNM_001159.42.1IL-12aTCA CAA CCA TCA GCA GAT CATCA CAG GCT CT TG GG CC CAA GC CNM_001380.1IL-12bATC CAG CGC CAA GAA AAAAT AGG GTC CGC CAA GAC CACNM_00159.3Cxcl10CGG CTA AA GCA AG CAA CCACT CGT CGG CCA AGA CCA CT CG CT CANM_00139.3Cxcl10CGG CTA AA GCA AG CA CT TTAGG GTC CT CA GG CT CA CT CG CT CA <td>iNos</td> <td>CCC TTC AAT GGT TGG TAC ATG G</td> <td>ACA TTG ATC TCC GTG ACA GCC</td> <td>NM_010927.3</td>	iNos	CCC TTC AAT GGT TGG TAC ATG G	ACA TTG ATC TCC GTG ACA GCC	NM_010927.3
CD86GAG CGG GAT AGT AAC GCT GAGGC TCT CAC TCCNML_019388.3Prgs2TCA TIC ACC AGA CAG ATT GCTAAG CGT TIG CGG TAC TCA TTNML_011198.3CD206CTT CGG CCC TTT GGA ATA ATTAG AAG AGC CC TT CG GT GANML_008625.2Arg1GTG AAG AAC CCA CGG TCT GTGCC AGA GAT GCT TCC AAC TGNML_007482.3Lgdl33GAT CAC AAT CAT GGG CAC AGATT GAA GCG GGG GTT AAA GTNML_010705.3Igf1TGG AG CTC TCT CA GT TCG TGGCA ACA CTC ATC CAC AAT GCNML_00172475.1Il/mTTG TG CC AAA GC CCC TGT GTA GCCAG CGG GTG GAA GG GTNML_001172475.1Il/mTTG TG CC AAG TCT CC GAC ATCTCT CTA GAG CGG AGA GAG TTNML_00100870.3Socs3CGT TGA CAG TCT TCC GAC AATTAT TCT GGG GGC GAA AGA ATNML_001707.3IL/bGGG CT CAA AGG AGT CCT TCG GAAGCC ATT CT TCT GGT ATT GCT TCG GT CCNML_001188.1IL/10CTC CCC TGT GAA AAT AAG AGCGCC ATT CT TCT GGA CT CT CG TG NML_001188.1NML_00131168.1IL/12aTCA CAA CCA GAA GAA CCATGC ACA GCT CT CG GC CT CAT MML_00139424.1NML_00139424.1IL-12aTCA CAA CCA TCA GCA GAT CATGC ACA CCT TGG CT CCNML_00139424.1IL-12aTCA CTA CAA CCA AAA AAAAAT ACC GGT CTG GG GTT CNML_008352.2IL-18TTC GTT GAC AAA AGA CAG CCTAT CAG GTC TGG GC CT AG AAC CTNML_001393.3Cxcl10GGG TAA AGG AGG CT TATAGG GTC TGG GGC TA GAA CTNML_001393.3Cxcl10GGG TAA AGG AGG CTG TTTCT CG CT CAA AGG CAAG CTNML_011333.2Cd2CAT CCA CCT TGT GG CTA TAAGG GTC T	CD32	CTG GAA GAA GCT GCC AAA AC	CCA ATG CCA AGG GAG ACT AA	NM_010187.2
Ptgs2TCA TTC ACC AGA CAG ATT GCTAAG CGT TTG CGG TAC TCA TTNM_011198.3CD206CTT CGG GCC TTT GGA ATA ATTAG AAG AGC CCT TGG GTT GANM_008625.2Lgals3GAT CAC CAC GG TCT GT GTGCC AGA GAT GCT TCC AAC TGNM_010705.3lgf1TGG ATG CTC TTC AGT TCG TGGCA AGA CTC ATC CAC AAT GCNM_010172475.1Il1mTTG TGC CAG ATC TG GG AGA GCCAG CGG GTT GAA GAG CTT GGT GGA GTG CTTG GA GA GA CCC GAG GT GAAG GTNM_001172475.1Il1mTTG TGC CAG ATC TGG AGA CCACT CTG GAG AGA GTT GGA GGG GTT GAA GANM_00108700.3Socs3CGT TGA CAG CCC GAG CACT CTG GGG GGG GAG AGA GTNM_007707.3IL-6CAA AGC CAG AGT CT TCA GACCC TTG GGA GAC CT TGG CNM_001548.2IL-6CAA AGC CAG AGT CT TCA GAGCC AGT CCT TGG GT CCNM_001548.2IL-12aTCA CAC ACC TCA GAG AGA CCCCT TG TAG CAC CTC TGC MN_001548.2IL-12IL-12bATC CAG CGC AAA AGA AAAAT AGC GAT CCT GGG GTC CT MN_001548.2IL-18IL-18TTC GTT GAC AAA CGA CG CCTAT CAG CTG GGG TTCNM_001830.3Cxc11GGC TCT TCT GGT CTG CTT TC GGA GACCTT GGT GGG GTG GAG CG TCTNM_001845.3Cxc110GGG TCA AAGC GAG GTG GAG AGGCT TAT TGA AGC GAG CG TG GGG GTG GAG CCNM_00133.3Cxc110GGG TCA AAGC GAG GTG GAG GG GTG GAG GC GTTA TTG CA GG GGG GTG TNM_00137.2MbpCCG GAC CCA AGA TGA TAACGT TGT GGG GGG GTG GG GG GGG GTG TNM_00137.2GfaqTTT TGA AAC CGA GG CG TTCGC CT CCA AGA CGC GG	CD86	GAG CGG GAT AGT AAC GCT GA	GGC TCT CAC TGC CTT CAC TC	NM_019388.3
CD206CTT CGG GCC TTT GGA ATA ATTAG AAG AGC CCT TGG GTT GANM_008625.2Arg1GTG AAG AAC CCA CGG TCT GTGCC AGA GAT GCT TCC AAC TGNM_007482.3Lgals3GAT CAC AAT CGG GCA CAGATT GAA CGC GGG GTT AAA GTNM_01075.3Igf1TGG ATG CTC TTC AGT TCG TGGCA ACA CTC ATC CAC AAT GCNM_01172475.1Sphk1TCC AGA AAC CCC TG GTA GCCAG CAG GTG GA GAG GTNM_001172475.1Il/mTTG CCAA ACC CCG AAG CACT CTG GAG GGA GTG CAG GTNM_001172475.1Il/aaGGA TAA GCA GAC CCG AAG CACT CTG GAG GGA GAC CTT GGT GGNM_001008700.3Socs3CGT TGA AG GTC TCC GAC AATAT TCT GGG GGC GAC AAG ATNM_007707.3IL/bGGG CT CAA AGG AAA GAA TCTCT TCT GGT ATT GCT TGGNM_008361.3IL-6CAA AGC CAG AGT CCT TCA GAGCC ACT CT TCT GG ACT CCNM_00115942.1IL10CTC CCC TGT GAA AAT AAG AGCGCC TGT TAG AGC CTT GG TCNM_001548.2IL-12aTCA CAA CCA GAA GAA AAAAT AGC GAT CCT TGG GT CCNM_001532.2IL-18TTC GTT GAC AAA AGA GAA GAAAT CG GT CGA GT GG GT CANM_00330.1Tr/faGCC TCT TCT TCT GGT TTAGG GTC TGG GCC ATA GAA CTNM_001369.3Cxcl10GGG TAA AGC GAG CCT TTAGG GT CGA GT CA GA CCG CANM_00137.2Cxl10GGG GTA AAGC GAG GTG CAT ACA TT GGA GAG CT TCNM_00137.2MbpCCG GAC CCA AGA TGA TAACCT TG GG GT GA GG GT GT ANM_010777.3GfapCTC CTG GTA CTG CT CTAGG GTC TAA GG GTG GTG TANM_010777.3GfapCTC CTG GTA CTG CC GT CT <td>Ptgs2</td> <td>TCA TTC ACC AGA CAG ATT GCT</td> <td>AAG CGT TTG CGG TAC TCA TT</td> <td>NM_011198.3</td>	Ptgs2	TCA TTC ACC AGA CAG ATT GCT	AAG CGT TTG CGG TAC TCA TT	NM_011198.3
Arg1GTG AAG AAC CCA CGG TCT GTGCC AGA GAT GCT TCC AAC TGNM_007482.3lgal3GAT CAC AAT CAT GGG CAC AGATT GAA GCG GG GT AAA GTNM_010705.3lgf1TGG ATG CTC TCA GT TCG TG GGGCA AAC AC CT ATC CAC AAT GCNM_001172475.1II1mTTC GAG AAC CCC TGT GTA GCCAG GCA GTG GCA GTG GAA GGNM_001172475.1II1mTTG TGC CAA GTC TGG AGA GCACT CTG GAG GA GT GAA GGNM_00108700.3Socs3GCT TGA CAG TCT CC GAC AATAT TCT GGG GCG GAG AAG ATNM_00108700.3Socs3GGG CTT GAA AGG AAA GAA TCTCT TCT TG GG ACT CGT GGNM_008361.3IL-6CAA AGC CAG AGT CCT TCA GAGCC TT CT TTG GG ATT GCT TGGNM_00168.2IL-10CTC CCC TGT GAA AAT AAG AGCGCC TT CT AG GA CCT TGG TCNM_001548.2IL-12aTCA CAA CCA CAA CCA AAAAT AGC GAT CCT GG CT MM_010548.2IL-18TTC GTT GAC AAA GAA AAAAT AGC GAT CT GG GG TTCNM_008360.1TrjaGCC TCT TCT GC TTAGG GTC TG GG GTT CG CAA ACT CANM_008361.3Cxcl10GG CCT CAT AAG GAA GAA AAAAT AGC GAT CT GG GG TTCNM_008361.3Cxcl10GG GTA AAC CGC AGG CG CTTAG GG TCT GG CC AT CAAA CTNM_00176.3Cxcl10GG GTA AAC CGC AGG CG CTTAG GG CCT CAG GG TCG CAG CG CG CAT CAG ACT CTNM_001876.3Cxcl10GG GTA AAC CGC GAG CG CT TTCG GC CAA GA CGC CT TTCG GC CAA GCG AG CCT TTNM_00137.3GdapCTC CCA GGA TGA AAACC CTT GG GA GG GTG AG CG CT CA CGNM_011333.3CCd13CC GG GAC CCA AGC CA CTCG GG CAA GCA GA CCA CTNM_01037.3 <td>CD206</td> <td>CTT CGG GCC TTT GGA ATA AT</td> <td>TAG AAG AGC CCT TGG GTT GA</td> <td>NM_008625.2</td>	CD206	CTT CGG GCC TTT GGA ATA AT	TAG AAG AGC CCT TGG GTT GA	NM_008625.2
Lgals3GAT CAC AAT CAT GGG CAC AGATT GAA GCG GGG GTT AAA GTNM_010705.3lgf1TGG ATG CTC TTC AGT TCG TGGCA ACA CTC ATC CAC AAT GCNM_010512.4Sphk1TCC AGA AAC CCC TGT GTA GCCAG CAG GTG GCA GT GT AGNM_001172475.1IllmTTG TG C CAA GTC TCG GAG ATGTTC TCA GAG CG GT GAA GG TNM_00108700.3Socs3GGT TGA CAG TCT TCC GAC AATAT TCT GGG GCG GAG AGA GTNM_0003861.3IL1bGGG CCT CAA AGG AAA GAA TCTCT TCT GGG ATT GGT TGGNM_00108700.3IL1bGGG CCT CAA AGG CAA GAA TCTCT TCT GTG ATT GCT TGGNM_008361.3IL-6CAA ACC CAG ACT CCT TCA GAGCC ACT CCT TCT GTG ACT CCNM_001168.1IL10CTC CCC TGT GAA AAT AAG AGCGCC TTG TAG ACA CCT TGG TCNM_001948.2IL-12aTCA CAA CCA TCA GCA GAA GAAACT CAG CAG CCT TG TG GC TTM M_00115942.4.1NM_0019548.2IL-12bATC CAG GC CAG AAG AAAAAT AGC GAT CT GG GCT TCNM_008350.1TnfaGCC TCT TCT CAT TCC TGC TTAGG GTC TGG GCC ATA GAA CTNM_008360.1TnfaGCC TCT TCT CAT TCC TGC TTAGG GTC CGG GCC ATA GAA CTNM_00176.3Cxcl10GGG TAA AGC GAG GTG GAG AGGCT TAT TGA AGC GGG GCA CAT CANM_001333.3Cxcl11GCA CCC AAC CG AGT CA TAGCG TCT GG GCC ATA GAA CTNM_001333.3Cxcl2CAT CCA CGT GTT GG GG AGA GAGCT TAT TGA AGC GGG GG GG GG GG MM_001331.3.2MbpCCG GAC CAA AGG CAG CCT TTCTG GCT CCA GG CTT ANM_010777.3GfapCTC CTG GTA ACG GG CG CT ACAG CCA GCA GG CT TA TG GA GAG CTG GT GT GT MM_01039168.1 <t< td=""><td>Arg1</td><td>GTG AAG AAC CCA CGG TCT GT</td><td>GCC AGA GAT GCT TCC AAC TG</td><td>NM_007482.3</td></t<>	Arg1	GTG AAG AAC CCA CGG TCT GT	GCC AGA GAT GCT TCC AAC TG	NM_007482.3
Igf1TGG ATG CTC TTC AGT TCG TGGCA ACA CTC ATC CAC AAT GCNM_010512.4Sphk1TCC AGA AAC CCC TGT GTA GCCAG CAG TGT GCA GTT GAT GANM_011172475.1II1mTTG TGC CAA GTC TGG AGA TGTTC TCA GAG CGG ATG AAG GTNM_031167.5II4raGGA TAA GCA CAC CCC AAG CACT CTG GAG AGA CTT GGT TGGNM_00108700.3Socs3CGT TGA CAG TCT TCC GAC AATAT TCT GGG GGC GAG AAG ATNM_00108700.3IL1bGGG CCT CAA AGG AAA GAA TCTCT TCT TG GGT ATT GCT TGGNM_003161.6IL-6CAA AGC CAG AGT CT TCA GAGCC ACT CCT TGT GG ACT CCNM_0115942.4IL10CTC CCC TGT GAA AAT AAG AGCGCC ATT GAG ACA CCT TGG TCNM_00115942.4IL-12aTCA CAA CAC AGA GAA AAAAT AGC GAT CCT GAG CTT GC CNM_00115942.4IL-12bATC CAG CGC AAG AAA AAAAT AGC GAT CT GG GG TTCNM_008352.2IL-18TCC GTT CT CAT TCC TGC TTAGG GTC TGG GG TTCNM_008360.1TnfaGCC TCT TCT CAT TCC TGC TTAGG GTC TGG GG TTCNM_00837.2Cxcl10GGG GTAA AGC GAG GTG GAGGCT TAT TGA AAC CGA GCT TANM_008176.3Cxcl22CAT CCA CGT GTG GG GAGGCT TAT TGA AAG CGG TGG GA CGNM_011337.2MbpCCG GAC CCA AGA AAA ACCTT GGG GG AGG CTG TG CG GG MM_011337.2MMp0GfapCTC TG GT AACG CGA CTCAG GCA GCA GCA GCT GAG GGG GTG GT GG GGG GTG TNM_010777.3GfapCTC CTG GT AACG CGA CTCAG GCA GCA GCA GCA GCA GCA GCA GCA GCA	Lgals3	GAT CAC AAT CAT GGG CAC AG	ATT GAA GCG GGG GTT AAA GT	NM_010705.3
Sphk1TCC AGA AAC CCC TGT GTA GCCAG CAG TGT GCA GTT GAT GANM_001172475.1II1mTTG TGC CAA GTC TGG ACA TGTTC TCA GAG CGG ATG AAG GTNM_031167.5II4raGGA TAA GCA GAC CCG AAG CACT CTG GAG AGA CTT GGT TGGNM_001008703.3Socs3GGT TGA CAG TCT TCC GAC AATAT TCT GGG GCG CAG AAG ATNM_007707.3IL1bGGG CCT CAA AGG AAA GAA TCTCT TCT TTG GGT ATT GCT TGGNM_008361.3IL-6CAA AGC CAG ACT CCT TCA GAGCC ACT CCT TGT GG ACT CCNM_01158.1IL10CTC CCT GT GAA AAT AAG AGCGCC ATT GCA GAC CCT TGG TCNM_001158.2IL-12aTCA CAA CCA CAG CA GAT CATGC AGA GCT TCA TTT CC GG CCNM_008352.2IL-18TTC GTT GAC AAA AGA CAG CCTAT CAG TCT GGG CCT GGG GTTCNM_008360.1TnfaGCC TCT TCT CAT TCC TGC TTAGG GTC TGG GCC ATA GAA CTNM_0011369.3Cxcl10GGG TCA AAGG AGA GTG CAGGCT TAT TGA AAG CGG TGG GG TTCNM_008176.3Cxcl2CAT CCA CGT GTT GGC TCATCA TTG GGA CCA GA GG CCTAT TGA GAG CGT GGA GCNM_01133.3Ccl3TTT TGA AAC CAG CAG CCT TTCTG CCT CCA AGA CTT CG GG TGA GGNM_01133.3Ccl3TTT TGA AAC CAG CAG CCT TTCTG GCT CCA AGG CTG CTANM_01137.2MbpCCG GCA CCA AGA TGA AAA ACCTT GG CTG CG GTG ACNM_01137.3GfapCTC CTG TA ACT GGC CGA CTAAG CCA AGC CAG AG CTA ACNM_010277.3GfapCTC GTG AACT GG CGA CTAAG CCA GCA GCA GCA GCA CT CA GG CTG CTA ACNM_01039168.1CD11BCTG GTG CTC TTG GCT CTC ATGCG CAG CTT CAT TCA TTG CT	Igf1	TGG ATG CTC TTC AGT TCG TG	GCA ACA CTC ATC CAC AAT GC	NM_010512.4
IIImTTG TGC CAA GTC TGG AGA TGTTC TCA GAG CGG ATG AAG GTNM_031167.5II4raGGA TAA GCA GAC CCG AAG CACT CTG GAG AGA CTT GGT TGGNM_001008700.3Socs3CGT TGA CAG TCT TCC GAC AATAT TCT GGG GGC GAG AAG ATNM_00777.3IL1bGGG CCT CAA AGG AAA GAA TCTCT TTT GG GT ATT GCT TGGNM_008361.3IL-6CAA AGC CAG AGT CCT TCA GAGCC ACT CCT TGT GG ACT CCNM_010548.2IL10CTC CCC TGT GAA AAT AAG AGCGCC ACT CCT TGT AG ACA CCT TGG TCNM_00115942.1IL-12aTCA CAA CCA TCA GCA GAT CATGC AGA GCT TCA TTT TCA CTCNM_008360.1IL-18TTC GTT GAC AAA GAA CAG CCTAT CAG GCT GG GT TCNM_008360.1TnfaGCC TCT TCT ATTC TGC TTAGG GTC TGG GGC ATA GAA CTNM_0013693.3Cxcl10GGG TAA AGG GAG GTG GAG AGGCT TAT TGA AAC CGG TG GG GT CCNM_0013693.3Cxcl10GGG TAA AGG GAG GTG GAG AGGCT TAT TGA AAC CGG TGA GAC CTNM_001333.3Ccl3TTT TG AA ACA GA GAA ACCTT GCT CTA GG GT GA GCNM_011333.3Cd3TTT GA AAC CA GA GC CT TTCTG GCA AGA GT CA GAA CTNM_011337.2MbpCCG GAC CCA AGA TGA AAA CCTT GGG ATA GG GTG GAG CTNM_011337.2MbpCCG GAC CCA AGA TGA CAA CCCTT GGG ATG GAG GTG GT A AGC CGA CTNM_01027.3GfapCTC CTG GTA ACT GGC CGA CTAAG CCA AGC AGC AGA CTA AGCNM_01027.3MeuNCAG TCC TGT GAG CTT CT GT GGCAG ATA GCC AGC AGC CTA AGC CAG AGC CTA AGCNM_001032960.1	Sphk1	TCC AGA AAC CCC TGT GTA GC	CAG CAG TGT GCA GTT GAT GA	NM_001172475.1
Il4raGGA TAA GCA GAC CCG AAG CACT CTG GAG AGA CTT GGT TGGNM_001008700.3Socs3GGT TGA CAG TCT TCC GAC AATAT TCT GGG GGC GAG AAG ATNM_007707.3IL1bGGG CCT CAA AGG AAA GAA TCTCT TCT TTG GGT ATT GCT TGGNM_00316.3IL-6CAA AGC CAG AGT CCT TCA GAGCC ACT CCT TGT GA CT CCNM_010870.3IL10CTC CCC TG GAA AAT AAG AGCGCC ACT CCT TG GACT CCNM_011548.2IL-12aTCA CAA CCA TCA GCA GAT CATGC AGA GCT TCA TTT TCA CTCNM_008352.2IL-18TTC GTT GAC AAA AGA CAG CCTAT CAG GCT GAG CTT G GG TTCNM_008360.1TnfaGCC TCT TCT CAT TCC TGC TTAGG GTC TGG GC CAT GA ACTNM_008363.3Cxcl10GGG TAA AGG GAG GTG GAG AGGCT TAT TGA AAG CAG CCNM_008176.3Cxcl10GGG TAA AGG CAG CCA TAAGG GTC TAT TGA AAG CAG CCNM_001333.3Ccl3TTT TGA AAC CAG CAG CAG CT TTCTG GCG TCA GGG TGA GGNM_011337.2MbpCCG GAC CAAA ACA AAA CCTT GGG ATG AGA CT CT CA GGNM_011337.3GfapCTC TG GTA ACT GGC CGA CTAAG CCA AGA CT ACA GG CAG CT TTNM_010277.3MbpCGA GC TGA GAG GA GA GA CTCAG GAG GAG GTG GAG CT TTNM_0101277.3GfapCTC GTA ACT GGC CGA CTAAG CCA AGC AGA CTA CAA CCA CCA AGA CTNM_0101277.3Ch11BCTG GTG CTC TTG GCT CT ATGGC AGC CAG CAG CTA CAA GCA CAG CAG CTA CAA GCA CAG CAG CAG CTA CAA GCA CAG AGC CTA CAG AGC CAA GCA CAG CAG CTA CAA GCA CAG CAG CTA CAA GCA CAG CAG CTA CAA GCA CAG AGC CAA GCA CAG CAG CAG CTA CAA GCA CAG CAG CAG CTA CAC GCA GAG CTA ACANM_0101277.3GfapCTC CTG	Il1rn	TTG TGC CAA GTC TGG AGA TG	TTC TCA GAG CGG ATG AAG GT	NM_031167.5
Socs3CGT TGA CAG TCT TCC GAC AATAT TCT GGG GGC GAG AAG ATNM_007707.3IL1bGGG CCT CAA AGG AAA GAA TCTCT TCT TTG GGT ATT GCT TGGNM_008361.3IL-6CAA AGC CAG AGT CCT TCA GAGCC ACT CCT TCT GG ACT CCNM_0131168.1IL10CTC CCC TGT GAA AAT AAG AGCGCC ATT CA CAC CCT TGG TCNM_001548.2IL-12aTCA CAA CCA TCA GCA GAT CATGC AGA GCT TCA TTT TCA CTCNM_001159424.1IL-12bATC CAG CGC AAG AAA GAA AAAAT AGC GAT CCT GAG CTT GCNM_008352.2IL-18TTC GTT GAC AAA AGA CAG CCTAT CAG TCT GGT CTG GGG TTCNM_008360.1TnfaGCC TCT TCT CAT TCC TGC TTAGG GTC TGG GCC ATA GAA CTNM_013693.3Cxcl1GCA CCC AAA CCG AAG TCA TAAGG TGC CAT CAG AGC AGT CTNM_018176.3Cxcl10GGG TAA AGG GAG GTG GAG AGGCT TAT TGA AAG CGG TGA GCNM_011333.3Cd2CAT CCA CGT GTT GGC TCATCA TTG CAT CCA GG CTT TC CAG GC TTATNM_011337.2MbpCCG GAC CCA AGA TGA AAA CCTT GGG ATG GAG GTG GTG TNM_010777.3McuNCGA TCG TGT AGG TTG CGC CATAAG CCA AGC AGA CTA AAG CCA CGA AGC CTA AAG CCA AGC AGT CTA CAG GTG GTG TNM_010177.3NeuNCGA TGC TGT AGG TTG CTG TGCAG ATA TGC TCA GCC AGA AGC AGA CTAGG CCA AGC AGA CTA AGC AGA CCANM_0101039168.1CD11BCTG GTG CTC TTG GCT CTC ATGGC AGC TTC ATT CAT CAT GCT CAT CAT GCTNM_001082960.1	Il4ra	GGA TAA GCA GAC CCG AAG C	ACT CTG GAG AGA CTT GGT TGG	NM_001008700.3
IL1bGGG CCT CAA AGG AAA GAA TCTCT TCT TTG GGT ATT GCT TGGNM_008361.3IL-6CAA AGC CAG AGT CCT TCA GAGCC ACT CCT TCT GTG ACT CCNM_031168.1IL10CTC CCC TGT GAA AAT AAG AGCGCC TTG TAG ACA CCT TGG TCNM_010548.2IL-12aTCA CAA CCA TCA GCA GAT CATGC AGA GCT TCA TTT TCA CTCNM_001159424.1IL-12bATC CAG CGC AAG AAA GAA AAAAT AGC GAT CCT GGG CTT GG CTT CCNM_008360.1IL-18TTC GTT GAC AAA AGA CAG CCTAT CAG TCT GGG GTT CNM_008360.1TnfaGCC TCT TCT CAT TCC TGC TTAGG GTC TGG GCC ATA GAA CTNM_008176.3Cxcl1GCA CCC AAA CGG AGG CTG GAG AGGCT TAT TCA AAG GGG TGA GCNM_011333.3Cxcl10GGG TAA AGG GAG CTG GAG AGGCT TAT TCA AAG CGG TGA GCNM_011333.3Cd2CAT CCA CGT GTT GGC TCATCA TTG GGA TCA TCT TCA GG TT CG GG TTNM_011337.2MbpCCG GAC CCA AGA TGA AAA CCTT GG ATT GCA GAG CTT ANM_010777.3GfapCTC CTG GTA ACT GGC CGA CTAAG CCA GCA AGC CA GCA AGC CTNM_01039168.1CD11BCTG GTG CTC TTG GCT CTC ATGGC AGC TTC ATT CAT CAT GCT NNM_001082960.1	Socs3	CGT TGA CAG TCT TCC GAC AA	TAT TCT GGG GGC GAG AAG AT	NM_007707.3
IL-6CAA AGC CAG AGT CCT TCA GAGCC ACT CCT TCT GTG ACT CCNM_031168.1IL10CTC CCC TGT GAA AAT AAG AGCGCC TTG TAG ACA CCT TGG TCNM_010548.2IL-12aTCA CAA CCA TCA GCA GAT CATGC AGA GCT TCA TTT TCA CTCNM_001159424.1IL-12bATC CAG CGC AAG AAA GAA AAAAT AGC GAT CCT GAG CTT GCNM_008352.2IL-18TTC GTT GAC AAA AGA CAG CCTAT CAG TCT GGC GTTG GGG TTCNM_008360.1TnfaGCC TCT TCT CAT TCC TGC TTAGG GTC TGG GCC ATA GAA CTNM_015943.3Cxcl1GCA CCC AAA CGG AGG TCA TAAGG TGC GGC CATA GAA CTNM_008176.3Cxcl10GGG TAA AGG GAG GTG GAG AGGCT TAT TGA AAG GGG TGA GCNM_021274.2Ccl2CAT CCA CGT GTT GGC TCATCA TTC GGA TCA TCT TGC TGNM_011333.3Cd3TTT TGA AAC CAG CAG CTT TTCTG GCT CCA AGA CTT CA GGNM_011337.2MbpCCG GAC CCA AGA TGA AAA CCTT GG ATG GAG GTG GTG TNM_01077.3GfapCTC CTG GT AGG TTG CTG TGAAG CCA AGC AGC AGC AGCNM_0101077.3NeuNCGA TGC TGT AGG TTG CTG TGCAG ATA TGC TCA GGC AGC AGANM_001039168.1CD11BCTG GTG CTC TTG GCT CTC ATGGC AGC TTC ATT CAT CAT GTNM_001082960.1	IL1b	GGG CCT CAA AGG AAA GAA TC	TCT TCT TTG GGT ATT GCT TGG	NM_008361.3
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CD11BCTG GTG CTC TTG GCT CTC ATGGC AGC TTC ATT CAT CAT GTNM_001082960.1	NeuN	CGA TGC TGT AGG TTG CTG TG	CAG ATA TGC TCA GCC AGC AG	NM_001039168.1
	CD11B	CTG GTG CTC TTG GCT CTC AT	GGC AGC TTC ATT CAT CAT GT	NM_001082960.1

(IL) IL-1 α , IL-1 β , IL-3, IL-4, IL-5, IL-6, IL-9, IL-10, IL-12(p40), IL-12 (p70), IL-13, IL-17, granulocyte colony stimulating factor (G-CSF), interferon (IFN) γ , tumor necrosis factor (TNF) α , chemokine C-X-C motif ligand (CXCL) 1 (KC), chemokine ligand (CCL) 2 (also known as MCP-1), CCL3 (also known as MIP1a), CCL4 (also known as MIP1b) and CCL5 (also known as RANTES). All samples were run in duplicate and data analysed with Bio-Plex Manager 6.0 software. Cytokines and chemokines were classified as cytotoxic, repair/ regeneration or immunomodulatory based on the literature (Colton and Wilcock, 2010; Ransohoff and Perry, 2009) and previous characterization in our lab (Chhor et al., 2013).

2.8. Statistics

Data are presented as means \pm SEM. No animals were excluded from any analysis. Numbers in each experiment are indicated within the text, or in the figure legends. Sample sizes were based on calculations of effect sizes from previous studies on this model within the laboratory (Kaindl et al., 2007). For two experimental groups, t-tests or Mann Whitney *U* test were performed. Where more than two experimental groups were compared an ANOVA was performed and when this was significant (p < 0.05) a *Bonferroni* post-test was performed. The appropriate statistical test was chosen based on data normality (Kolmogorov-Smirnov test). The statistical test performed on each data set (using GraphPad 5.0 software [San Diego, CA, USA]) is indicated in the figure legend or within the text.

3. Results

3.1. TBI increases early cell death, microglial number and cortical cytokine / chemokine levels

Following TBI, CCasp3-positive cells were present in the underlying cortex, thalamic nuclei, hippocampal dentate gyrus, subiculum and striatum, and increased numbers of Iba-1 positive cells were observed mainly in the underlying cortex, hippocampus and striatum, and a qualitative representation is found in Fig. 1. Specifically for Iba-1, numbers of positive cells were increased in the cortex following TBI at P1 by $\approx 150\%$ (contralateral 30.86 ± 2.24 versus ipsilateral 46.64 ± 2.41 ; n = 36) and in the striatum by $\approx 350\%$ (contralateral 12.05 ± 0.77 versus ipsilateral 47.23 ± 2.36 ; n = 36). We analysed the number of Iba-1 positive cells by sex and found no difference between control values for the cortex (contralateral male 25.21 ± 0.74 , n = 20 versus contralateral female 23.97 ± 1.17, n = 16; p = 0.35, *t*-test) or striatum (contralateral male 11.68 ± 1.07 , n = 17 versus contralateral female 12.47 ± 1.15, n = 16; p = 0.55, *t*-test), or in the response to injury in the cortex (ipsilateral male 53.71 ± 2.31, n = 17 versus ipsilateral female 51.66 ± 2.48, n = 16; p = 0.52, *t*-test) and striatum (ipsilateral male 47.97 ± 3.38 , n = 17 versus ipsilateral female 46.43 ± 3.37 , n = 16; p = 0.76, *t*-test). As such, we grouped males and females for the following analysis. The effects of TBI on the expression of 20 cytokines and chemokines were measured in the ipsilateral hemisphere at 6, 14 h and 24 h, and compared to levels in sham animals (Fig. 2). Expression increased for markers associated with each MG/M Φ phenotype at all three time points. We observed that relative to levels in a sham hemisphere, the pro-inflammatory/cytotoxic phenotype markers IL-1ß and CCL3 (MIP1 α) showed the greatest and most persistent increases in expression over time (>5 fold). The prototypical antiinflammatory or reparatory/regenerative cytokine IL-4 and the immunomodulatory cytokine IL-10 were also increased at all three time points in the ipsilateral hemisphere. TNF α and IL-12 (p70) were the only markers that did not significantly increase at any time point.

3.2. TBI induces $MG/M\Phi$ expression of markers of a regenerative/ immunomodulatory phenotype

CD11B-positive MG/M Φ were isolated from whole cortices using MACS technology at 2, 6 and 14 h and 1 and 5 days post-TBI, and gene expression of 12 phenotype markers was measured (Fig. 3). Expression of the prototypical cytotoxic MG/M Φ markers



Fig. 2. Expression of cytokines and chemokines from the ipsilateral hemisphere over time post-TBI. Genes are grouped based on predicted role in inflammation: cytotoxic (CytoT), reparatory/regenerative (R-Regen), and immunomodulatory (Immu-M) based on (Colton, 2009; Prinz et al., 2011). Data are normalized to sham group expression (Sham = 1) and are indicated as means \pm SEM (n = 6–8 animals/group). Data were compared to the corresponding sham group using a Mann-Whitney *U* test. *p < 0.05, **p < 0.01, ***p < 0.001.

CD86 and CD32 was decreased by TBI, and iNOS showed no increase at any time point examined. The cross-phenotype marker Cox-2 (cytotoxic-immunomodulatory) and IL1Rn, which has immunomodulatory functions, were persistently increased by TBI. Two additional immunomodulatory markers, SOCS3 and IL-4r α , showed early increases, but by 1-day post-TBI were reduced to below non-TBI levels. Among the reparatory/regenerative MG/ M Φ markers, Arg1 and Gal3 showed persistent increases following TBI, but IGF-1 and CD206 were decreased.

3.3. Blocking $MG/M\Phi$ activation with minocycline causes improvements in neuropathology at 1 day post-TBI

Early brain injury was assessed via CCasp3 cell counts and using ventricular dilatation, calculated as the ratio of the ventricular size in the ipsilateral vs. the contralateral hemisphere. Animals treated with minocycline had reduced numbers of CCasp3-positive cells in the cortex, hippocampus and striatum (Fig. 4C and D). In agreement with these data, minocycline treatment led to less ventricular dilatation compared with the untreated group at 1 day post-TBI (Fig. 4A and B). Sham groups treated with either vehicle or minocycline displayed no change in ventricular size and had very low levels of CCasp3-positive cells (data not shown).

3.4. Improved neuropathology due to minocycline is accompanied by reduced MG number and altered $MG/M\Phi$ activation

To characterize any relationship between the MG/M Φ inflammatory response and neuroprotection, 1 day following TBI, MG cell numbers were quantified using Iba-1 immunolabeling and the phenotype of isolated CD11B-positive MG/M Φ was assessed. As expected based on previous reports of the effects of minocycline, the numbers of Iba-1-positive cells were decreased in the cortex, hippocampus and striatum of minocycline-treated animals (Fig. 5). Minocycline induced complex changes in the phenotype and cytokine/chemokine expression of MG/M Φ that were isolated post-TBI (Fig. 6). The cytotoxic phenotype markers iNOS and IL-6 were reduced from TBI only levels by minocycline treatment but IL-1 β was increased. Among repair/regeneration markers, treatment with minocycline stimulated a further increase in the expression of Gal3 and Arg1, and reduced the typical loss of IGF1. Among immunomodulatory markers, minocycline increased the expression of IL-1Rn above typical levels but led to an even greater decrease in the expression of IL10.



Fig. 3. Expression of phenotype markers by MG/M Φ isolated at various times post-TBI. Genes are grouped based on predicted role in inflammation: cytotoxic, reparatory/ regenerative (R-Regen), and immunomodulatory based on (Colton, 2009; Prinz et al., 2011). Data are normalized to sham group expression (Sham = 1) and are means ± SEM (n = 5–6 animals/group). Gene expression over time was analysed with a two way repeated measures ANOVA, with a Bonferroni post-test to compare the relative expression for each hemisphere at each time point. Summary of the ANOVA results are presented on each panel (effects of TBI [HEMI]. Results of the post-test are indicated with: p < 0.05, p < 0.01.



Fig. 4. Minocycline improves neuropathology 1-day post-TBI. A) Quantification of ventricular volume at 1 day post-TBI, and B) representative images of cresyl-violet-stained sections from both groups indicating ventricular size. C) Quantification of CCasp-3-positive cell number. D) Representative images of CCasp-3 immunolabeling in the striatum from both groups, scale bar 50 μ m. Data are indicated by means ± SEM (n = 9–18 animals/group), and PBS- and minocycline-treated groups were compared via a Student's *t*-test. *p < 0.05, **p < 0.01.

3.5. Early improvements in neuropathology due to minocycline treatment are lost by 5 days post-TBI

Immunolabeling for MAP2 and MBP were used as surrogates for damage to neurons and myelination by oligodendrocytes respectively. TBI decreased the volume of tissue immunolabeled for MAP2 and MBP, and persistently caused ventricular dilatation in the injured hemisphere at 5 days post-TBI (P14; Fig. 7). MAP-2 immunolabeling was similar to that reported previously in the immature brain (Carlsson et al., 2011; Lingwood et al., 2008), displaying a more diffuse pattern and with less cytoplasmic intensity than in the adult. Despite improvements at 1-day post-TBI, at 5 days post-TBI, in animals treated with minocycline, ventricle size was identical to that in PBS treated TBI animals (Fig. 7A). In accordance with the ventricular data, the loss of MAP2 and MBP immunolabeling was not prevented by treatment with minocycline (Fig. 7B–E).

4. Discussion and conclusions

4.1. Principal results and the TBI model

In our closed-contusion model of paediatric TBI, injury modestly increased the levels of both pro- and anti-inflammatory cytokines/chemokines in the brain as well as the number of MG. Isolated MG/M Φ had only moderate changes in gene expression, and increases specifically in markers for the repair/regeneration and immunomodulatory phenotypes. Blocking inflammation/MG/ $M\Phi$ activation with minocycline decreased MG number, reduced expression of some pro-inflammatory cytokines but was only transiently neuroprotective.

We chose for this study a closed-contusion weight-drop TBI model as it has injury mechanisms similar to those seen in paediatric TBI (Xiong et al., 2013). In particular, within the first 30 days of life, at 7-day old, mice display the most widespread apoptotic injury following TBI (Bittigau et al., 1999). This is also the period of greatest vulnerability to excitotoxic lesion in the rodent (Ikonomidou et al., 1989; McDonald et al., 1988) a likely effect of the reduced compensatory anti-oxidant defences of the immature as opposed to the adult brain (Fan et al., 2003). As apoptotic cell death, excitotoxicity and oxidative stress play crucial roles in the pathogenesis of TBI in the neonate (Ruppel et al., 2001; Zhang et al., 2005); this adds weight to the relative usefulness of modelling TBI at this period of rodent development.

4.2. Cortical tissue damage and injury response following paediatric TBI

Following TBI we observed increased total cortical expression of cytokines and chemokines, as well as dilated ventricles and obvious tissue injury in the thalamus and hippocampus of the injured hemisphere. These observations are generally in agreement with



Fig. 5. Minocycline decreases MG cell number 1-day post-TBI. A) quantification of the number of Iba-1-positive cells at 1 day following TBI in PBS- and minocycline-treated mice within the cortex, hippocampus and striatum. B) Representative Iba-1-positive cells in animals from each group from within the striatum, scale bar 50 μ m. Data are indicated by means ± SEM (n = 9–16 animals/group). PBS- and minocycline-treated groups were compared via a Student's *t*-test. *p < 0.05.

previous reports from this model and TBI in large animal models and humans (Helmy et al., 2011; Kaindl et al., 2007; Moretti et al., 2016; Xiong et al., 2013). Specific comparisons across studies are hampered by differences in models and methods, specifically the use of protein versus gene analysis. However, specific comparisons for gene expression can be made to published gene array data. In this context levels of almost all of the jointly reported cytokines and chemokines are considerably lower in our model than that previously reported in two adult rat models of controlled cortical impact (Matzilevich et al., 2002; Raghavendra Rao et al., 2003; White et al., 2013), and one rat model of fluid-percussion injury (Truettner et al., 2005), but were comparable to that reported in a model of adult rat bilateral prefrontal cortical contusion (He et al., 2004). Protein levels for chemokines and cytokines are also consistently and persistently high in previous reports from adult models of contusion and blast-induced TBI (Bye et al., 2007; He et al., 2004; Kumar et al., 2015; Williams et al., 2007). We suggest that further work is needed to ascertain if any smaller magnitude of cortical pro-inflammatory cytokine release is linked to our observation that neonatal TBI induces a predominantly reparatory/ regenerative or immunomodulatory MG/M Φ phenotype. As MG/ $M\Phi$ are the chief drivers of neuroinflammation, a predominantly anti-inflammatory response might prevent cortical inflammation reaching the levels seen in adult injury models, in which there is a robust cytotoxic/pro-inflammatory MG profile (Kumar et al., 2015).

We also wish to briefly discuss the effects of TBI in the contralateral hemisphere. We noted that gene expression was lower than sham level even in the contralateral hemisphere in MG/MΦ for cytotoxic markers (including CD86 and CD32) and also for the repair-regeneration marker CD206 and the immunomodulatory marker SphK1. This type of remote gene expression change has been previously reported in an adult cortical contusion model where it was reported that these are not simply reduced magnitude changes spilling over from the ipsilateral cortex but that some effects are specific (White et al., 2013). Remote tissue changes (such as in the cortex following spinal cord injury) are considered to be crucial mediators of sensorimotor dysfunction and cognitive impairments (Ajao et al., 2012; Kamper et al., 2013; Kim et al., 2006). The diffusion of inflammatory products setting up a chain reaction, signalling via gap junctions in astrocytes and changes in neuronal activity patterns in distant areas are hypothesized to underpin these remote effects. Disruption of the developmental functions of MG (such as synaptogenesis) is considered to underpin some of the injury associated with damage to the immature brain at the injury site and in the remote regions, see (Tremblay et al., 2011). Furthermore, on going changes in homeostatic functions are associated with neurodegeneration such as in aging (Grabert et al., 2016; Griffin et al., 2006; Hart et al., 2012; Lourbopoulos et al., 2015). As such, additional longitudinal studies in this model are warranted to explore the remote and persisting effects of TBI.

4.3. $MG/M\Phi$ phenotype in paediatric vs. adult TBI and other injury models

This is the first study to assess the phenotype of ex-vivo MG/M Φ over such a comprehensive time course (and with so many markers) after an acute neonatal injury. However, there are interesting temporal studies on isolated MG in an adult TBI model (Kumar et al., 2015), and total cortical inflammation in an adult TBI model (Wang et al., 2013) although the methods of analysis vary making it difficult to directly compare data. However, in adult TBI, the protein expression of *ex vivo* MG for classical pro-inflammatory markers increased over time such that at +5 days all markers were robustly increased, compared to our gene expression data wherein only three cytotoxic markers were moderately increased, with no cohesive time point of change. The authors of the adult TBI study sought to describe the robust predominantly pro-inflammatory or mixed



Fig. 6. Effects of minocycline treatment on the expression of phenotype markers from MG/M Φ isolated 24 h post-TBI. Data are shown normalized to expression in a sham group (Sham = 1) and as means ± SEM (n = 9–16 animals/group). Genes are grouped based on predicted role in inflammation: cytotoxic, reparatory/regenerative (R-Regen), and immunomodulatory based on (Colton, 2009; Prinz et al., 2011). Data were compared with a two way ANOVA, with a Bonferroni post-test to compare the relative expression for each hemisphere (PBS versus TBI). Summary of the ANOVA results are presented on each panel (effects of interaction between variable [INTx], effects of time [TIME] and effects of TBI [HEMI]. Results of the post-test comparing each hemisphere are indicated with: p < 0.05, p < 0.01, m p < 0.001.

phenotype that replaced a transient but specific repair/regeneration and immunomodulatory phenotype as "Mtran" (Kumar et al., 2015). In our paediatric model, since cytotoxic MG/M Φ gene expression was generally low, including at 5 days post-TBI, we conclude it is unlikely that this phenotype occurs in this model, although we would need to validate our data with the same FACS based protein analysis. However, when comparing neonatal and adult studies it is worth considering that robust age-dependent differences in MG/ M Φ gene expression have been reported (Bennett et al., 2016; Butovsky et al., 2014). In brief, the stage of development is likely important for MG/M Φ reactivity and is an important consideration for studies of neuroinflammation.

Another key point in the interpretation of these data is the ability of cells to co-express markers, reflecting the *in vivo* complexity of phenotype descriptors. Our gene expression analysis dose not allow up to determine if there are discrete populations of cells switching phenotype or cell co-expressing different category marker as has been previously reported in adult studies (Bedi et al., 2013; Li et al., 2014; Vogel et al., 2013). Co-expression pattern are likely in our paediatric TBI model, but from the paucity of gene expression changes overall any robust co-expression of markers seems unlikely.

4.4. Mode of action of minocycline as a neuroprotective agent

Minocycline is a second-generation semi-synthetic tetracycline that is best known for reducing pro-inflammatory responses via its effects on MG/M Φ (Homsi et al., 2010; Kobayashi et al., 2013; Ng et al., 2012). Minocycline has been used successfully to reduce brain damage across a diverse range of injury/disease models, such as multiple sclerosis (experimental autoimmune encephalitis), term and preterm brain injury (excitotoxicity and hypoxiaischemia, respectively) and Alzheimer's disease, for review see (Garrido-Mesa et al., 2013). Minocycline successfully reduced TBI severity at our early time point of 1 day post-lesion, despite there being little pro-inflammatory response from MG/M Φ . It is not clear from the literature whether minocycline can act directly on astrocytes (Kernt et al., 2010; Yoon et al., 2012) to facilitate any effect. Nevertheless, several pathological mechanisms involved in TBI are counteracted by minocycline, possibly accounting for the neuroprotection. These include that minocycline increases levels of the anti-apoptotic protein Bcl-2 (Wang et al., 2004) and the chelation of magnesium and calcium (Gonzalez et al., 2007) and also decreases activation of MMPs (Koistinaho et al., 2005) and caspase-1 and caspase-3 (Sanchez Mejia et al., 2001). The



Fig. 7. Lack of improvement in neuropathology in minocycline treated animals 5 days post-TBI. A) Quantification of ventricular size 5 days post-TBI and B) quantification of the volume of tissue loss in the traumatized hemisphere 5 days post-TBI based on MAP-2 immunoreactivity, D) representative photomicrographs of MAP2-immunolabeled sections, scale bar 50 μ m. C) and E) quantification and representative images of the volume of white matter loss in the traumatized hemisphere based on MBP immunoreactivity, scale bar 200 μ m. Data are indicated by means ± SEM (n = 12–18 animals/group). PBS- and minocycline-treated groups were compared via a Student's *t*-test.

protective effect of inhibiting MMPs in this model of paediatric TBI has been previously demonstrated (Sifringer et al., 2007).

4.5. Reasons for absence of long-term neuroprotection by minocycline

This study is not the first to report a limited neuroprotective effect of minocycline (Fernandez-Gomez et al., 2005; Fox et al., 2005; Sriram et al., 2006; Yang et al., 2003). Of particular interest is a transient neuroprotective effect reported in an adult closedcontusion TBI model that is strikingly similar to what we observed. In this adult TBI model, behavioural improvements and reduced lesion volume at 1-day post-TBI were lost by 4 days post-TBI (Bye et al., 2007). An early but transient therapeutic effect of minocycline has also been reported following hypoxic-ischemic injury in the mouse (Fox et al., 2005; Nijboer et al., 2010), indicating that this effect of minocycline is not specific to TBI. A limitation of these previous studies and our current study is that we did not test whether behavioural outcomes were improved, despite no change in neuropathology. The concept of a protective phase of the MG/M Φ response after injury has gained enormous support from studies of adult and neonatal models (Faustino et al., 2011; Hernandez-Ontiveros et al., 2013; Hu et al., 2012; Lalancette-Hebert et al., 2007). We speculate that the lack of persistent neuroprotection with minocycline in this model might be because microglia are attempting to repair the brain. As such, there are short-term positive effects (that might relate to positive effects of minocycline on other cells Koistinaho et al., 2005), but when MG are prevented from attempting to repair the brain in the longer term due to exposure to minocycline these positive gains are neutralised by 5 days. What is apparent however is that as outlined in a (non exhaustive) list of studies in Table 1 there are unclear influences on outcome of species (rat versus mouse) and treatment regime (immediate/early only, versus immediate and continuing). The multitude of differences in experimental conditions and outcome measures preclude any firm conclusions on the influences of these factors on the true neuroprotective ability of minocycline.

5. Conclusions

In summary, despite cortical inflammation and cell death following TBI, MG/MΦ retain the expression of markers of an endogenous repair and regenerative phenotype in this model. Also, there are only moderate increases in total cortical inflammatory markers compared to adult injury models. We identified that using minocycline to modify the activity of MG/MΦ had positive early effects on injury, but did not persistently improve outcome. This work adds considerably to our understanding of neuroinflammation after TBI in a neonatal model by suggesting that further therapy design should focus on supporting repair and regeneration type MG/MΦ activation states rather than blanket immunosuppression.

Conflict of Interest Statement

All authors declare that there are no conflicts of interest.

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