Acute injury in the peripheral nervous system triggers an alternative macrophage response

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Background

Injury to the peripheral nervous system (PNS) induces a well-orchestrated cellular process that leads to the complete disintegration of the nerve segment distal to the lesion site, termed Wallerian degeneration (WD) [1].

Background

The activation of the immune system in neurodegeneration has detrimental as well as beneficial effects. Which aspects of this immune response aggravate the neurodegenerative breakdown and which stimulate regeneration remains an open question. To unravel the neuroprotective aspects of the immune system we focused on a model of acute peripheral nerve injury, in which the immune system was shown to be protective.

Methods

To determine the type of immune response triggered after axotomy of the sciatic nerve, a model for Wallerian degeneration in the peripheral nervous system, we evaluated markers representing the two extremes of a type I and type II immune response (classical vs. alternative) using real-time quantitative polymerase chain reaction (RT-qPCR), western blot, and immunohistochemistry.

Results

Our results showed that acute peripheral nerve injury triggers an anti-inflammatory and immunosuppressive response, rather than a pro-inflammatory response. This was reflected by the complete absence of classical macrophage markers (iNOS, IFNγ, and IL12p40), and the strong up-regulation of tissue repair markers (arginase-1, Ym1, and Trem2). The signal favoring the alternative macrophage environment was induced immediately after nerve damage and appeared to be established within the nerve, well before the infiltration of macrophages. In addition, negative regulators of the innate immune response, as well as the anti-inflammatory cytokine IL-10 were induced. The strict regulation of the immune system dampens the potential tissue damaging effects of an over-activated response.

Conclusions

We here demonstrate that acute peripheral nerve injury triggers an inherent protective environment by inducing the M2 phenotype of macrophages and the expression of arginase-1. We believe that the M2 phenotype, associated with a sterile inflammatory response and tissue repair, might explain their neuroprotective capacity. As such, shifting the neurodegeneration-induced immune responses towards an M2/Th2 response could be an important therapeutic strategy.

Keywords: Innate immune system, Negative regulation, M2, RT-qPCR, Neuroprotection, Wallerian degeneration
Subsequently, these cytokines induce the expression of additional immune mediators such as IL-6, GM-CSF, and IL-10 in both Schwann cells and fibroblasts [3,4,6,7]. The production of MCP-1 and MIP-1α, which reaches a maximum at 1 day after injury, promotes the recruitment of macrophages to the damaged nerves [4,8]. Moreover, mast cells accumulate in the endoneurium of injured nerves [9] and release mediators that contribute to the recruitment of macrophages and neutrophils [9,10]. Infiltration of blood-borne monocytes, which spread over the entire nerve, starts from 2 to 3 days after injury and macrophage accumulation peaks at 7 days post injury [11-14]. These infiltrated immune cells take over the cytokine production and are responsible for the rapid clearance of myelin debris. Before the infiltration of the hematogenous macrophages, local macrophages proliferate and undergo morphological changes consistent with immunophenotypic signs of activation [15]. These resident macrophages, together with SCs, readily contribute to myelin phagocytosis [12,16]. Two to three weeks after injury, the inflammatory response is turned off and macrophages are rapidly eliminated [12,13,17]. Generally, WD is believed to induce a strong pro-inflammatory response, as reflected by the induction of cytokines such as TNF and IFNγ [18-20], and the reported elevation of iNOS [21]. Still, one would expect that neurodegeneration might trigger a more dampened immune response, which is typically associated with sterile inflammation. To address this question we used a model of WD (axotomy of the sciatic nerve) to analyze which type of immune response is being induced. Our results revealed that WD leads to the expression of several negative regulators of the innate immune system. In addition, a predominant M2-like macrophage response could be observed, reflecting the presence of an immunosuppressive milieu necessary to initiate wound repair and restore tissue homeostasis.

**Material and methods**

**Mice work and induction of peripheral nerve injury**

All animal experiments were approved by the local ethics committee (University of Antwerp), and conducted according to the guidelines of the Federation of European Laboratory Animal Science Associations (FELASA). Axotomy experiments of the *N. ischiadicus* (sciatic nerve) were conducted in 6- to 8-week-old C57BL/6 mice as previously described [22]. Briefly, mice were anesthetized with a single intraperitoneal injection of ketamine (Ketalar; Pfizer; 150 mg kg⁻¹) and xylazine (Rompun; Bayer; 10 mg kg⁻¹). An incision was made at the right thigh, and gluteal and hamstring muscles were carefully separated to expose the sciatic nerve. The sciatic nerve was transected and the wound was closed by sutures. The contralateral side was left untouched. For analgesia, bupronorphinum (Temgesic; Schering-Slough; 0.1 mg kg⁻¹) was injected after surgery. Sham operation experiments were performed to evaluate the effect of damage around the nerve, inflicted by the operation, on the gene expression profile.

**Intravenous injection of TLR ligands**

Lipopolysaccharide (LPS; TLR4-ligand) (Sigma; 10 mg kg⁻¹) or triacyl lipopeptide (Pam3Cys; TLR1/2-ligand) (Sigma; 10 mg kg⁻¹) were injected intravenously in 6- to 8-week-old C57BL/6 mice. PBS was injected in the control mice.

**RNA isolation and RT-qPCR**

At defined time points after sciatic nerve transection, the mice were euthanized by inhalation of CO₂. The distal part of the transected *N. ischiadicus* and the contralateral control side were removed; snap frozen and stored at −80 °C until use. The nerves were homogenized in Trizol with a Potter Elvehjem homogenizer, and small fragments were further homogenized by sonication. Total RNA was extracted using the RNeasy Lipid Tissue kit (Qiagen) according to the manufacturer's protocol. The quality of the RNA was verified by gel electrophoresis. DNase treatment was performed with TURBO DNase (Ambion). cDNA was produced using the Superscript III first strand synthesis system for RT-PCR (Invitrogen). Real-time quantitative polymerase chain (RT-qPCR) reactions were done with 10 ng cDNA in SYBR Green I mix and run on an ABI Prism 7900 HT Sequence Detection System (Applied Biosystems). All PCR reactions were performed in triplicate. Primers were designed making use of Primerbank (www.pga.mgh.harvard.edu/primerbank). Primer sequences are listed in Table 1. The RT-qPCR data were normalized according to the method described by Vandesompele et al. [23], by geometric averaging of multiple internal control genes. Processing the raw data and normalization of the relative quantities were computed with an improved version of the ΔΔCt-method [24]. The mRNA expression levels are expressed relative to the basal condition (0 h time point; not operated mice or not injected mice).

**Cell isolation**

As a control for the western blot analysis, peritoneal macrophages were stimulated to induce the M1 or M2 expression profile. Peritoneal macrophages were isolated from adult mice that were injected with 3 % thioglycolate [25]. On day 4 after injection, mice were euthanized and the peritoneal cavity rinsed with ice-cold PBS. Macrophages were collected and resuspended in RPMI with 1 % FCS. Erythrocytes stayed in suspension and were removed after 45 min. The remaining cells were kept in RPMI with 10 % FCS. The next day, dendritic cells in suspension were removed and macrophages were
| Table 1 Primer sequences |
|-------------------------|
| **Negative regulators** |
| A20                     | GAACAGCGATCAGGCCAGG   | GGACAGTTGCGGTTCTTCACATT |
| IkBα                    | TGAAGCGAGAGGACTGACGC  | TCTGAGGATGATTGCAAAGTG   |
| IL-1RA                  | GCTCATTGTGGGATTCTCAA  | CCCAGACAGAACAGAGAGC     |
| MyD88s                  | TGAAGTCGCGCATCGGAC    | CGCGACACCTTTTCTCAAT     |
| SIGIRR                  | GTGCACATGCGCCCTAATTTCC | ATGGACAGACTCTTTCAGCC    |
| SOCS1                   | CTCCGGCTGCTATTTGGGAC  | AAAAGGCACTGAAAGGCTCG    |
| **Immune mediators**    |
| Cox2                    | TGACCACTATTCCTGAAAACCACG | GCACTATCGCTTTCCACATTCAC |
| MIP-1α                  | TTCTCTGATCCATGACACTG  | CGTGGAAGCTCTCGCTGAGTC   |
| IL-1β                   | GCAACTTGTTCGAGACTCACT | ATCTTTCGGGTCCGCTCAACT  |
| IL6                     | TAGTCTCTTCTACCTTCCABA | TTGGTCTTCTAGACCTTCTTC  |
| MCP-1                   | TTTAAAACTCTGGATCGAACC | GCATTAGCTTACATTTCGGGT  |
| **Macrophage markers and M1/M2 markers** |
| AIF1/Il-1β              | ATCAACAAGCAATATCTCTTGATGA | CAGCATTGCCTTACAGGACATA |
| Arg1                    | TGCTCGCGGAGCTGAGAC    | GCCTAGTGAAATCGGGCTTTTT |
| CD11b                   | CCGTGACCTTGGAAGGACTGC | ACCGGCGTGCTGCTGAGTC    |
| Cd41                    | GTCTCAAAACTGCAGCCTGGA | GGCTACGGCGTGAGGAACAT   |
| F3/80                   | ATGGGACACCCACTTTCAAGGC | GCAGACTGAGTGGACACACAA |
| Fizz2                   | CAAACACTARGTACATCTTCC | ACCAGATGCAGTACCCCACCA |
| IFNγ                    | ATGGAGCTCAGCTTCTGATAC | CCATCTCTTGGCCATGCTTC   |
| IL4                     | GGTTCTAACCCTGAGCTGTAG | GGGATAGCTTCTCAGGAGTAT  |
| IL10                    | ATGGGGCAAGAGCCCTTTATTTC | AGCATGTGTTGGACACCTTGTA |
| IL12p40                 | ATGGGTGAGGAAGGCTCTTTT | CGCAAGATCGACGGGGACT    |
| IL-13                   | GCAAAACATCAGAAGACAGCA | GTACAGGAATCCAGGGCTAC   |
| iNOS                    | GTCTCCAGGCAACATACAGAA | GTGGAGGCGTGATGCTAC     |
| MRC1                    | CTCTCGAAGCTTGGACGC    | GGAATTTTCGGTGAGCTGCTTC |
| TREM2                   | GAACCCTCACCATCTCTGTA  | CTCGAAACTCATGACTCT    |
| Ym1                     | AAGAGGAGTTTCCAAAATCGGT | CTTCTGGTATGGTGTTAAGTGA |
| IFNγR1                  | GTCGAGCTTGGAGCGGACT   | ATCCCGAGCATACGAGGCGT   |
| IL4Rα                   | ATTTTCGCTTGGAGCTGACTGA | CGTGGAGAAGCTGAGATTGAT |
| IL10R                   | CCTATCCCCACACGCTGAGA | CGAGTTGATCTTCCGCTACTGT |
| IL13Rβ                  | TACGCCACGCTTGAGGGATT | TGAAGATCGAATTGGCTCGG    |
| **Scavenger receptors** |
| Clec7a                   | ATTTTGGCGGACAAATCTCAGG | GCAAGACTGAGAAAAACTCCTC |
| Ragε                     | ATGCACAGAAGGGGATGCTTT | CTGCTTTGGAATAGACACTCCG |
| **Housekeeping genes**  |
| ACTB                    | GCTTCATTGGGACTTGCATTGA | GCATGCAATGGTTCTTCTTAT |
| B2M                     | ATGCAGGGAGAAAGAATACGGA | AGCTATCTAGGATATTCAATAAGTGGAAA |
| HMB5B                   | GAAACTCGCTTCGCGCTGATT | GTGCGCGTTCCTACATCCTATG |
| RPL13a                  | CCTGCTGCTTCAAGAGTTTGTT | TGTTGTCACTCGCTGGTACCTT |
| TBP                     | TCTACTGTTGAGCTGCGGTAAA | TTCTCATTGATCTGCGAGAAA |
kept in culture in RPMI with 10% FCS. Macrophages were stimulated with either LPS (Sigma; 100 ng ml⁻¹) and IFNγ (R&D Systems; 100 ng ml⁻¹) or IL-4 (R&D Systems; 10 ng ml⁻¹) and IL-13 (R&D Systems; 10 ng ml⁻¹) to generate M1 or M2 macrophages, respectively.

**Western blot analysis**

For western blot analysis, the distal part of the transplanted *N. ischiadicus* and the contralateral control side were carefully removed; snap frozen and stored at −80 °C until use. Protein lysates were prepared in E1A lysis buffer (1% NP-40, 20 mM HEPES (pH 7.9), 250 mM NaCl, 20 mM β-glycerophosphate, 10 mM NaF, 1 mM sodium orthovanadate, 2 mM dithiothreitol, 1 mM EDTA, and a protease inhibitor cocktail) by homogenization in a Potter Elvehjem homogenizer. Total protein concentration was determined by Bradford. Equal amounts of protein lysates (20 to 30 μgram) were separated on NuPAGE gels, transferred to nitrocellulose membranes and analyzed by immunoblotting. Briefly, membranes were blocked using blocking buffer (5% milk in PBS containing 0.1% Tween-20), and incubated overnight at 4 °C with a primary antibody. Secondary HRP-conjugated antibodies were used to visualize antibody signals on films using the ECL system (Thermo Scientific). Antibodies used were anti-arginase-1 (Santa

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**Figure 1** Peripheral nerve injury triggers a transient immune response. RT-qPCR analysis showing the transient induction of immune mediators at different time points after peripheral nerve injury (a), and the induction of negative regulators of the innate immune response (b), as determined by RT-qPCR analysis. RT-qPCR data are analyzed using the ΔΔCt method and mRNA expression levels are expressed relative to the basal condition (=0 h time point). Data presented are the compiled data of different independent experiments, with n representing the number of independent experiments for each gene.
Cruz; 1/1,000), anti-iNOS (BD Transduction Laboratories; 1/1,000) and anti-β-tubulin (Abcam; 1/10,000).

**Immunohistochemistry**
The dissected distal part of the sciatic nerve was fixed overnight in Bouin’s fixative (saturated picric acid, formaldehyde, and glacial acetic acid). The nerves were dehydrated, embedded in paraffin for longitudinal sectioning, and stored until processing. The 4 μm sections were immunohistochemically stained using the avidin-biotin staining technique. Briefly, slides were rehydrated and antigen retrieval was performed by boiling the slides in a citrate buffer (0.1 M sodium citrate buffer; pH 6.0). Endogenous peroxidase activity was blocked by treating the slides with 2 % hydrogen peroxide for 20 min. Next, the slides were incubated for 30 min in blocking buffer (20 % normal goat serum in PBS with 1 % BSA) and incubated overnight with primary antibody at 4 °C. The antibodies used were arginase-1 (Santa Cruz) and iNOS (BD Transduction Laboratories). Sections were treated with avidin-conjugated secondary antibodies for 30 min at room temperature before adding an avidin-biotin complex solution (Vector Laboratories) for 30 min. The signal was visualized by incubating the sections with 3,3’-diaminobenzidine (DAB) in PBS containing 0.1 % hydrogen peroxide. Negative control sections were handled the same way, but in the absence of primary antibody. All sections were counterstained with hematoxylin and dehydrated before mounting. Sections were analyzed with a standard light microscope (Zeiss Axioskop with Olympus UC30 camera). For double immunofluorescence staining, the 4 μm sections of paraffin-embedded sciatic nerves, which were processed for antigen retrieval as described above, were incubated in a blocking solution (1 % milk in PBS) for 30 min and incubated at 4 °C with goat-anti-arginase-1 antibody (Santa Cruz). The next day, a donkey-anti-goat Alexa fluor 488 conjugated secondary antibody (Life Technologies) was applied. After stringent washing, the staining with the second marker was performed with the same procedure, using a marker for macrophages (F4/80; Santa Cruz) and a marker for Schwann cells (S100; Dako) and an Alexa fluor 594 conjugated secondary antibody (Life Technologies). Negative controls, excluding one or both of the primary antibodies, were included in the experiments. The immunofluorescence images were captured on a Zeiss LSM700 confocal microscope, using a 40× (1.3 NA) objective. Frame-by-frame scanning with standard emission settings and excitation with a 488 nm or 555 nm diode laser was used to discriminate the two fluorophores.

**Results**
Wallerian degeneration (WD) induces an immune response that is considered to be predominantly pro-inflammatory by expressing several pro-inflammatory molecules such as TNF, IFNγ, and iNOS [19-21]. To confirm the pro-inflammatory environment, we isolated total RNA of the distal segment of four sciatic nerves isolated at different time points upon axotomy.

### Table 2 Overview of the main functions of immune mediators and negative regulators of the immune system

| Immune mediators | Function |
|------------------|----------|
| MCP-1 (CCL2)     | Monocyte chemoattractant protein-1 (CC-chemokine 2) Chemokine, emigration of monocytes from bone marrow; monocyte recruitment [27] |
| COX2             | Cyclo-oxygenase 2 Production of prostaglandins; vasodilatation, inflammation, platelet disaggregation [28], neuropathic pain [29] |
| IL-1β            | Interleukin-1 beta Pro-inflammatory cytokine; many biological functions: up-regulation of adhesion molecules, influx of neutrophils, induction of additional inflammatory mediators, important in sterile inflammation [30] |
| IL-6             | Interleukin 6 Pleiotropic cytokine; wide range of biological activities in immune regulation, hematopoiesis, and inflammation [31] |
| MIP-1α (CCL3)    | Macrophage inflammatory protein 1 alpha (CC-chemokine 3) Chemokine, recruitment of monocytes to the inflamed tissue [4] |

| Negative regulators | Function |
|---------------------|----------|
| A20                 | - Negative regulator of TLR-signaling. Blocks TLR-mediated signaling by blocking NFκB signaling [32] |
| IkBα                | Inhibitor kappa B alpha Inhibits NFκB signaling by masking the nuclear localization signal, keeping NFκB in its inactive state [33] |
| IL-1RA              | IL-1 receptor antagonist Antagonist of IL-1; binds to IL-1R1 thereby blocking signaling [30] |
| MyD88s              | Myeloid differentiation 88 small Antagonist of the adaptor protein MyD88 [26] |
| SIGIRR              | Single immunoglobulin IL-1R-related molecule Inhibits IL-1 signaling, orphan receptor of IL-1 family with antagonistic properties [34] |
| SOCS1               | Suppressor of cytokine signaling 1 Suppression of cytokine signaling by inhibiting JAK tyrosine kinase [35] |
analyzed three independent experiments and measured the expression of several cytokine and chemokine transcripts using RT-qPCR. In line with literature data, the inflammatory mediators IL-1β, Cox2, MCP-1, and MIP-1α were strongly up-regulated, with maximum expression levels at 24 h after axotomy (Figure 1a) [4]. Strikingly, the expression levels of the inflammatory genes dropped at later time points after axotomy, with most pro-inflammatory genes returning to the basal condition at 48 h. We determined whether this transient immune response was accompanied with the induction of several negative regulators of the immune system and found that IL-1RA displayed a high induction (Figure 1b). Moreover, MyD88small (MyD88s) and IκBα, both negatively regulating NF-κB activation [26], were up-regulated already 4 h after injury. In contrast, two other negative regulators of the NF-κB pathway, A20 and SIGIRR, were not induced. Suppressor of cytokine signaling 1 (SOCS1) was only weakly induced after axotomy at these early time points (Figure 1b). Functions of the immune mediators and negative regulators are shown in Table 2.

While many reports already described the induction of cytokines and chemokines in WD, it is less obvious what type of immune response is triggered by injury in the PNS. Therefore, we decided to focus on gene expression profiles for genes associated with M1 vs. M2 macrophages, representative for the two extremes of a merely pro-inflammatory vs. a merely anti-inflammatory/wound healing phenotype [25,36,37]. The main functions of these genes are described in Table 3. We first determined when macrophages start to accumulate in our model, by analyzing the presence of three universal markers for macrophages (Iba1, CD11b, and F4/80) using RT-qPCR [38,39]. In general, it is considered that a first contribution to the immune response in the nerve is

Table 3 Overview of the main functions of M1 and M2 associated markers

| General macrophage markers | M1 associated markers | M2 associated markers |
|---------------------------|-----------------------|-----------------------|
| CD11b | Cluster of differentiation 11b | Expressed on all myeloid lineage cells |
| F4/80 (EMR1) | EGF-like module-containing mucin-like hormone receptor-like 1 | Expressed on most tissue macrophages |
| Iba1 | Ionized calcium binding adaptor molecule 1 | Expressed on macrophages (and microglia) |
| IFNγ | Interferon gamma | Polarize macrophages towards M1; critical for innate and adaptive immunity against viral and intracellular bacterial infections (also produced by macrophages) [40] |
| IL-12p40 | Interleukin 12 subunit p40 | Pro-inflammatory cytokine; induces Th1 [41], production of cytokines (mainly IFNγ) [42], stimulation cell-mediated immunity against microbial pathogens [43] |
| iNOS | Inducible nitric oxide synthase | Production of nitric oxide (NO) using arginine as a substrate; defense against microorganisms [44] |
| Arg1 | Arginase 1 | Production of L-ornithine and urea using arginine as a substrate, first step to production of polyamines (growth factors in the nervous system) [45]; counteracts iNOS |
| CDH1 | Cadherin-1 | Cell-cell adhesion [46], interaction with CD103 of regulatory T-cells [47] |
| Clec7a (dectin1) | C-type lectin domain family 7, member a | Pattern recognition; non-opsonic beta-glucan receptor [48] |
| FIZZ1 | Found in inflammatory zone | Lipid and sugar metabolism; angiogenesis-promoting factor [49] |
| IL-4 | Interleukin 4 | Th2 effector cytokine: polarize macrophages to M2 [50], IgE class switching, differentiation of CD4+ T-cells into Th2 cells [51] |
| IL-10 | Interleukin 10 | Anti-inflammatory cytokine [52] |
| IL-13 | Interleukin 13 | Th2 effector cytokine: polarize macrophages to M2; fibrosis; defense against parasite infections [53] |
| MRC1 | Mannose receptor type C1 | Pattern recognition, type II response promoting effect [54] |
| Rage | Receptor for advanced glycosylation end products | Pattern recognition; binds endogenous ligands such as S100B to stimulate trophic effects on neurons [55] |
| Trem2 | Triggering receptor expressed on myeloid cells 2 | Pattern recognition [56]; augmentation of apoptotic neuron phagocytosis and attenuation of proinflammatory cytokine secretion [57] |
| Ym1 (CH3L3) | Chitinase 3-like-3 | Weak eosinophils chemoattractant properties [58]; binds heparan sulfate, heparin, GlcN oligers, Th2 promoting effect [59] |
Figure 2 (See legend on next page.)
mediated by resident cells (such as Schwann cells, resident macrophages, and fibroblasts) because blood-borne monocytes infiltrate the nerve only 2 to 3 days after injury [12]. Macrophages, expressing Iba1, CD11b, and F4/80, start to accumulate in the injured nerves from day 3 onwards as determined by RT-qPCR (Figure 2a) and immunohistochemistry (Figure 3). Coinciding with the accumulation of macrophages, a second peak in the immune response could be observed, as shown by the bi-phasic induction of IL-6 and IL-1β expression (Figure 2b). As expected, MCP-1, a chemoattractant for macrophages produced by Schwann cells, is expressed right before macrophage accumulation (Figure 2b). In order to determine the phenotype of the macrophages present in the peripheral nerve after injury, we analyzed markers typically associated with M1 vs. M2 macrophages. None of the M1 markers such as iNOS, IL-12p40, and IFNγ were induced after axotomy at any time point investigated (Figure 2c). On the other hand, the M2-associated genes, arginase-1 and Ym1, were clearly induced. The expression of these genes reached a maximal value at 1 day after axotomy and returned to basal level at day 7. Another typical marker for M2 macrophages, Trem2, was induced from day 3 onwards and its expression level remained elevated till day 14 after axotomy (the latest time point investigated in this study) (Figure 2d). The expression of Trem2 appeared to be mediated by the accumulating macrophages, as its expression level displayed a similar pattern as the general macrophage markers (Figure 2b and d). Some markers (arginase-1 and Ym1) were also slightly induced in sham-operated animals, however this induction was only minor compared to the induction seen after axotomy (Figure 2e). Altogether, these data suggest that acute peripheral nerve injury favors an M2 macrophage environment. Additional analyses confirmed this hypothesis. We found that receptors known to trigger M2 cells (such as IL-4Ra and IL-13Ra1), and to stimulate macrophage suppressor function (such as IL-10R), were induced in injured peripheral nerves at 7 and 14 days after injury (Figure 4a). The IFNγR1 receptor, which characterizes M1 macrophages, was not enhanced (Figure 4b). Moreover, scavenger receptors (such as Clec7a and Rage), which are typically expressed by M2 macrophages, showed an increased expression level after axotomy at the late time points relative to the uninjured control nerve (Figure 4c).

The M2 gene expression profile is typically triggered by the cytokines IL-4 and/or IL-13 [60]. In order to determine if these cytokines play a role in the induction of the alternative macrophage environment after axotomy, their expression level was investigated at early time points using RT-qPCR. The IL-4 expression was hardly detectable at the mRNA level in our model of acute peripheral nerve injury and did not seem to be induced. The IL-13 expression, however, was induced upon axotomy at the earliest time point (4 h) investigated (Figure 4d). Importantly, also the anti-inflammatory cytokine IL-10 was induced after injury (Figure 4d). The high IL-10 and low IL-12p40 expression levels are representative of a typical M2 activation profile (Figures 2c and 4d).

Next we analyzed the macrophage phenotype at protein level by using western blot and immunohistochemistry. As the balance between arginase-1 and iNOS expression is highly indicative of the macrophage phenotype, these two markers were used in the following experiments [36]. Western blot analysis of protein lysates of the distal segment of the sciatic nerve showed an induction of arginase-1 protein after axotomy. Arginase-1 protein was detectable from day 1 after injury and reached a maximal signal at day 3. Albeit showing a small decrease over time, the arginase-1 protein level remained high until day 14 after axotomy (Figure 5a). iNOS was not detectable at any time point by western blot analysis, confirming our RT-qPCR data (Figure 5b). As a positive control, peritoneal macrophages were stimulated in vitro with either IL-4/IL-13 or LPS/IFNγ to obtain M2 and M1 macrophages, respectively. As expected, the M2 macrophages expressed arginase-1 and the M1 macrophages expressed iNOS protein (Figure 5a, b). Immunohistochemistry of paraffin-embedded sciatic nerves confirmed the temporal expression profile for arginase-1 shown by western blot. Arginase-1 is rapidly expressed throughout the entire injured nerve. The expression level peaked at 3 days post injury and remained high until day 14 (Figure 5c). Double immunofluorescence staining revealed that arginase-1 was present in F4/80 positive cells and not in S100 positive Schwann cells, which identifies macrophages as the main source for arginase-1 (Figure 6). While at earlier time points all cells that expressed F4/80 were found to be positive for arginase-1, at later time points arginase-1 negative macrophages were present as

(See figure on previous page.)

**Figure 2** Peripheral nerve injury induces M2 associated genes. RT-qPCR analysis of the immune response up to 14 days post injury, showing macrophage markers (a), cytokine expression (b), M1 associated genes (c), and M2 associated markers (d). Sham operation induces M2 markers, but only to a minor extent compared to acute neurodegeneration (e). The mRNA levels are expressed relative to the basal condition (=0 h time point). Data presented are the compiled data of different independent experiments, with n representing the number of independent experiments for each gene. For sham operation one representative experiment out of two independent experiments is shown.
well. Immunohistochemical staining for iNOS confirmed that this protein was not induced after axotomy (Figure 5c). We only observed strong iNOS staining in blood capillaries in particular regions on the nerve that was present independently of the axotomy, showing that the antibody staining was working properly (Figure 5d).

Finally, we determined whether the M2-predominated immune response triggered after nerve injury is typical for the PNS or whether it is specific for neurodegeneration. To this end, we investigated at different time points the expression of M1 and M2 markers in sciatic nerves from mice intravenously injected with TLR-ligands. We used lipopolysaccharide (LPS), a TLR4 ligand known to induce a classical type 1 immune response, and Pam3Cys, a TLR1/2 ligand. Intravenous injection of LPS as well as Pam3Cys elicited a rapid and strong immune response in the sciatic nerve, as shown by the induction of inflammatory genes such as IL-1β, Cox2, MIP-1α, and MCP-1 (Figure 7a). Interestingly, the pro-inflammatory cytokine IL-12p40 and typical M1 immune mediator iNOS, both representative for a type I immune response, were induced after LPS injection (Figure 7b). Several negative regulators, such as IL-1RA, MyD88s, and SOCS1, which mediate a negative feedback loop, were also induced by LPS injection (Figure 7c). Injection with Pam3Cys, however, clearly induced a mixed immune response as reflected by the expression of the M1 associated cytokine IL-12p40 and the expression of Ym1, which is an M2-associated macrophage marker (Figure 7b). iNOS was not detectable after Pam3Cys injection (Figure 7b) and none of the other M2-associated genes such as arginase-1 and Trem2 were induced (Figure 7b). These data show that a prototypical type I immune response can be observed in the nerve after injection of LPS, while Pam3Cys seems to induce a mixed immune response. Both TLR-mediated responses clearly differed from the immune response induced after acute peripheral nerve injury.

**Discussion**

In response to an infection, a strong pro-inflammatory immune response is triggered. The recruited inflammatory cells are activated when they encounter pathogen-associated molecular products (PAMPs) such as LPS. Hereupon, these cells phagocytose infectious agents and produce pro-inflammatory mediators such as iNOS, IL-12, ROS, and RNS to fight off the invading pathogen. These agents, however, can also cause tissue damage. The innate immune system also detects the presence of endogenous molecules; so-called danger associated molecular patterns (DAMPs) that are only exposed in conditions of injury. Under conditions of cellular stress or injury, one might expect a more dampened, strictly controlled immune response as the cost-benefit ratio is higher. As we and others have shown, pro-inflammatory
mediators such as IL-1β and Cox2 and chemokines such as MIP-1α and MCP-1 are rapidly induced in WD, a model of sterile inflammation in the nerve [3,4]. In the present study we show that the expression of these inflammatory genes is strictly controlled as the mRNA levels of all cytokines and chemokines return to basal level at 48 h. Negative regulators of the pro-inflammatory signaling pathways are induced prior to the decline in inflammatory gene expression, thereby limiting the pro-inflammatory immune response and also the excessive damage caused by the immune system.

Although PAMPs and DAMPs are recognized by the same set of receptors, such as TLRs, they can induce a different set of genes [61]. Bacterial compounds trigger a microbicidal environment and classically activated M1 macrophages; while endogenous molecules seem to activate an inflammatory response associated with genes that mediate tissue repair [61,62]. Since most studies so far focused specifically on the induction of pro-inflammatory mediators, WD in the PNS has always been associated with the induction of a strong pro-inflammatory immune response. We found, however, by analyzing genes associated with M1 and M2 macrophages, that acute peripheral nerve injury rather induces an M2-like macrophage environment. None of the typical pro-inflammatory markers of the M1 subtype of macrophages such as iNOS, IFNγ, and IL-12p40 could be detected, while M2 markers such as arginase-1, Ym1, and Trem2 were highly up-regulated. Intriguingly, other M2 markers like Fizz1 and Cdh1 were not induced. Van den Bossche et al. showed that some M2 markers like Cdh1 are strongly down-regulated by the presence of pro-inflammatory cytokines [63]. This could be the case here as well. The stimulation of the alternative macrophage environment in the nerve appeared to be controlled at the level of IL-13. This cytokine was readily detectable from 4 h after the onset of neurodegeneration, and prior to the expression of arginase-1 and Ym1. IL-13, which is together with IL-4 a central master switch in the M2 phenotype, is typically expressed by macrophages, basophils, mast cells, or activated T cells [64]. Since we detect accumulation of macrophages only from days 2 to 3 onwards, it is less clear at the moment which cells

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**Figure 4** Peripheral nerve injury is associated with the expression of M2-associated cytokines and cytokine receptors. RT-qPCR analysis of the sciatic nerve showing cytokine receptor expression profiles upon injury (a, b), scavenger receptor expression (c), and expression of M2 cytokines (d). The mRNA levels are expressed relative to the basal condition (=0 h time point). Data presented are the compiled data of different independent experiments, with n representing the number of independent experiments for each gene.
Figure 5 Protein analysis of M1 versus M2 markers after PNS injury. Western blot analysis showing the arginase-1 protein induction in axotomized sciatic nerves. The IL-4/IL-13 stimulated peritoneal macrophages were used as a positive control (M2) (a). Western blot analysis showing iNOS protein induction in axotomized sciatic nerves. Peritoneal macrophages stimulated with IFNγ/LPS served as a positive control (M1) (b). For western blot analysis, MW values expressed in kDa are shown at the right side of the blots. Immunohistochemical staining of sciatic nerve slices for arginase-1 and iNOS at different time points after injury (c). Blood vessels in the nerves stain positive for iNOS by immunohistochemical staining (d). Scale bar 50 μm.
are responsible for the early onset expression of IL-13, arginase-1, or Ym1. In the peripheral nerve resident macrophages, mast cells or SCs could be engaged in the expression of IL-13, while neutrophils could contribute to the expression of arginase-1 and Ym1. Neutrophils are recruited to the damaged nerves at day 1 after injury [65], and are suggested to contribute to the expression of tissue repair genes [66]. Our results demonstrate that damage to the nerve establishes a rapid immunosuppressive reaction within the nerve, and this from very early time points on, which appears to be in contrast with another recent report [67]. Shechter et al. described that axotomy of the optic nerve creates a pro-inflammatory environment in the nerve that was later turned into an anti-inflammatory one by infiltrating macrophages [67]. Macrophages have been shown before to play a beneficial role in WD in the PNS, as depleting them impaired functional recovery [68]. By phagocytosing debris, macrophages contribute to regeneration by removing inhibitory myelin debris and paving the way for neurite outgrowth. Pre-existing auto-antibodies have been shown to play an important role in clearance of myelin debris by promoting a macrophage influx and stimulating their phagocytic activity [69]. In addition, macrophages produce neurotrophic factors, thereby supporting regeneration [70]. The protective role of macrophages in WD might also be explained by their phenotype. The M2 macrophages were shown to be neuroprotective in vitro by stimulating neurite outgrowth, while M1 macrophages were neurotoxic to neuronal cell cultures [71]. Moreover, potent inducers of a systemic Th2 switch, such as glatiramer acetate and statins, support the neuroprotection and/or nerve regeneration [72-74]. The Th2-inducing adjuvants, such as IFA and Alum, promote axon regeneration better than the Th1-inducing adjuvant CFA [75,76]. Also Th2 cells support neuronal survival in vitro to a greater extent than Th1 cells [77].

In autoimmune diseases of the PNS such as Guillain-Barré Syndrome (GBS) and chronic inflammatory demyelinating polyneuropathy (CIDP), a Th1 response is associated with the early stages of the disease. During recovery of GBS and CIDP, a shift towards a Th2 response is observed, suggesting a protective role for Th2 responses in these diseases [78-80]. Also from animal models it is apparent that type II immune responses are beneficial; as nasal administration of recombinant IL-4 ameliorates ongoing experimental autoimmune neuritis.
(EAN) and inhibits demyelination [81]. The self-limiting clinical course of GBS might be explained by the induction of IL-4 and IL-10. The role of the immune system in hereditary neuropathies is less well studied. Patients suffering from inherited neuropathies show endoneurial T-cells in their nerve biopsies and some patients even show inflammatory infiltrates [82,83]. Studies with animal models such as the heterozygote P0 mice, a model of Charcot-Marie-Tooth (CMT) 1B neuropathy, clearly show a functional degenerative role for macrophages and
T-cells [84,85]. Unfortunately, the type of immune response triggered in hereditary neuropathies has not been addressed.

In CNS injury, macrophages have been implicated in both exacerbating as well asameliorating tissue damage at the injury site. Kigerl et al. showed that spinal cord injury (SCI) initially induces both M1 and M2 macrophages, however, the M1 phenotype predominates the lesion site after 1 week [71]. The presence of both phenotypes might explain the dual effect of macrophages in this model. Moreover, axonal regeneration after SCI is prevented by an inhibitory environment due to myelin inhibitors. Qui et al. showed that elevating cAMP was sufficient to overcome the myelin-mediated inhibition [86]. Subsequent studies showed that arginase-1 and polyamines played a major protective role downstream of CAMP [45]. By using PNS grafts together with acidic fibroblast growth factor in a model of SCI, the recruited macrophages produced large amounts of arginase-1 and were involved in polyamine synthesis [87]. This strategy significantly improved functional recovery [88]. Altogether these data show a central role for type II immune responses, arginase-1 and downstream polyamines in regeneration.

Lessons from graft implantation in CNS injury showed that PNS tissue induces a permissive environment for regeneration. Macrophages contribute to this permissive environment as spinal cord-injured axons failed to regenerate through peripheral nerve grafts in the absence of CD11b+ cells [68]. We now hypothesize that PNS injury triggers an inherent protective environment by inducing an M2 phenotype of macrophages and arginase-1 expression. This model can further be used to unravel how the environment is induced and to elucidate which protective program needs to be elicited. Currently, it remains to be shown how the alternative macrophage environment is established; however there might be a role for IL-13, a typical M2 inducer, as this cytokine is up-regulated very early after injury and prior to the M2-associated gene expression.

Finally, the induction of the alternative macrophage environment appeared to be triggered specifically in response to neurodegeneration. Our results show that when challenged with bacterial products such as LPS, a typical pro-inflammatory immune response, as reflected by a strong IL-12p40 and iNOS signal and the absence of Ym1 or arginase-1, could be detected within the PNS. Intriguingly, injection with Pam3Cys, a TLR1/2 ligand, induced a mixed response marked by presence of both IL-12p40 and Ym1 induction. TLR2, the co-receptor for TLR1, has been associated before with the induction of a type II gene expression [89,90]. In a recent study we showed that specifically TLR1 was highly induced after acute peripheral nerve injury and hypothesized that it might play a role in detecting neuronal injury [22]. The possible involvement of TLR1/2 in the detection of peripheral nerve injury and in the switch towards the type II gene expression is currently under investigation.

**Conclusion**

In conclusion, we demonstrate that acute peripheral nerve injury induces an inherent protective response with the initiation of several negative feedback loops, limiting excessive tissue damage. Furthermore, we show that an M2-like anti-inflammatory environment is induced, rather than a pro-inflammatory one. Since type II responses have been shown before to be neuroprotective, we believe that instead of inhibiting the immune responses, shifting the macrophage phenotype or type of immune response towards an alternative activation state or type II response would be a better therapeutic strategy to stimulate repair, as this would create a permissive environment for neuronal regeneration.

**Abbreviations**

CFA: Complete freund’s adjuvant; CIDP: Chronic inflammatory demyelinating polyneuropathy; CMT: Charcot-Marie-Tooth; CNS: Central nervous system; DAB: 3,3’-Diaminobenzidine; DAMP: Danger associated molecular pattern; EAN: Experimental autoimmune neuritis; GBS: Guillain-Barré Syndrome; IFA: Incomplete freund’s adjuvant; LPS: Lipopolysaccharide; MyD88s: Myeloid differentiation 88 small; PAMP: Pathogen associated molecular pattern; PNS: Peripheral nervous system; RNS: Reactive nitric species; ROS: Reactive oxygen species; RT-qPCR: Real-time quantitative polymerase chain reaction; SC: Schwann cell; SCI: Spinal cord injury; SOCS1: Suppressor of cytokine signalling 1; WD: Wallerian degeneration.

**Competing interests**

The authors declare that they have no competing interests.

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**Authors’ contributions**

EY carried out the experiments and wrote the manuscript, together with SJ. AC performed the intravenous injections. BA provided expertise on microscopy. SG, LP, and GL assisted in the animal surgery and experiments. LAS and JAVG contributed to the interpretation of the results and discussed analyses. VT and SJ designed the methods and experiments. All the authors read and approved the final version of the manuscript.

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