Accumulation and therapeutic modulation of 6-sulfo LacNAc⁺ dendritic cells in multiple sclerosis

ABSTRACT

Objective: To examine the potential role of 6-sulfo LacNAc⁺ (slan) dendritic cells (DCs) displaying pronounced proinflammatory properties in the pathogenesis of multiple sclerosis (MS).

Methods: We determined the presence of slanDCs in demyelinated brain lesions and CSF samples of patients with MS. In addition, we explored the impact of methylprednisolone, interferon-β, glatiramer acetate, or natalizumab on the frequency of blood-circulating slanDCs in patients with MS. We also evaluated whether interferon-β modulates important proinflammatory capabilities of slanDCs.

Results: SlanDCs accumulate in highly inflammatory brain lesions and are present in the majority of CSF samples of patients with MS. Short-term methylprednisolone administration reduces the percentage of slanDCs in blood of patients with MS and the proportion of tumor necrosis factor-α- or CD150-expressing slanDCs. Long-term interferon-β treatment decreases the percentage of blood-circulating slanDCs in contrast to glatiramer acetate or natalizumab. Furthermore, interferon-β inhibits the secretion of proinflammatory cytokines by slanDCs and their capacity to promote proliferation and differentiation of T cells.

Conclusion: Accumulation of slanDCs in highly inflammatory brain lesions and their presence in CSF indicate that slanDCs may play an important role in the immunopathogenesis of MS. The reduction of blood-circulating slanDCs and the inhibition of their proinflammatory properties by methylprednisolone and interferon-β may contribute to the therapeutic efficiency of these drugs in patients with MS. *Neural Neuroimmunol Neuroinflammation 2014;1:e33; doi: 10.1212/NXI.0000000000000033*

GLOSSARY

CTL = cytotoxic T lymphocyte; DC = dendritic cell; FCS = fetal calf serum; GA = glatiramer acetate; IFN = interferon; Ig = immunoglobulin; IL = interleukin; LPS = lipopolysaccharide; MP = methylprednisolone; MS = multiple sclerosis; NA = natalizumab; PBMC = peripheral blood mononuclear cell; PBS = phosphate-buffered saline; pDC = plasmacytoid DC; RRMS = relapsing-remitting MS; slan = 6-sulfo LacNAc⁺; Th = T helper; TNF = tumor necrosis factor.

Multiple sclerosis (MS) is a frequent chronic inflammatory disease of the CNS that is characterized by damage of myelin sheaths and axonal loss. Various studies revealed that activated CD4⁺ T helper (Th) cells and CD8⁺ cytotoxic T lymphocytes (CTLs) essentially contribute to the inflammatory processes underlying MS.¹ ² Dendritic cells (DCs) display a unique capacity to induce and expand CD4⁺ T cells and CD8⁺ CTLs.³ Recent mouse models demonstrated that DCs essentially contribute to T-cell-mediated inflammation and damage in experimental autoimmune encephalomyelitis.⁴–¹⁰ However, little is known about the role of distinct human DC subsets in MS pathogenesis.

Previously, we described 6-sulfo LacNAc⁺ (slan) DCs (formerly termed M-DC8⁺ DCs) as a particular proinflammatory subset of human blood DCs.¹¹–¹³ We demonstrated that activated slanDCs produce large amounts of various proinflammatory cytokines, display a marked capability to handle immunoglobulin (Ig) G–complexed antigens, and exhibit direct cytotoxic activity.¹²–¹⁶ They also efficiently induce and expand CD4⁺ T cells and CD8⁺ CTLs, promote the differentiation

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of naive CD4+ T lymphocytes into Th1 or Th17/Th1 cells, and activate natural killer cells. SlanDCs can be detected in affected tissues of patients with various inflammatory diseases.

Here, we determined the presence of slanDCs in demyelinated brain lesions and CSF of patients with MS. Furthermore, the impact of drugs approved for MS treatment on the frequency of circulating-blood slanDCs in patients with MS was explored. In addition, the influence of interferon (IFN)-β on various properties was evaluated.

METHODS Study approval. All experiments were approved by the institutional review boards of the University Medical Centre Göttingen or the University Hospitals of Ulm or Dresden. Patients or healthy donors gave their written informed consent.

Brain tissues, CSF, and blood from patients or healthy donors. Brain biopsies of patients with MS were used for immunohistochemical staining. One to 2 MS lesions of different brain regions from 15 patients were analyzed. MS lesions were evaluated based on neuropathologic criteria.

CSF samples from 9 patients diagnosed with MS according to the revised McDonald criteria, 9 patients with noninflammatory neurological disorders (primary headache, musculoskeletal pain, CNS ischemia), 5 patients with viral meningitis, and 5 patients with neuroborreliosis were analyzed for slanDCs.

To assess the impact of immunomodulatory drugs on circulating-blood slanDCs and monocytes from patients with MS, groups of different treatment modalities were selected. Patients diagnosed with relapsing remitting MS (RRMS) or natalizumab (NA; Tysabri, Biogen-Idec, Neustadt, Germany), 2 mGL-glutamine, 100 U/mL penicillin, and 100 µg/mL streptomycin (Biochrom). PBMCs (2 × 10^5 cells/well) were plated on round-bottomed 96-well plates. After 6 hours, PBMCs were stimulated with 100 ng/mL lipopolysaccharide (LPS; Sigma-Aldrich) and T cells were identified by using the monoclonal mouse anti-KiM1P antibody (1:5,000; Department of Neuropathology, University Medical Centre, Göttingen, Germany) and T cells were identified by using the monoclonal mouse anti-CD3 antibody (1:200; Serotec, Düsseldorf, Germany), which were applied overnight at 4°C. After washing procedure, staining with biotin-labeled anti-mouse IgG antibody (GM1P) or anti-rat IgG antibody (CD3) was performed for 30 minutes followed by ExtrAvidin-Peroxidase (all Sigma–Aldrich, Steinheim, Germany) for 60 minutes. Antibody binding was detected with the peroxidase substrate 3,3-diaminobenzidine (Sigma–Aldrich). Degree of demyelination was evaluated by luxol fast blue–periodic acid–Schiff staining. Stained sections were analyzed by Olympus BX60 microscope (Olympus, Hamburg, Germany) and evaluated with CellA software.

For determination of cell in tissue sections, positive stained cells were counted at 100× magnification and average density was calculated per 100 leukocytes detected on the slides.

Fluorescence-activated cell sorting. Peripheral blood mononuclear cells (PBMCs) were prepared by Ficoll-Hypaque (Biochrom, Berlin, Germany) density centrifugation. To determine the frequency of slanDCs, PBMCs were incubated with the slan-reactive antibody M-DC8 followed by FITC-conjugated anti-mouse IgM-specific goat F(ab)2 (Beckmann Coulter, Marseille, France) and PerCP-conjugated anti-HLA-DR (BD Biosciences, Heidelberg, Germany), each for 15 minutes at 4°C. FITC-conjugated anti-mouse IgM without primary antibody was used as negative control. Monocytes were characterized by PerCP-conjugated anti-HLA-DR and APC-conjugated anti-CD14 (BD Biosciences). Negative controls included directly labeled or unlabeled isotype-matched irrelevant antibodies (BD Biosciences).

For further characterization of slanDCs and monocytes during relapse treatment, PBMCs were suspended in culture medium consisting of RPMI 1640 (Biochrom), 5% human AB serum (CC pro, Neustadt, Germany), 2 mM L-glutamine, 100 U/mL penicillin, and 100 µg/mL streptomycin (Biochrom). PBMCs (2 × 10^6 cells/well) were plated on round-bottomed 96-well plates. After 6 hours, PBMCs were stimulated with 100 ng/mL lipopolysaccharide (LPS; Sigma–Aldrich) or 10 µM R848 (InvivoGen, Toulouse, France) in the presence of 0.2 µM Monensin (Biomol, Hamburg, Germany) for tumor necrosis factor (TNF)-α staining and harvested 12 hours later. For analysing CD150 expression, cells were incubated with LPS or R848 for 24 hours.

Intracellular molecules were investigated by cell fixation with ice-cold 4% paraformaldehyde (Merck) for 15 minutes and permeabilized with 0.1% saponin (Merck) in phosphate-buffered saline (PBS; Biochrom) plus 1% fetal calf serum (FCS; Biochrom) for 3 minutes at 4°C. Afterward, cells were incubated with a PE-conjugated anti-TNF-α or PE-conjugated isotype-matched irrelevant antibody (BD Biosciences) for 15 minutes. Direct immunofluorescence staining of the cell surface molecule CD150 was performed using...
PE-conjugated anti-CD150 or PE-conjugated isotype-matched irrelevant antibody (BD Biosciences). Cells were evaluated on a FACSCalibur cytometer (BD Biosciences).

**Immunomagnetic isolation of slanDCs and CD4⁺ T cells.** Blood samples were drawn from healthy donors. Isolation of slanDCs was performed as described previously. The purity of the isolated slanDCs was >90% as determined by flow cytometry. The isolation of CD4⁺ T cells, CD8⁺ T cells, and naïve CD45RA⁺ CD4⁺ T cells was conducted by depletion from PBMCs utilizing immunomagnetic separation according to the manufacturer’s instructions (Miltenyi Biotec, Bergisch-Gladbach, Germany). Flow cytometric analysis showed a purity of >90% of the separated T cells.

**Cytokine assay.** SlanDCs from healthy donors were cultured for 24 hours in the presence or absence of 10³ or 10⁴ U/mL IFN-β. Cells were stimulated with LPS to release cytokines. After another 18 hours supernatants were collected. A commercial ELISA kit (BD Biosciences) was used to measure concentration of cytokines including TNF-α, interleukin (IL)-1β, IL-6, and IL-12p70, following manufacturer’s instructions.

**T-cell proliferation and programming.** Separated slanDCs were cultured in the absence or presence of 10³ or 10⁴ U/mL IFN-β for 6 hours and washed. To perform T-cell proliferation assay, IFN-β-treated slanDCs (1 × 10⁴ cells/well) and allogeneic CD4⁺ T cells or CD8⁺ T cells (1 × 10⁵ cells/well) were co-incubated for 4 days. ³H-thymidine (1 μCi, Hartmann Analytic, Braunschweig, Germany) was added for another 18 hours. Then, harvested cells were evaluated for ³H-thymidine incorporation by using a beta counter (PerkinElmer Life Sciences, Rodgau-Jügesheim, Germany).

For evaluation of T-cell programming, IFN-β-treated slanDCs (1 × 10⁴ cells/well) were co-cultured with allogeneic naïve CD45RA⁺ CD4⁺ T cells (1 × 10⁵ cells/well) and stimulated with LPS. After 8 days, 10 ng/mL phorbol myristate acetate (Merck)
and 1 μg/mL ionomycin (Sigma–Aldrich) were used to restimulate T cells for another 4 hours. To evaluate IFN-γ and IL-4 release, cells were fixed with ice-cold 4% paraformaldehyde for 15 minutes and incubated with 0.1% saponin in PBS (Biochrom) plus 1% FCS for permeabilization for 3 minutes at 4°C. Afterward, cells were incubated with FITC-conjugated anti-IFN-γ and PE-conjugated anti-IL-4 antibody or a PE- or FITC-conjugated isotype-matched irrelevant antibody (all BD Biosciences). After 15 minutes, cells were washed and analyzed by flow cytometry.

**Statistical analysis.** Student t test was used to assess the significance of the results. Values of \( p < 0.05 \), \( p < 0.01 \), and \( p < 0.001 \) were considered as significant.

**RESULTS** Accumulation of slanDCs in highly inflammatory brain lesions of patients with MS. To gain novel insights into the potential role of slanDCs in MS immunopathogenesis, we evaluated the presence of this DC subset in 18 inflammatory demyelinating brain lesions (figure 1A), which were infiltrated by macrophages (figure 1B) and CD3+ T cells (figure 1C). SlanDCs were detected in 17 brain lesions and appeared as irregular-shaped cells with long cytoplasmic processes or round cells in areas with less cell infiltration (figure 1, D and E, table). It is interesting that significantly higher numbers of slanDCs were found in highly active lesions (mean: 155 slanDC/mm², range: 83–211) compared to lesions with less inflammatory activity (moderate activity—mean: 77 slanDC/mm², range: 19–192; mild activity—mean: 38 slanDC/mm², range: 0–77), as shown in figure 1F and the table. Macrophages and CD3+ T cells were also detected at significant frequencies in highly active lesions (figure 1, G and H, table).

Detection of slanDCs in CSF of patients with MS. We also determined the presence of slanDCs in CSF samples of 9 patients with MS and 9 control patients with noninflammatory neurologic diseases. SlanDCs were present in 6 out of 9 CSF samples of patients with MS (figure 2, A and B). In contrast, slanDCs were absent in all analyzed control CSF samples (figure 2B). In addition to the accumulation of slanDCs in highly inflammatory brain lesions, these results provide evidence for a contribution of slanDCs to MS immunopathogenesis. For comparison, the presence of slanDCs in CSF samples of patients with other inflammatory disorders was investigated. As depicted in figure 2B, slanDCs were detectable in CSF samples of 4 out of 5 patients with viral meningitis and 3 out of 5 patients with neuroborreliosis.

MP decreased the percentage of blood slanDCs in patients with MS. In further experiments, the impact of short-term administration of MP on the percentage of blood slanDCs and monocytes in PBMCs from 11 patients with RRMS with disease relapse was explored. Remarkably, the percentage of slanDCs decreased significantly at day 1 and day 2.

**Table**

| Patient no. | Lesion no. | SlanDC/mm² | T cell/mm² | T-cell cuffs | Macrophages/mm² | Demyelinization/remyelination | Degree of inflammation |
|-------------|------------|------------|------------|--------------|-----------------|-------------------------------|-----------------------|
| 1           | 1          | 109        | 228        | +++          | 1,800           | CD/ERM                       | Highly active          |
| 2           | 1          | 32         | 128        | +            | 1,344           | CD/ERM                       | Moderately active      |
| 3           | 1          | 38         | 90         | +            | 1,496           | CD/ERM                       | Mildly active          |
| 4           | 1          | 92         | 120        | ++           | 1,280           | CD/ERM                       | Moderately active      |
| 5           | 1          | 77         | 80         | +            | 1,536           | CD/ERM                       | Mildly active          |
| 5           | 2          | 19         | 170        | ++           | 1,588           | CD/ERM                       | Moderately active      |
| 6           | 1          | 26         | 154        | +            | 1,200           | CD/ERM                       | Mildly active          |
| 7           | 1          | 83         | 256        | +            | 1,680           | CD/ERM                       | Highly active          |
| 8           | 1          | 147        | 320        | ++           | 2,240           | CD/no RM                     | Highly active          |
| 9           | 1          | 77         | 294        | ++           | 1,500           | SD/ERM                       | Moderately active      |
| 10          | 1          | 28         | 166        | +++          | 1,568           | SD/no RM                     | Moderately active      |
| 11          | 1          | 192        | 128        | +            | 1,760           | CD/no RM                     | Moderately active      |
| 12          | 1          | 173        | 243        | +++          | 1,720           | SD/ERM                       | Highly active          |
| 13          | 1          | 0          | 83         | +            | 800             | SD/LRM                       | Mildly active          |
| 14          | 1          | 166        | 115        | +++          | 1,800           | SD/no RM                     | Highly active          |
| 15          | 1          | 96         | 166        | +            | 2,000           | CD/ERM                       | Highly active          |
| 15          | 2          | 51         | 96         | +            | 1,400           | CD/ERM                       | Mildly active          |

Abbreviations: CD = complete demyelination; ERM = early remyelination; LRM = late remyelination; no RM = no remyelination; SD = scattered demyelination; slanDC = 6-sulfo LacNAc+ dendritic cell.
IFN-β reduced the percentage of blood slanDCs in patients with MS. We also explored the impact of long-term treatment of patients with RRMS with IFN-β, GA, or NA on the percentage of slanDCs and monocytes in PBMCs. GA and NA therapy did not influence the percentage of slanDCs and monocytes compared to untreated patients with MS (figure 4, A and B). In contrast to monocytes, IFN-β administration significantly reduced the percentage of slanDCs in PBMCs of all patients with RRMS compared to untreated patients with MS (figure 4, A and B). Notably, the percentage of slanDCs in PBMCs of patients with RRMS who developed neutralizing antibodies during IFN-β treatment was comparable to that of untreated patients with RRMS (figure 4A), indicating that neutralizing anti-IFN-β antibodies can abrogate the effect of IFN-β on slanDCs.

IFN-β impaired cytokine production and slanDC-mediated T-cell proliferation and programming. In further studies, the impact of IFN-β on important immunostimulatory properties of slanDCs was investigated. IFN-β significantly impaired the capacity of LPS-stimulated slanDCs to produce TNF-α, IL-1β, IL-6, and IL-12p70 (figure 4, C–F). SlanDC-mediated proliferation of CD4+ and CD8+ T cells was also significantly reduced by IFN-β (figure 4G and H). In addition, the ability of LPS-stimulated slanDCs to promote the differentiation of naïve CD45RA+ CD4+ T lymphocytes into IFN-γ–expressing proinflammatory Th1 cells was markedly inhibited by IFN-β (figure 4I). These results indicate that IFN-β efficiently impairs important immunostimulatory properties of slanDCs.

DISCUSSION Emerging evidence revealed that human DCs are present in brain lesions and CSF of patients with MS. Thus, the presence of rare DCs expressing the maturation marker CD83 in small numbers of perivascular cuffs in 8 out of 51 tissues from patients with MS has been reported. Furthermore, it was demonstrated that mature CD83+ DCs preferentially located in perivascular cuffs of active MS lesions can produce the proinflammatory cytokine IL-23p19. DCs expressing the costimulatory molecule CD86 were also detected in perivascular cuffs of MS lesions, which can be explained by recent data, indicating that human monocytes migrate across the human blood-brain barrier and differentiate into CD83+ DC-SIGN+ DCs.

Whereas these studies are based on the detection of general marker molecules for DCs, little is known about the presence of distinct human DC subsets in brain tissues of patients with MS. Previously, human plasmacytoid DCs (pDCs) were reported, which are implicated in the pathogenesis of several inflammatory diseases and accumulate in active lesions and inflamed meninges of patients.
with MS. Here, we found that human myeloid slanDCs are detectable in 17 of 18 brain lesions. Remarkably, a significant increase in slanDC number was observed in highly inflammatory brain lesions compared to lesions with less inflammatory activity. Recently, we have documented that slanDCs express the chemokine receptor CXCR4 and that the corresponding ligand CXCL12 promotes their migration. Various studies revealed that CXCL12 is expressed in active as well as inactive MS lesions. However, the expression level of CXCL12 was shown to be higher in active MS lesions than inactive lesions. Thus, the elevated CXCL12 expression and/or the presence of additional chemokines such as CXCL13 in active MS lesions may essentially contribute to the pronounced recruitment of slanDCs in such lesions.
Figure 4  Impact of IFN-β on relevant proinflammatory capabilities of slanDCs

The percentages of (A) 6-sulfo LacNAc⁺ dendritic cells (slanDCs) or (B) monocytes in peripheral blood mononuclear cells (PBMCs) derived from blood of 30 patients with relapsing-remitting multiple sclerosis (RRMS) treated with interferon (IFN)-β, glatiramer acetate (GA), or natalizumab (NA); untreated patients with RRMS (de novo); and healthy donors (CTRL) are depicted. In addition, the percentages of (A) slanDCs or (B) monocytes in PBMCs derived from blood of 4 IFN-β-treated patients with RRMS who developed neutralizing anti-IFN-β antibodies (Abs) are shown. Horizontal bars show the mean values. After immunomagnetic cell sorting, slanDCs were stimulated with lipopolysaccharide (LPS) and cultured with or without IFN-β for 24 hours. Supernatant concentrations of (C) tumor necrosis factor (TNF)-α, (D) interleukin (IL)-1β, (E) IL-6, and (F) IL-12p70 were evaluated by ELISA. IFN-β-pretreated slanDCs were co-cultured with allogeneic (G) CD4⁺ or (H) CD8⁺ T cells. After 4 days, ³H-thymidine incorporation was examined to assess T-cell proliferation. Findings of 3 different analyses are presented as mean ± SEM of triplicate determinations. (I) IFN-β-treated slanDCs were co-cultured with allogeneic naive CD45RA⁺ CD4⁺ T cells and stimulated with LPS. After 8 days of cell culture, expression of IFN-γ or IL-4 of CD4⁺ T cells was evaluated by flow cytometry. Three independent analyses presented similar results. Results of one representative donor are presented. Asterisks indicate statistical significance (* p < 0.05, ** p < 0.01, *** p < 0.001).
SlanDCs were also detectable in the majority of CSF samples. This observation further substantiates previous studies, indicating that numbers of pDCs in CSF of patients with MS are higher than in noninflammatory diseases.\(^{34,35}\) The presence of slanDCs and pDCs in active brain lesions and CSF provide evidence that these human DC subsets may participate in MS immunopathogenesis.

In further studies, the impact of various drugs approved for MS treatment on frequency and properties of slanDCs was explored. Short-term administration of MP significantly decreased the percentage of slanDCs in PBMCs from patients with RRMS. This is in agreement with a previous report demonstrating that MP therapy in patients with MS resulted in a decreased percentage of blood-circulating pDCs and CD11c\(^+\) myeloid DCs.\(^{36}\) The MP-mediated reduction of circulating-blood DCs may be caused by inducing apoptosis.\(^{37}\) MP treatment of patients with RRMS also significantly reduced the proportion of TNF-\(\alpha\)– or CD150-expressing slanDCs after stimulation compared to slanDCs from untreated patients with RRMS. The inhibitory effects of glucocorticoids have also been reported by other studies, demonstrating that these agents can efficiently impair differentiation, maturation, and functional properties of DCs.\(^{38,39}\)

Here, we also described that long-term IFN-\(\beta\) treatment in patients with RRMS significantly decreases the percentage of slanDCs in PBMCs compared to untreated patients with MS, which can be explained by IFN-\(\beta\)-mediated induction of apoptosis.\(^{40}\) Notably, the proportion of slanDCs from patients with RRMS who developed anti-IFN-\(\beta\) antibodies during treatment was comparable to that of untreated patients with RRMS. These findings indicate that neutralizing anti-IFN-\(\beta\) antibodies can abrogate the effect of IFN-\(\beta\) on slanDCs. In addition, we found that IFN-\(\beta\) significantly inhibits the secretion of proinflammatory cytokines by activated slanDCs. The capacity of slanDCs to stimulate CD4\(^+\) and CD8\(^+\) T-cell proliferation and to promote the polarization of naïve CD45RA\(^+\) CD4\(^+\) T lymphocytes into proinflammatory Th1 cells was also markedly impaired by IFN-\(\beta\). These results are in line with previous reports demonstrating that IFN-\(\beta\) impairs the maturation and cytokine release of pDCs and their capacity to stimulate T-cell responses.\(^{41,42}\)

Here, we demonstrate that slanDCs accumulate in highly inflammatory brain lesions and are detectable in CSF samples from patients with MS. These novel findings provide evidence that slanDCs may play a role in MS immunopathogenesis. MP and IFN-\(\beta\) significantly decrease the frequency of blood-circulating slanDCs in patients with MS and efficiently impair proinflammatory properties of this DC subset. These effects may contribute to the therapeutic efficiency of these drugs in patients with MS.
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Accumulation and therapeutic modulation of 6-sulfo LacNAc\(^+\) dendritic cells in multiple sclerosis

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