Immunomodulation of inflammatory leukocyte markers during intravenous immunoglobulin treatment associated with clinical efficacy in chronic inflammatory demyelinating polyradiculoneuropathy

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Abstract

Objective: The objective of the study was to profile leukocyte markers modulated during intravenous immunoglobulin (IVIg) treatment, and to identify markers and immune pathways associated with clinical efficacy of IVIg for chronic inflammatory demyelinating polyradiculoneuropathy (CIDP) with potential for monitoring treatment efficacy.

Methods: Response to IVIg treatment in newly diagnosed IVIg-naïve and established IVIg-experienced patients was assessed by changes in expression of inflammatory leukocyte markers by flow cytometry. The adjusted INCAT disability and Medical Research Council sum scores defined clinical response.

Results: Intravenous immunoglobulin modulated immunopathogenic pathways associated with inflammatory disease in CIDP. Leukocyte markers of clinical efficacy included reduced CD185⁺ follicular helper T cells, increased regulatory markers (CD23 and CD72) on B cells, and reduction in the circulating inflammatory CD16⁺ myeloid dendritic cell (mDC) population and concomitant increase in CD62L and CD195 defining a less inflammatory lymphoid homing mDC phenotype. A decline in inflammatory CD16⁺ dendritic cells was associated with clinical improvement or stability, and correlated with magnitude of improvement in neurological assessment scores, but did not predict relapse. IVIg also induced a nonspecific improvement in regulatory and reduced inflammatory markers not associated with clinical response.

Conclusions: Clinically effective IVIg modulated inflammatory and regulatory pathways associated with ongoing control or resolution of CIDP disease. Some of these markers have potential for monitoring outcome.

Keywords: autoimmune neuropathies, chronic inflammatory demyelinating polyradiculoneuropathy, dendritic cells, disease pathways, Fc-gamma receptors, immunophenotyping, intravenous immunoglobulin G
INTRODUCTION

Chronic inflammatory demyelinating polyradiculoneuropathy (CIDP) is an immunologically heterogeneous autoimmune inflammatory disease resulting in peripheral nerve demyelination, causing profound disability in more than 50% of cases (Franssen & Straver, 2014; Mathey et al., 2015). High-dose intravenous immunoglobulin (IVIg), with a superior safety profile to steroids (Dalakas, 2012), is a preferred first-line treatment with established evidence of efficacy (Hughes et al., 2008). The kinetics of response to IVIg in CIDP is similar to plasma exchange (Dalakas, 2012) and is faster than that reported for corticosteroids (Van den Bergh et al., 2010). Short-term prednisolone or pulsed dexamethasone may induce long-term remission (Eftimov, Vermeulen, van Doorn, Brusse, & van Schaik, 2012), whereas IVIg does not eliminate the underlying cause of disease and ongoing IVIg is required to prevent relapse. Individual optimization of IVIg dosing is required because of individual rates of IgG catabolism (Eftimov, Vermeulen, van Doorn, Brusse, & van Schaik, 2012), whereas IVIg does not eliminate the underlying cause of disease and ongoing IVIg is required to prevent relapse. Individual optimization of IVIg dosing is required because of individual rates of IgG catabolism in order to achieve a pharmacokinetic profile supporting treatment efficacy (Kuitwaard et al., 2013; Rajabally, Wong, & Kearney, 2013).

To accommodate patient-specific metabolism of IVIg, early high-dose IVIg is recommended for initial recovery followed by titration to the minimal effective dose (Mathey & Pollard, 2013), whereas low-dose initial treatment may be insufficient to prevent irreversible axonal loss in some CIDP variants (Dalakas, 2011, 2012; Mathey & Pollard, 2013). However, there are no definitive strategies to predict the optimal IVIg dose nor the response to treatment, suggesting frequent monitoring is required to establish optimal treatment in new patients.

Immunomodulatory effects of IVIg have been observed across diverse leukocyte populations, including immunopathogenic disease pathways characteristic of CIDP (Dalakas, 2011; Ephrem et al., 2005; Mathey & Pollard, 2013; Mathey et al., 2015). Based on preliminary phenotyping of activated leukocytes cultured with IVIg, we hypothesized that biomarkers of efficacy of IVIg may be found on multiple leukocyte populations detected in peripheral blood. This study reports immunophenotyping of major leukocyte populations aimed at identifying changes in leukocyte surface marker expression during IVIg treatment. Markers altered by IVIg treatment were then tested for association with clinical efficacy. The data provide evidence that immune phenotyping before and after IVIg treatment can be used to distinguish responders from nonresponders, and suggest targeting-specific markers and pathways for monitoring clinical response to IVIg in CIDP is feasible.

METHODS

2.1 Ethical review

Recruitment of CIDP patients in New South Wales was approved by the Sydney Local Health District (RPAH Zone) HREC, and in Victoria by individual institutional HRECs. All study participants gave informed consent.

2.2 CIDP patients

Newly diagnosed (IVIg-naive) patients (n = 7) and established (IVIg-experienced) patients (n = 14) were examined over the course of two consecutive IVIg treatment cycles. Diagnosis was by clinical and electrophysiological criteria (typical or atypical, sensory and motor, pure motor, or sensory ataxic) according to European Federation of Neurological Societies/Peripheral Nerve Society guidelines (Van den Bergh et al., 2010). Inclusion and exclusion criteria for newly diagnosed CIDP were as used for the ICE trial (Hughes et al., 2008), excepting patient N9 with sensory CIDP. One patient (E14) received low-dose prednisolone (12 mg/day) and methotrexate (20 mg/week); no other patient received steroids.

2.3 IVIg treatment

The two treatment cycles were included, according to patient group: (1) induction and initial maintenance cycles in newly diagnosed patients and (2) two maintenance cycles in patients on established dose-titrated regimens. Standard-of-care IVIg induction dosing was 2 g/kg over 3–5 consecutive days; individually optimized maintenance doses ranged from 0.4 to 1 g/kg every 2–4 weeks according to the Australian guidelines (National IVIg Criteria Review Working Group, 2012).

2.4 Blood collection and clinical assessment

Blood samples were collected prior to IVIg infusion (day 0) and on day 7 during each cycle, and leukocyte marker expression was quantified by flow cytometry. Clinical scores were measured at the start and end of each treatment cycle. For the adjusted INCAT disability score (Hughes et al., 2001), a clinically meaningful change was defined as one point (change between 0 and 1 for upper limb function was not clinically significant in the adjusted INCAT scale). For the Medical Research Council (MRC) sum score (Kleyweg, van der Meche, & Schmitz, 1991), a two-point change was determined to be clinically meaningful. Response in newly diagnosed patients was defined as at least 1 INCAT point improvement during the induction cycle and stable or improved in the second. Patient N9, with a predominantly sensory deficit, was also assessed by the INCAT sensory sum score (Merkies, Schmitz, van der Meche, & van Doorn, 2000). Patients responding to established IVIg regimens were scrutinized for response versus relapse during individual treatment cycles, according to stable versus a clinically meaningful deterioration in either neurological score as defined above.

2.5 Full blood counts and flow cytometry panels

Full blood counts were performed on a CellDyn Ruby automated counter (Abbott Diagnostics, Lane Cove, NSW, Australia) using EDTA-anticoagulated blood. Whole blood flow cytometry was performed on heparinized blood. Antibody clones chosen for the leukocyte markers are listed in Table 3, and all flow cytometry
reagents were sourced from BD Biosciences (North Ryde, NSW, Australia) unless indicated. A large number of markers were screened in order to identify potential changes in population and functional markers, with the expectation that only a few markers would have changes detected in peripheral blood. Flow cytometry was performed by two scientists at a single site using the same flow cytometer and standardized procedures. Antibodies were incubated with blood (100 μL/test) for 15 min before treating with FacsLyse. Tubes for intracellular markers in Treg cells were treated with FacsPerm, washed, and incubated with antibodies for 30 min. Washed leukocytes were run on a BD FACS Canto II, with appropriate compensation settings. Raw.fcs files were analyzed with FlowJo software (Tree Star Inc, Ashland, OR, USA), gates were set against appropriate isotype controls. Marker expression refers to % positive cells in the parent leukocyte population.

2.6 Statistical analysis

Statistical analysis was performed using Prism software (GraphPad, San Diego, CA, USA). An initial screen of markers influenced by IVIg treatment was determined by the Wilcoxon signed-rank test, with significance defined by \( p < .05 \). Markers that changed after IVIg treatment were associated with clinical efficacy after segregating data from responder versus nonresponse/relapse cycles, reporting mean ± SD and paired t tests to determine the magnitude of change in marker expression associated with clinical efficacy. Correction for multiple markers associated with clinical efficacy was applied within each leukocyte population. Difference in marker expression before IVIg treatment between response and relapse cycles was determined by two-tailed Mann–Whitney test. Association between change in marker expression and clinical outcome used Fisher’s exact test. Association between the magnitude of change in marker expression and change in neurology scores was tested by Spearman’s rank correlation coefficient.

3 RESULTS

3.1 Clinical response to IVIg treatment

On the expectation that clinical response could be determined in new CIDP patients after only two IVIg treatment cycles (Hughes et al., 2008), the two initial treatment cycles were used to determine clinical response defined by the disability scores. Two consecutive treatment cycles from patients on established IVIg regimens were studied to compare marker responses with new patients, to determine the stability of markers of clinical response, and to identify markers associated with potential episodes of clinical relapse. Patient details, IVIg regimens, and clinical scores for each treatment cycle are listed in Table 1. Clinically effective IVIg treatment was recorded in 11 of 17 treatment cycles in newly diagnosed and 27 of 32 cycles in established patients. Isolated cycles characterized by clinical relapse were recorded, suggesting that some IVIg doses may have been at the threshold of clinical efficacy in some patients. Clinical response to each treatment cycle was not associated with IVIg dose, pretreatment variables including disability scores, or leukocyte counts (Table 2), although mean lymphocyte count tended to be higher in established patients that relapsed.

3.2 Leukocyte markers influenced by IVIg treatment

The effect of IVIg treatment on circulating leukocyte populations was screened across a wide range of surface antigens representing both subpopulation and functional markers. Multiple redundant markers were tested because it was not known which would be detectable in peripheral blood during an in vivo response to IVIg, compared to our preliminary data from cultured leukocytes. For example, changed expression of CD25, CD38, CD69, CD71, and CD95 were identified after in vitro T-cell activation and subsequent exposure to IVIg. Therefore, only one or two robust markers representing T-cell responses, identified by the initial screen of response to IVIg, were needed for evaluation of clinical efficacy of IVIg treatment. Despite broad changes in markers representing diverse leukocyte populations in our preliminary in vitro cultures, only a few markers changed in peripheral blood sampled after IVIg treatment (Table 3). However, changes in marker expression 7 days after IVIg treatment, measured during two treatment cycles per patient, were found in each major leukocyte population. These changes included decreased follicular helper T cells (CD185+ CD4 T cells), increased Treg cell circulation (reduced CD184) and increased activation (HLA-DR), and naïve B cells with increased inhibitory and regulatory markers (CD23+ and CD72+), but reduced inhibitory CD32b+ B cells. IVIg treatment resulted in a decrease in the inflammatory CD16+ myeloid dendritic cell (mDC) population and a corresponding increase in markers defining a noninflammatory mDC population (CD62L and CD195), whereas monocytes responded with reduced CD32a and CD32b expression.

The rationale to analyze both newly treated and patients on established IVIg regimens together was the need to first identify markers influenced by IVIg, and also IVIg treatment is required to maintain clinical effect through frequent modulation of pathogenic processes. After segregating responder from nonresponse/relapse data, markers associated with clinical efficacy were determined according to increased significance of association (lower \( p \) values) compared with the combined data or opposing direction of change in marker expression between responder and nonresponse/relapse cycles. From the 13 markers found in 7 leukocyte populations that were influenced by IVIg treatment, only 6 markers were specifically associated with clinical efficacy of IVIg treatment in CIDP (Table 4). Correction for multiple testing in these markers using Bonferroni
correction within each leukocyte population resulted in an adjusted level of significance, indicated in Table 4. If a conservative correction was based on all 13 markers, the adjusted level required for significance was \( p = .0038 \), and only four markers remained significant (Table 4). We decided to apply Bonferroni correction within leukocyte populations, because it is immunologically implausible to classify follicular helper T cells and CD16\(^+\) mDCs as linked variables, whereas CD16, CD62L, and CD195 were linked markers on mDCs and required appropriate correction. These markers associated with clinical efficacy signified a change away from an inflammatory profile. But none of these markers were capable of predicting clinical outcome before IVIg treatment. Baseline CD184 expression on Treg

### Table 1: Patient details, intravenous immunoglobulin (IVIg) dose, and neurological response for each treatment cycle

| Patient | Age | Gender | IVIg regimen\(^a\) | Neurological response to IVIg\(^b\) | Observed relapse or nonresponse |
|---------|-----|--------|-------------------|----------------------------------|-------------------------------|
|         |     |        | Cycle 1 | Cycle 2 | Clinical score | Pre-IVIg | Post cycle 1 | Post cycle 2 | |
| Newly diagnosed (IVIg-naïve) patients |
| N1  | 43  | F      | 2.0/4 W | ND     | NCAT\(^c\) | 3 | 2 | ND\(^d\) |
| N2  | 70  | M      | 2.0/4 W | 0.4/4 W | MRC | 52 | 57 |
| N3  | 56  | M      | 2.0/4 W | 0.4/4 W | INCAT | 9 | 5 | 5 |
| N4  | 56  | F      | 2.0/4 W | 0.4/4 W | MRC | 44 | 44 | 44 |
| N5  | 78  | M      | 2.0/3 W | 1.0/3 W | INCAT | 5 | 3 | 4\(^d\) |
| N6  | 78  | M      | 2.0/3 W | 1.0/3 W | MRC | 60 | 60 | 60 |
| N7  | 65  | M      | 2.0/3 W | 1.0/3 W | INCAT | 3 | 2 | 1 |
| N8  | 35  | M      | 1.0/4 W | 0.4/4 W | MRC | 55 | 58 | 56.5 |
| N9  | 75  | M      | 2.0/4 W | 0.4/4 W | INCAT | 5 | 5 | 5 |
| Established (IVIg-experienced) CIDP patients |
| E1  | 76  | F      | 0.2/4 W | 0.4/4 W | MRC | 52 | 52 | 54 |
| E2  | 68  | F      | 1.0/3 W | 1.0/3 W | INCAT | 3 | 3 | 3 |
| E3  | 52  | M      | 0.7/4 W | 0.7/4 W | MRC | 60 | 60 | 60 |
| E4  | 77  | F      | 0.4/4 W | 0.4/4 W | INCAT | 5 | 5 | 4 |
| E5  | 67  | F      | 0.9/4 W | 0.9/4 W | MRC | 54 | 59 | 56 |
| E6  | 67  | M      | 0.4/4 W | 0.4/4 W | INCAT | 6 | 6 | 6 |
| E7  | 70  | M      | 0.8/4 W | 0.8/4 W | MRC | 58 | 60 | 60 |
| E8  | 76  | M      | 0.4/4 W | 0.4/4 W | INCAT | 1 | 1 | 0 |
| E9  | 67  | M      | 0.8/3 W | 0.8/3 W | MRC | 60 | 60 | 60 |
| E10 | 73  | M      | 1.0/3 W | 1.0/3 W | INCAT | 6 | 6 | 6 |
| E11 | 37  | F      | 1.0/3 W | 1.0/3 W | MRC | 57 | 57 | 58 |
| E12 | 80  | M      | 0.4/4 W | 0.4/4 W | INCAT | 5 | 3 | 3 |
| E13 | 66  | M      | 0.4/2 W | 0.4/2 W | MRC | 56 | 56 | 56 |
| E14 | 72  | M      | 0.7/3 W | 0.7/3 W | INCAT | 3 | 3 | 3 |

(Continues)
cells was higher in nonresponders, but this marker was not specifically associated with clinical efficacy (Table 4).

### 3.3 Decreased CD16\(^+\) mDC population is associated with clinical outcome

We next investigated the suitability of these markers to provide early indication of outcome from a treatment cycle, aimed at monitoring efficacy when comparing treatment options, including dose titration. Six markers associated with clinical efficacy were observed during two treatment cycles per patient. A single treatment cycle that defined clinical outcome for each patient was selected (the relapse cycle was chosen in response-discordant patients), and increase versus decrease in marker expression was tested for predicting clinical outcome measured at the end of that cycle (Fisher’s exact test). Of these six markers, only CD16\(^+\) mDCs were directly associated with clinical outcome using Fisher’s exact test for sensitivity and specificity (Fig. 1). Decreased CD16\(^+\) mDCs in responders provided sensitivity for predicting outcome (100%; 95% CI = 80.5–100), but specificity was weak (Fig. 1B), also driving down expression in some relapsing patients. However, the magnitude of decreased CD16\(^+\) mDCs correlated with improved neurological disability scores (MRC and INCAT; Fig. 1C). The MRC sum score defined relapse in established patients, whereas the INCAT generally defined nonresponse in new patients (Table 1).

The immunological relevance of reduced CD16\(^+\) mDCs was further investigated. The CD16\(^+\) mDC population was also characterized as HLA-DR\(^{low}\). Stable clinical response was associated with loss of this population after IVIg, along with a reduced proportion of HLA-DR\(^{++}\) cells expressing CD16 (Fig. 1D). Clinical relapse was associated with either no change or an increase in the CD16\(^+\)HLA-DR\(^{low}\) population (Fig. 1E). Phenotypic profiling of the CD16\(^+\)HLA-DR\(^{low}\) and the CD16\(^-\)HLA-DR\(^{++}\) populations suggested that clinically effective IVIg treatment decreased an activated (CD40\(^+\)CD86\(^+\)), nonlymphoid (CD62L\(^-\), CD195\(^-\), CD197\(^-\)) circulating population that expressed activating Fc\(\gamma\) receptors CD16 and CD32a, but not the inhibitory CD32b. This CD16\(^+\)HLA-DR\(^{low}\) population lacked the activating Fc\(\gamma\)R CD64 associated with monocyte-derived DCs. Although decreased after clinically effective IVIg treatment, this CD16\(^+\) mDC population cycled back to baseline levels measured immediately prior to the following treatment cycle. This suggests de novo regeneration of new

| TABLE 1 (Continued) |
|----------------------|
| | IVIg regimen\(^a\) | Neurological response to IVIg\(^b\) | Observed relapse or nonresponse |
|----------------------|----------------|------------------|--------------------------|
| Patient | Age | Gender | Cycle 1 | Cycle 2 | Clinical score | Pre-IVIg | Post cycle 1 | Post cycle 2 | |
| E15 | 70 | M | 0.4/4 W | 0.4/4 W | INCAT | 3 | 3 | 3 |
| E16 | 52 | M | 0.4/4 W | 0.4/4 W | MRC | 59 | 59 | 59 |
| | | | | | INCAT | 2 | 2 | 1 |
| | | | | | MRC | 60 | 60 | 60 |

ND, second treatment cycle not given.
\(^a\)IVIg dose (g/kg) and cycle length in weeks (W).
\(^b\)Neurological scores that defined relapse or nonresponse are given in bold.
\(^1\)ICAT disability score (0–10; 0 = normal function), MRC sum score (max. 60 = normal function), and INCAT-S (sensory sum score) (0–20; 0 = normal function).
\(^2\)Missing data: N1 withdrew from the study after the first cycle and blood samples were not collected from the second cycle of N3.

| TABLE 2 Pretreatment variables, disability scores, and leukocyte counts were not associated with clinical response |
|---------------------------------------------------------------|
| Pretreatment variables | Newly diagnosed (IVIg-naïve) CIDP | Established (IVIg-experienced) CIDP |
|------------------------|---------------------------------|---------------------------------|
| IVIg cycles (n) | 11 | 6 | 27 | 5 | .880 |
| IVIg dose | 1.35 ± 0.73 | 1.35 ± 0.79 | .903 | 0.59 ± 0.26 | 0.61 ± 0.22 | .815 |
| Steroid use | Nil | Nil | 6 | 64.8 ± 7.5 | .415 |
| Age at treatment | 66.1 ± 9.4 | 69.3 ± 10.8 | .641 | 68.0 ± 11.6 | .415 |
| INCAT disability | 4.6 ± 2.4 | 4.5 ± 1.0 | .903 | 2.8 ± 1.3 | 3.0 ± 1.9 | .815 |
| MRC sum score | 51.6 ± 7.1 | 55.5 ± 3.0 | .216 | 53.0 ± 5.5 | 54.4 ± 6.6 | .692 |
| Lymphocytes (10\(^9\)/L) | 1.77 ± 0.32 | 1.64 ± 0.64 | .713 | 2.03 ± 0.81 | 3.07 ± 0.95 | .070 |
| Monocytes (10\(^9\)/L) | 0.54 ± 0.13 | 0.71 ± 0.48 | .522 | 0.64 ± 0.45 | 0.79 ± 0.53 | .580 |

\(^a\)Mean ± SD and Mann–Whitney test.
inflammatory mDCs between treatment cycles is a possible cause of ongoing inflammation in CIDP requiring life-long IVIg treatment.

## 4 | DISCUSSION

This study confirmed some of the known immunomodulatory action of IVIg on disease pathways of inflammatory neuropathies, and identified markers associated with clinical outcome. Changes in disease pathways associated with clinical efficacy included reduced capacity for autoantibody production associated with decreased follicular helper T cells and increased expression of regulatory and tolerance markers on B cells. A decrease in circulating inflammatory CD16+ mDCs was associated with clinical efficacy. In addition, markers associated with a reduced inflammatory profile, independent of clinical response, may signify the overall nonspecific anti-inflammatory effects of IVIg which in combination may contribute to clinical efficacy. Although IVIg may induce long-term

| Leukocyte population (phenotype) | Investigational markers | Mab clone   | Changed in marker expressiona |
|----------------------------------|-------------------------|-------------|-----------------------------|
| T cells (CD45+CD3+CD4+)          |                         | CD4+        | CD4-                        |
| CD25                             | M-A251                  | 0.819       | 0.330                       |
| CD26                             | M-A261                  | 1.000       | 1.000                       |
| CD27                             | M-T271                  | 1.000       | 0.750                       |
| CD28                             | CD28.2                  | 0.160       | 0.164                       |
| CD38                             | HIT2                    | 0.895       | 0.090                       |
| CD69                             | L78                     | 0.055       | 0.055                       |
| CD71                             | M-A712                  | 0.313       | 0.742                       |
| CD95                             | CX2                     | 0.755       | 0.319                       |
| CD103                            | Ber-ACT8                | 0.844       | 0.469                       |
| CD120b                           | hTNFR-M1                | 0.859       | 0.190                       |
| CD184                            | 12G5                    | 0.335       | 0.516                       |
| CD185                            | 51505                   | 0.042       | 0.213 Decrease              |
| CD195                            | 2D7/CCR5                | 0.960       | 0.402                       |
| CD197                            | 150503                  | 0.282       | 0.013 Decrease              |

### Regulatory T cells (CD45+CD4+CD25Hi CD127Lo)

| Leukocyte population (phenotype) | Investigational markers | Mab clone   | Changed in marker expressiona |
|----------------------------------|-------------------------|-------------|-----------------------------|
| CD62L                            |                         | DREG-56     | 0.192                       |
| CD184                            |                         | 12G5        | 0.034 Decrease              |
| CD195                            |                         | 2D7/CCR5    | 0.140                       |
| HLA-DR                           |                         | G46-6       | 0.011 Increase              |
| CD152                            |                         | BN13        | 0.470                       |
| FoxP3                            |                         | 259D/C7     | 0.110                       |
| TGFβ                             |                         | 9016        | 0.695                       |

### Naive and memory B cells (CD19+ CD27+)

| Leukocyte population (phenotype) | Investigational markers | Mab clone   | Changed in marker expressiona |
|----------------------------------|-------------------------|-------------|-----------------------------|
| CD23                             | M-L233                  | 0.0097      | 0.976 Increase              |
| CD32/32b                         | FL18.26/X63             | 0.012       | 0.0004 Decrease             |
| CD32a                            | IV                      | 0.899       | 0.577                       |
| CD40                             | 5C3                     | 0.967       | 0.095                       |
| CD69                             | L78                     | 0.599       | 0.969                       |
| CD70                             | Ki-24                   | 0.898       | 0.247                       |
| CD72                             | J4-117                  | 0.007       | 0.375 Increase              |
| CD80                             | L307.4                  | 0.641       | 0.358                       |
| CD86                             | 2331 FUN-1              | 0.979       | 0.360                       |
| HLA-DR                           |                         | G46-6       | 0.054                       |

### Myeloid dendritic cells (CD45+HLA-DR+CD11c lineage-neg)

| Leukocyte population (phenotype) | Investigational markers | Mab clone   | Changed in marker expressiona |
|----------------------------------|-------------------------|-------------|-----------------------------|
| CD16                             | 3G8                     | 0.0017      |                             |
| CD32/32a                         | FL18.26/IV.3            | 0.468       |                             |
| CD32b                            | X63                     | 0.946       |                             |
| CD40                             | 5C3                     |             |                             |
| CD62L                            | DREG-56                 | 0.0006      |                             |
| CD64                             | 10.1                    | 0.368       |                             |
| CD83                             | HB15e                   | 0.091       |                             |
| CD86                             | 2331 FUN-1              | 0.814       |                             |
| CD184                            | 12G5                    | 0.995       |                             |
| CD195                            | 2D7/CCR5                | 0.0077      |                             |
| CD300f                           | UP-D2                   | 0.356       |                             |

(Continues)
immunomodulatory changes, including inhibitory Fcγ receptor expression, or autoreactive T-cell populations (Klehmet et al., 2015), most of the anti-inflammatory effects of IVIg observed in our study were short-lived, suggesting the underlying causes of CIDP are modulated but not removed by IVIg in most patients. It is thought that the transitory effect of IVIg on clinical outcome may be a consequence of catabolism of infused IgG, impacting on direct competition with pathogenic autoantibodies and complement inactivation (Berger, McCallus, & Lin, 2013). The transitory effect of IVIg treatment on markers identified in this study may likewise reflect the decline in IVIg interactions with Fcγ receptors and the downstream regulation of B cell, mDC and follicular helper T-cell-mediated responses.

The phenotypic profile of the CD16⁺HLA-DRlow mDC population suggests these were mature (CD195⁻) and inflammatory (CD40⁺CD86⁺), but unlike conventional mature mDCs, locked in peripheral circulation with impaired lymphoid migration (CD62L⁻CD185⁻CD195⁻; Kim & Diamond, 2015; Lutz & Schuler, 2002; Martin-Fontecha, Lanzavecchia, & Sallusto, 2009; Sallusto et al., 1998). Furthermore, these CD16⁺HLA-DRlow mDCs had receptors for inflammatory (CD16, CD32a/b) but not regulatory (CD32b) antibody interaction. Reduction of this CD16⁺HLA-DRlow mDC population resulted in a relative increase of lymph-node homing markers (CD62L and CD195) in the remaining CD16⁺HLA-DR++-enriched mDC population, conferring reduced inflammatory potential (Wildenberg et al., 2008). The origin of this CD16⁺HLA-DRlow population is controversial, some suggesting a monocytic origin (Kim & Diamond, 2015; Randolph, Sanchez-Schmitz, Liebman, & Schakel, 2002). However, low CD205 expression and lack of CD64 on this population suggest otherwise. Regardless of the source, accumulation between treatment cycles of a circulating activated inflammatory mDC population deficient for lymph-node homing markers may be a major immunopathogenic pathway of CIDP disease, and depletion of this population may be a critical immunoregulatory action of IVIg. The relative increase in CD195 and CD62L defining the CD16⁺HLA-DR++ mDC population after clinically effective IVIg suggests increased lymph-node homing and reduced peripheral tissue inflammation is another immunoregulatory action of IVIg. However, lack of key homing markers on the CD16⁺HLA-DRlow mDC population does not explain removal from circulation after IVIg.

B-cell activation and generation of autoantibodies are considered critical immunopathogenic pathways in CIDP and other autoimmune neuropathies. Data from this study showed that IVIg increased both regulatory (CD23) and peripheral tolerance (CD72) markers on naïve B cells. The inhibitory Fcγ receptor CD32b was expressed on more memory B cells before IVIg treatment in responders than nonresponders, but was not directly associated with clinical outcome. CD185 expression defines follicular helper T cells with increased homing for B-cell germinal centers and is elevated in a number of autoimmune diseases (Park et al., 2014). The observed decrease in follicular helper T cells after IVIg may reduce autoantibody production as well as macrophage activation pathways (Schmitt, Bentebibel, & Ueno, 2014; Slight et al., 2013).

Inhibitory Fcγ receptor-mediated pathways of immunomodulation in B cells and monocytes are considered primary mechanisms of IVIg immunomodulation (Tackenberg et al., 2009), and was confirmed by data from this study. Expression of CD32b decreased after IVIg on both monocytes and memory B cells, as reported on monocytes in IVIg-treated immune thrombocytopenia (Shimomura et al., 2012). Downregulation of CD32b on B cells may result from increased CD40 ligation (Zhang et al., 2013). Alternatively, an increase in the balance of type-2 and regulatory cytokines induced by IVIg treatment may promote CD32b expression (Boruchov et al., 2005), and transitory CD32b downregulation observed in this study

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**TABLE 3** (Continued)

| Leukocyte population (phenotype) | Investigational markers | Mab clone | Changed in marker expression |
|----------------------------------|-------------------------|-----------|-----------------------------|
| Monocytes (CD45⁺CD14⁻)           | CD16                    | B73.1     | 0.528                        |
| CD32⁺/CD32a/b                    | FL1B.26/IV.3            | 0.0057    | Decrease                    |
| CD32b                            | X63                     | 0.050     | Decrease                    |
| CD38                             | HIT2                    | 0.149     |                              |
| CD40                             | SC3                     | 0.179     |                              |
| CD64                             | 10.1                    | 0.229     |                              |
| CD69                             | L7B                     | 0.632     |                              |
| CD80                             | L307.4                  | 0.549     |                              |
| CD86                             | 2331 FUN-1              | 0.843     |                              |
| CD163                            | GHI/61                  | 0.872     |                              |
| CD184                            | 12G5                    | 0.159     |                              |
| CD195                            | 2D7/CCR5                | 0.547     |                              |
| CD204a/b                         | SR-41/MSR1              | 0.139     |                              |
| CD206                            | 19.2                    | 0.484     |                              |
| CD300f                           | UP-D2                   | 0.077     |                              |
| HLA-DR                           | G46-6                   | 0.547     |                              |

a Change in expression was determined by Wilcoxon’s signed-rank test; significant changes (p ≤ 0.05) are in bold font.

All antibodies were from BD Pharmingen unless indicated: *R&D Systems; *Biolegend; and *CD32 was replaced during the study with isoform-specific CD32a and CD32b antibodies produced by HMT and PMH (Ramsland et al., 2011). CD32 expression on dendritic cells and monocytes is equivalent to CD32b, and CD32 on B cells is equivalent to CD32b, and therefore data were pooled where appropriate.

b Gated on cells negative for lineage markers CD3, CD14, CD19, CD20, CD34, CD56, and CD66.
### TABLE 4  Effect of intravenous immunoglobulin (IVIg) treatment on leukocyte populations and markers associated with clinical efficacy in chronic inflammatory demyelinating polyradiculoneuropathy

| Population         | Marker                 | Wilcoxon test | Response to IVIg (all patients) | Responders (clinical efficacy) | Nonresponse/relapse | Pre-IVIg predictor (response vs. relapse) |
|--------------------|------------------------|---------------|---------------------------------|--------------------------------|---------------------|------------------------------------------|
|                    |                        |               | Wilcoxon test | Pre-IVIg | Post IVIg | Paired t test | Pre-IVIg | Post IVIg | Paired t test | Pre-IVIg | Post IVIg | Paired t test | Mann–Whitney |
| CD4 T cells        | CD185                  | 0.042         | 19.2 ± 7.4     | 17.5 ± 7.3 | 0.013    | 19.4 ± 8.0     | 17.2 ± 7.9 | 0.008†     | 18.4 ± 4.8     | 18.8 ± 4.1 | 0.645     | 0.616         |
|                    | CD197                  | 0.013         | 27.7 ± 14.7    | 25.4 ± 14.8 | 0.020    | 27.6 ± 15.2    | 25.9 ± 15.9 | 0.120     | 28.3 ± 13.5    | 23.5 ± 10.1 | 0.059     | 0.885         |
| Tregs              | CD184                  | 0.034         | 64.6 ± 19.6    | 59.0 ± 20.8 | 0.033    | 61.1 ± 20.8    | 55.9 ± 20.4 | 0.097     | 74.7 ± 12.0    | 68.1 ± 20.3 | 0.190     | 0.029         |
|                    | HLA-DR                 | 0.011         | 33.8 ± 14.7    | 36.1 ± 13.5 | 0.016    | 34.0 ± 15.1    | 35.9 ± 13.6 | 1.000     | 33.2 ± 14.5    | 36.4 ± 13.9 | 0.026     | 0.914         |
| Naïve B cells      | CD23                   | 0.010         | 54.8 ± 23.9    | 60.3 ± 19.6 | 0.008    | 54.2 ± 25.0    | 61.6 ± 20.8 | 0.009**    | 56.6 ± 20.6    | 55.7 ± 14.8 | 0.884     | 0.811         |
|                    | CD72                   | 0.007         | 84.2 ± 12.9    | 86.1 ± 11.4 | 0.159    | 83.9 ± 13.2    | 87.3 ± 11.3 | 0.010     | 85.4 ± 12.4    | 86.1 ± 11.6 | 0.426     | 0.739         |
|                    | CD32b                  | 0.020         | 43.0 ± 20.3    | 36.6 ± 22.5 | 0.012    | 43.2 ± 20.4    | 36.3 ± 22.7 | 0.031     | 42.4 ± 19.6    | 37.3 ± 23.1 | 0.202     | 0.918         |
| Memory B cells     | CD32b                  | 0.0002        | 54.0 ± 21.7    | 44.8 ± 24.5 | 0.0004   | 55.1 ± 22.2    | 46.9 ± 25.4 | 0.002     | 49.9 ± 20.3    | 37.6 ± 20.6 | 0.082     | 0.521         |
| Myeloid dendritic cells | CD16 | 0.002          | 57.1 ± 18.1    | 48.5 ± 20.9 | 0.0004   | 58.0 ± 18.4    | 46.4 ± 21.7 | <0.0001*   | 53.8 ± 17.2    | 56.5 ± 16.0 | 0.433     | 0.500         |
|                    | CD62L                  | 0.0006        | 26.0 ± 14.6    | 32.5 ± 15.8 | 0.0003   | 27.4 ± 15.4    | 35.0 ± 16.7 | 0.0005*    | 20.7 ± 10.1    | 23.4 ± 6.7  | 0.315     | 0.117         |
|                    | CD195                  | 0.008         | 13.1 ± 11.1    | 17.5 ± 10.3 | 0.002    | 13.4 ± 10.7    | 18.8 ± 14.8 | 0.001*     | 12.1 ± 13.0    | 12.7 ± 10.3 | 0.823     | 0.777         |
| Monocytes          | CD32a                  | 0.006         | 74.0 ± 30.9    | 67.0 ± 35.9 | 0.014    | 75.4 ± 29.8    | 67.8 ± 35.8 | 0.026     | 69.8 ± 35.1    | 64.5 ± 37.8 | 0.338     | 0.657         |
|                    | CD32b                  | 0.050         | 50.8 ± 38.9    | 40.7 ± 36.7 | 0.079    | 53.7 ± 36.9    | 38.1 ± 36.5 | 0.032     | 42.2 ± 47.9    | 48.7 ± 40.5 | 0.148     | 0.642         |

*aSignificance of paired data based on the direction of change from initial screen of IVIg response.

*bSignificance of paired data based on the mean difference in expression before and after treatment.

*cExpression of markers before IVIg treatment were compared (responders vs. nonresponders/relapse) to identify any markers capable of predicting clinical response.

*dCorrection for multiple testing according to Bonferroni was based on the number of markers analyzed in each leukocyte population (two markers per population: p = .025; three markers per population: p = .017; conservative correction for all 13 markers: p = .0038). Bold markers were associated with clinical efficacy and passed correction at the population level; an asterisk identified markers that passed conservative Bonferroni correction.
FIGURE 1  Decreased CD16+ myeloid dendritic cell population after intravenous immunoglobulin (IVIg) was associated with clinical efficacy. (A) Change in CD16+ mDCs during the treatment cycle that defined response from relapse or nonresponse in each patient. (B) Predictability of decreased CD16+ mDCs on clinical outcome was supported by sensitivity but not specificity (Fisher’s exact test). (C) The magnitude of CD16 downregulation on mDCs correlated with increased MRC sum scores, and decreased INCAT disability scores (Spearman’s rank correlation). Decreased CD16 expression was associated with loss of the HLA-DRlow CD16+ mDC population after IVIg in a responder patient (D), but retention of this population after IVIg in a relapsing patient (E). The HLA-DRlowCD11c+ population is outlined by the elliptical gate in the middle panels, and the CD16/HLA-DR gate was set according to this HLA-DRlow population and isotype control for CD16. The pre-IVIg phenotype of this HLA-DRlow CD11c+ population was the same for both responding patient (F) and relapsing patient (G), which defined an activated mature but circulating inflammatory mDC population that declined after clinically effective therapy.
may be a temporary effect of cytokine neutralization after IVIg infusion.

In searching for biomarkers to monitor and predict outcome of IVIg therapy, an ideal biomarker should be sensitive to changes in IVIg dosing and predict clinical outcome soon after treatment, as measured 7 days after infusion in this study. Useful biomarkers should be able to guide optimal IVIg dosing for new CIDP patients, and provide early warning of relapse when performing dose tapering or testing for IVIg dependence when in remission. The six markers associated with clinical efficacy identified in this study may be suitable for monitoring therapeutic response in combination. They signified a reduced inflammatory profile and correlated with clinical outcome, albeit a temporary change for most markers, reverting toward baseline before the next IVIg infusion. A decreased CD16+ mDC population was associated with response to IVIg and demonstrates the feasibility of using this marker in a test to establish optimal IVIg dosing for new CIDP patients, although specificity was insufficient to predict relapse during dose tapering. Additionally, the small number of patients in this study means that these data cannot be used to guide clinical decisions on IVIg dosing. Confirmation of the critical components of these immunomodulatory pathways may facilitate development of novel therapeutics aimed at replacing this expensive and nonspecific anti-inflammatory treatment, and such markers may provide the means to test efficacy.

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CONFLICT OF INTEREST

None declared.

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