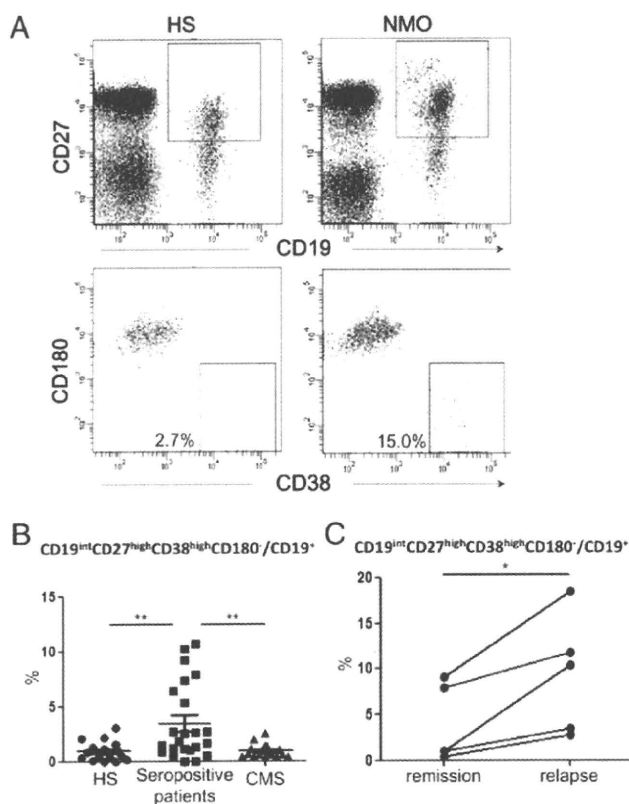


**Results**

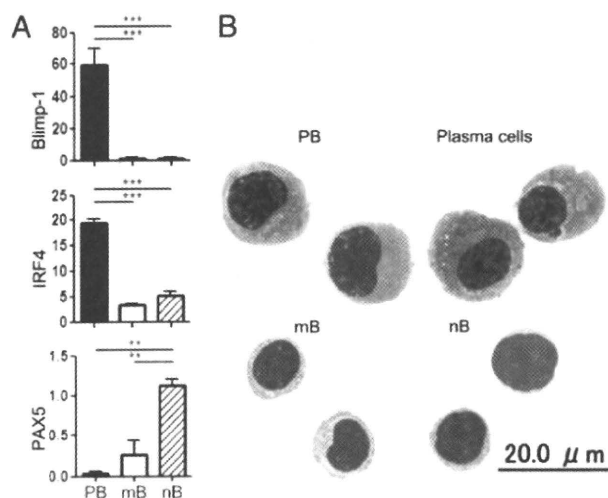
**CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> B Cells Were Increased in the Peripheral Blood of NMO Patients.** Although AQP4-Abs are identified as IgGs (18), no prior study has focused on proportional changes of B-cell subsets in NMO. We therefore performed multicolor flow cytometric analysis of peripheral blood mononuclear cells (PBMC) derived from patients and controls. After starting the study, we soon noticed a remarkable expansion of a distinct B-cell subset in some patients with NMO. The expanded B cells were identified as a population of CD27<sup>high</sup>, CD38<sup>high</sup>, and CD180<sup>-</sup>, and showed lower expression of CD19 than other B cells (Fig. 1A). Notably, this population did not express the B-cell marker CD20 (Fig. S1). First, we collected samples from patients in remission and analyzed the pooled data. We found that the proportion of this subpopulation among CD19<sup>+</sup> B cells was significantly increased in AQP4-Ab seropositive patients with NMO or NMO spectrum disorder (Fig. 1B) compared with HS or CMS patients. There was no significant difference in the proportion of this B-cell subpopulation between those with typical NMO and those with NMO spectrum disorder. Furthermore, the frequency of this B-cell subpopulation was correlated with the serum AQP4-Ab titer (Fig. S2). Comparison of paired samples obtained from the same patients during relapse and in

remission showed that the CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> B cells further increased during relapse (Fig. 1C). In contrast, the frequencies of CD27<sup>-</sup> naïve B cells (nB) and CD27<sup>+</sup>CD38<sup>-low</sup> memory B cells (mB) were not altered in AQP4-Ab seropositive patients compared with controls (Fig. S3). The large majority of seropositive patients were treated with corticosteroids. However, the frequency of CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> cells among CD19<sup>+</sup> B cells was not correlated with the daily corticosteroid dose given to patients (Fig. S4). Moreover, the increase in cells in NMO patients was still evident compared with that in CMS patients similarly treated with corticosteroids (Fig. S5). Taken together, the selective increase in CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> B cells in seropositive patients was thought to reflect their role in the pathogenesis of NMO but not to be an effect of the corticosteroid treatment.

**Expanded Cells Resemble Early Plasma Cells in Gene Expression and Morphology.** To gain insights into the developmental stage of the CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> B cells, we quantified the mRNA expression of B-cell-associated transcription factors in sorted cell populations. Compared with nB and mB, this population showed much higher expression of B-lymphocyte-induced maturation protein 1 (Blimp-1) and IFN regulatory factor 4 (IRF4), which are essential for the regulation of plasma cell differentiation (19, 20) (Fig. 2A). In contrast, the expression of paired box gene 5 (PAX5), known to be down-regulated in early plasma cell differentiation (21), was reciprocally reduced in the B-cell subset. This gene expression pattern is very similar to that of plasma cells. However, it was notable that the cells of interest expressed CD19, which is not detected in mature plasma cells. Moreover, only 40% of this population expressed the most reliable plasma cell marker CD138 (22). Morphological analysis also confirmed the similarity of this population to plasma cells: they exhibit eccentric nucleus, perinuclear hof region, and abundant cytoplasm. However, they possess a larger nucleus with a lower extent of chromatin clumping compared with CD138<sup>+</sup> plasma cells derived from HS (Fig. 2B). Notably, the CD138<sup>+</sup> population among CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> cells in NMO patients was



**Fig. 1.** CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> B cells increased in NMO patients. (A) A flow cytometric scheme for the analysis of B-cell subpopulations. PBMC from HS and NMO in remission were stained with fluorescence-conjugated anti-CD19, -CD27, -CD38, and -CD180 mAbs. CD19<sup>+</sup>CD27<sup>+</sup> cells were partitioned (Upper) and analyzed for expression of CD38 and CD180 (Lower). Values represent the percentages of CD38<sup>high</sup>CD180<sup>-</sup> cells within CD19<sup>+</sup>CD27<sup>+</sup> cells. (B) Analysis of the pooled data derived from patients in clinical remission. This shows the percentages of CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> cells within CD19<sup>+</sup> cells from HS, seropositive patients, and CMS patients (\*\**P* < 0.01; Tukey's post hoc test). (C) Comparison of remission and relapse of NMO. Data obtained from the same patients are connected with lines (\**P* < 0.05; Wilcoxon signed rank test).



**Fig. 2.** Resemblance of CD19<sup>int</sup>CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> cells to plasma cells. (A) mRNA expression of Blimp-1, IRF4, and PAX5. B-cell subpopulations [CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> (PB), CD27<sup>-</sup> naïve (nB), CD27<sup>+</sup>CD38<sup>-low</sup> memory (mB)] were sorted by FACS and total RNA was extracted for qRT-PCR analysis. RNA levels were normalized to ACTB for each sample (\*\**P* < 0.01; \*\*\**P* < 0.001; Tukey's post hoc test). (B) May-Grünwald-Giemsa staining of B-cell subpopulations. PB (Upper Left), mB (Lower Left), and nB (Lower Right) from NMO are presented along with morphologically identified plasma cells (CD19<sup>int</sup>CD27<sup>high</sup>CD38<sup>high</sup>CD138<sup>+</sup>) from HS (Upper Right).

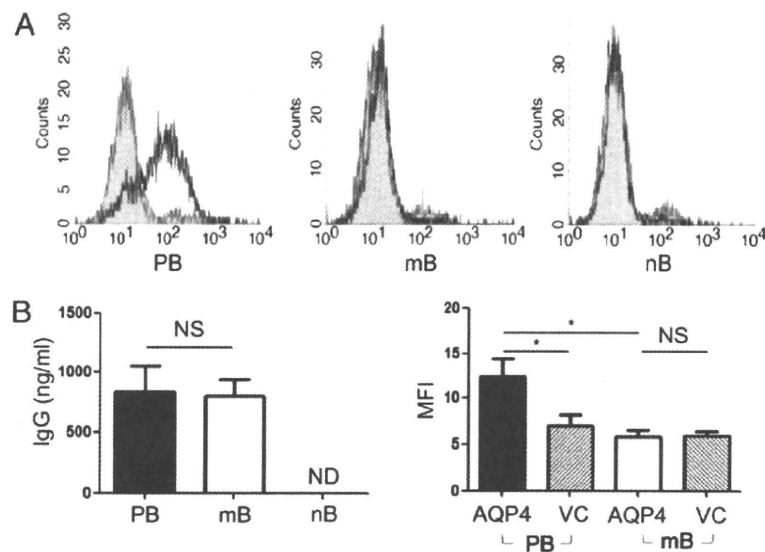
morphologically indistinguishable from the CD138<sup>+</sup> population in NMO patients or HS, indicating the immature characteristic of CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> cells (Fig. S6). These phenotypical and morphological characteristics as well as the results of the quantitative real-time PCR (qRT-PCR) analysis indicate that this B-cell population is equivalent to plasmablasts (PB) (22–26). Hereafter, we use the term “PB” to distinguish this population from other B cells.

**Expression of B-Cell Cytokine Receptors on PB.** Prior studies have identified cytokines that are critically involved in the differentiation and/or survival of plasma cells, including IL-6 and B-cell-activating factor (BAFF). IL-6 induces B-cell differentiation into plasma cells, maintains early plasma cell survival, and enhances plasma cell IgG secretion (24). Besides, IL-6 is elevated in the cerebrospinal fluid (CSF) or peripheral blood of NMO patients compared with that of CMS patients and HS (27, 28). In a rodent autoimmunity model, IL-6 deficiency caused impaired antibody secretion by B cells (29). Given the potential role of IL-6 in NMO, we performed flow cytometry analysis for the expression of IL-6R. Results showed remarkable expression of IL-6R on PB, although it was only marginal or absent on mB and nB (Fig. S7). Because BAFF and A proliferation-inducing ligand (APRIL) can also promote the survival of PB (25, 26), we next evaluated the expression of the receptors for BAFF and APRIL, BAFF receptor (BAFF-R), B-cell maturation antigen (BCMA), and transmembrane activator and calcium modulator and cyclophilin ligand interactor (TACI). Expression of BCMA and TACI was selectively up-regulated in PB in parallel with IL-6R. In contrast, BAFF-R was up-regulated in mB and nB, but not in PB (Fig. S7).

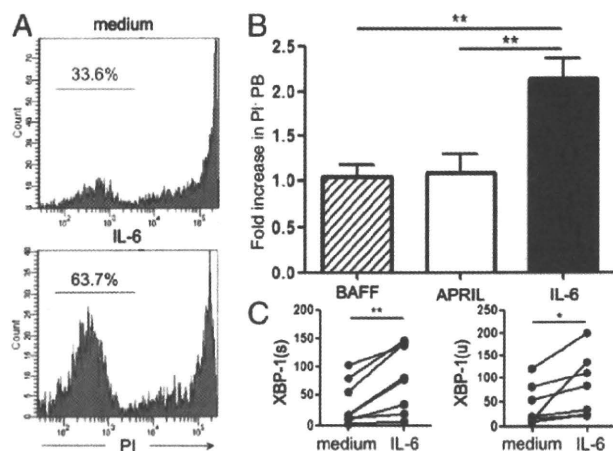
**PB Is a Selective Source of AQP4-Abs in Peripheral Blood.** We were interested to know whether PB were capable of producing AQP4-Abs upon stimulation with cytokines and, therefore, examined the

ability of IL-6, BAFF, and APRIL to enhance AQP4-Ab secretion by PB. We cultured the isolated PB for 6 d in the presence or absence of each cytokine, and evaluated the presence of AQP4-Abs in the supernatants by measuring IgG binding to Chinese hamster ovary (CHO) cells transfected with the human AQP4 vector (CHO<sup>AQP4</sup>) or the vector control (CHO<sup>VC</sup>). We found that IL-6, but not BAFF or APRIL, could significantly enhance AQP4-Ab secretion from PB (Fig. S8), as assessed by specific IgG binding to CHO<sup>AQP4</sup>. Further study focusing on IL-6 showed that exogenous IL-6 promoted the production of AQP4-specific IgGs from PB (Fig. 3A), but not from the other B-cell subpopulations. Similar results were obtained from six independent experiments (Fig. S9), indicating that PB could be major AQP4-Ab producers in PBMC. In the absence of addition of IL-6, supernatants from PB did not show any significant reactivity to CHO<sup>AQP4</sup>. To further analyze the AQP4-Ab-secreting potential of each B-cell subpopulation, we next stimulated the cells with a combination of IL-6, IL-21, and anti-CD40 that efficiently induces B-cell differentiation and IgG production (30). This polyclonal stimulation induced the secretion of similar amounts of IgGs from mB and PB. However, only the supernatant of PB specifically reacted to CHO<sup>AQP4</sup> cell transfectants, indicating that AQP4-Ab-producing B cells were highly enriched in PB (Fig. 3B).

**Survival and Functions of PB Depend on IL-6 Signaling.** We evaluated the influence of IL-6, BAFF, and APRIL on the survival of PB after 2 d of in vitro culture (Fig. 4A). Among the added cytokines, only IL-6 was found to significantly promote the survival of PB (Fig. 4B). We also assessed the expression levels of X-box-binding protein 1 (XBP-1) in PB by qRT-PCR after 24 h of culture with or without IL-6. XBP-1 is a transcription factor critical for IgG secretion (31), and the splicing process of XBP-1 mRNA yields a more active and stable protein. We found that the expression of both unspliced [XBP-1(u)] and spliced [XBP-1(s)] forms of XBP-1 mRNA was augmented in PB by the ad-



**Fig. 3.** Production of AQP4-Abs by PB. (A) Using flow cytometry, we examined whether AQP4-Abs could be produced by PB, mB, or nB cells. FACS-sorted cells were cultured with IL-6 (1 ng/mL) for 6 d and supernatants were collected. Supernatant IgGs reactive to CHO<sup>AQP4</sup> (open histogram) and CHO<sup>VC</sup> cells (closed histogram) were detected by anti-human IgG secondary antibody. The supernatant from PB (Left), but not from mB or nB, contains IgGs reactive to CHO<sup>AQP4</sup>, indicating that only PB secrete AQP4-Abs after stimulation with IL-6. (B) Memory B cells (mB) produce IgG but not AQP4-Abs. B-cell subpopulations were cultured in the presence of IL-6 (1 ng/mL), IL-21 (50 ng/mL), and anti-CD40 mAb (1  $\mu$ g/mL) for 6 d. IgGs in the culture supernatants were measured by sandwich ELISA (Left) (each assay was performed in quadruplicate). Data from three patients are expressed as mean  $\pm$  SD. The activity of AQP4-Abs in the culture supernatants from PB and mB was also measured by flow cytometry (Right). Aliquots of CHO<sup>AQP4</sup> cells (AQP4) and CHO<sup>VC</sup> cells (VC) ( $n = 4$  for each) were stained with the supernatant of PB or mB from every patient. Data are expressed as median fluorescence intensity values from the results of three patients (\* $P < 0.05$ ; Tukey's post hoc test). ND, not detected; NS, not significant.

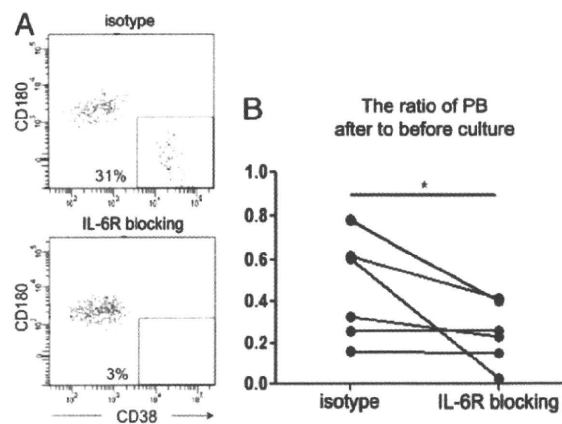


**Fig. 4.** Effect of exogenous IL-6 on PB. (A) IL-6 promotes the survival of PB. FACS-sorted PB were cultured in the presence or absence of recombinant IL-6 (1 ng/mL) for 2 d. PI staining of cultured PB showed that exogenous IL-6 increased the percentage of surviving cells (Lower) compared with cells cultured in the medium alone (Upper). Values shown are percentages of unstained cells. (B) Comparison of IL-6 with BAFF and APRIL. Here we show that only IL-6 could promote cell survival. Data are expressed as fold increase of % PI<sup>-</sup> cells following the addition of each cytokine. At least four independent experiments were performed to obtain the results (\*\* $P < 0.01$ ; Tukey's post hoc test). (C) Effect of IL-6 on XBP-1 expression. FACS-sorted PB were cultured with or without IL-6 for 24 h, and total RNA was extracted from the cells to quantify expression levels of XBP-1(u) and XBP-1(s) by qRT-PCR. Each line connects values obtained from seven independent experiments (\* $P < 0.05$ ; \*\* $P < 0.01$ ; Wilcoxon signed-rank test).

dition of IL-6. These results suggest that IL-6 promoted the survival of PB and enhanced IgG secretion from PB, leading to an increased production of AQP4-Abs in NMO patients (Fig. 4C). In addition, we found that the frequency of PB tended to be increased when serum IL-6 levels were higher than the mean  $\pm$  2 $\times$  SEM of those in HS [PB/PBMC (%) for IL-6 high group  $0.62 \pm 0.47$  (%); PB/PBMC (%) for IL-6 low group  $0.15 \pm 0.05$  (mean  $\pm$  SD)]. These observations prompted us to address whether the blockade of IL-6R signaling could exhibit any influence on PB. We cultured PBMC derived from AQP4-Ab seropositive patients in the presence of 20% autologous serum and examined the effect of adding anti-IL-6R antibody by counting the number of surviving PB. We found that the frequency of PB among total B cells decreased significantly in the presence of anti-IL-6R mAb (Fig. 5A and B). Among six patients examined, the PB reduction was remarkable in three patients, but was only marginal in the other patients. Notably, the former group of patients showed higher IL-6 levels in the serum (4.69, 6.47, and 25.5 pg/mL for each patient), compared with the latter (1.42, 1.43, and 2.91 pg/mL). The frequency of other B-cell subpopulations did not change with the addition of anti-IL-6R mAb. These results led us to postulate that in vivo administration of anti-IL-6R mAb may ameliorate NMO.

## Discussion

A growing body of evidence suggests that AQP4-Abs play a pathogenic role in NMO (6, 7, 10–12). Here we report that a B-cell subpopulation bearing the CD19<sup>int</sup>CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> phenotype is responsible for the selective production of AQP4-Abs. The cells that we call PB are vulnerable to IL-6R blockade by anti-IL-6R mAb, leading us to propose anti-IL-6R mAb as a therapeutic option for NMO. Bennett et al. recently reported that plasma cells in CSF are a potential source of pathogenic AQP4-Abs (10). However, this study has not excluded a possible role of AQP4-Abs produced in the peripheral blood. It has been



**Fig. 5.** IL-6R blockade selectively inhibits the survival of PB. (A) PBMC were cultured in a medium containing 20% autologous serum in the presence of IL-6R-blocking antibody or its isotype control mAb for 2 d. The cells were stained and analyzed as described in the experiment in Fig. 1A. Data represent the percentages of PB within CD19<sup>+</sup>CD27<sup>+</sup> cells. A representative pair of six independent experiments is shown. (B) The percentage of PB within CD19<sup>+</sup> B cells (PB%) was determined for each pair of cultures either with anti-IL-6R mAb (IL-6R-blocking) or isotype control mAb (isotype) before and after starting the culture. Then, the PB survival ratio was calculated for each culture by dividing the PB% at the end of the culture by the PB% obtained before starting the culture. Lines connect the PB survival ratios of six independent experimental pairs to clarify that IL-6R blockade reduces PB survival (\* $P < 0.05$ ; Wilcoxon signed-rank test).

repeatedly shown that the passive transfer of pathogenic autoantibodies, including AQP4-Abs (10–12, 32), augments the formation of inflammatory lesions in EAE. Therefore, once T-cell-mediated inflammation takes place in the CNS, pathogenic autoantibodies produced outside the CNS are able to enter the CNS compartment. It is also notable that AQP4-Abs are more abundant in the peripheral blood of NMO patients than in their CSF (33). Taken together, we speculate that PB that are expanded in the peripheral blood during relapse may play a critical role in the pathogenesis of NMO by producing AQP4-Abs, although more work is necessary to explore whether PB can enter the CNS.

It is generally thought that circulating IgGs are mainly secreted by long-lived plasma cells residing in healthy bone marrow. It remains unclear how PB secreting AQP4-Abs can differentiate and survive in the peripheral circulation. It has been previously shown that autoantibodies producing plasma cells accumulate in peripheral lymphoid organs (34). It would be interesting to investigate which organs blood PB move to during the course of NMO. The levels of IL-6 in the serum and CSF are elevated in NMO compared with HS or CMS patients (27, 28). In this regard, it is of note that blocking IL-6R signaling was found to dramatically reduce the survival of PB ex vivo, which was dependent on the presence of autologous serum containing IL-6. These results suggest that the increase of PB in AQP4-Ab seropositive patients could be attributed to the increased IL-6 in the serum. We also demonstrated that improved PB survival in the presence of exogenous IL-6 was accompanied by up-regulated expression of XBP-1. It is noteworthy that wild-type and XBP-1<sup>-/-</sup> B cells start to produce more IL-6 after forced overexpression of XBP-1(s), which results in the operation of a positive feedback loop controlling IgG secretion (31). Treatment with anti-IL-6R is promising because IL-6R blockade could terminate this vicious loop that controls the production of autoantibodies.

It has been reported that NMO patients have higher levels of BAFF in the serum or CSF compared with CMS patients (35). BAFF is also known to support plasma cell differentiation and survival of PB induced in vitro (25). However, in our ex vivo

study, BAFF did not promote the survival of PB, indicating that PB were not a target for BAFF. We speculate that BAFF might specifically act on an early process of plasma cell differentiation and does not have an influence on cells like PB that have entered a later stage.

IL-6R blockade by humanized mAb against IL-6R (tocilizumab) has proven to be useful for treating immune-mediated diseases, including rheumatoid arthritis (36) and Castleman's disease (37). Here we propose that IL-6R-blocking antibody treatment should be considered as a therapeutic option for NMO. Currently, most NMO patients are being treated with corticosteroids in combination with immunosuppressive drugs and plasma exchange (38). Anti-CD20 mAb, which causes B-cell depletion, has also been used for serious cases of NMO. Because the level of B-cell depletion appears to correlate with the suppressive effects of anti-CD20 in NMO (39), it has been argued that B cells are essential for the pathogenesis of NMO, either via acting as antigen-presenting cells or as autoantibody producers. Weber et al. recently reported that activated antigen-specific B cells serve as antigen-presenting cells and polarize proinflammatory T cells in EAE (40), supporting the view that the therapeutic effects of anti-CD20 might be attributable to the depletion of antigen-presenting B cells. Notably, they also cautioned that elimination of CD20<sup>+</sup> cells might deplete nonactivated cells as well as regulatory B cells possessing anti-inflammatory potentials. Although the effect of anti-CD20 on AQP4-Ab-secreting cells has not been reported, it is likely that the majority of PB are not affected because they do not express CD20. Consistent with our prediction, anti-CD20 treatment was not effective in aggressive cases of NMO (41, 42). It appears that selective depletion of activated antigen-specific B cells could be a more promising strategy to improve the efficacy of B-cell-targeted therapies for NMO. In this regard, PB-targeting therapy is a promising approach. Given the efficacy of IL-6R blockade in reducing the number of PB *ex vivo*, we find it very interesting to explore the effect of anti-IL-6R mAb on NMO.

## Materials and Methods

**Patients and Controls.** A cohort of 24 AQP4-Ab seropositive patients was recruited at the Multiple Sclerosis Clinic of the National Center of Neurology and Psychiatry (NCNP). Among these, 16 met the revised NMO diagnostic criteria (3). The other 8 were diagnosed with NMO spectrum disorder (1) because they did not develop both myelitis and optic neuritis (optic neuritis alone in 6 cases; myelitis alone in 2 cases). Seventeen age- and sex-matched CMS patients and 20 HS were enrolled as controls. Serum AQP4-Ab levels were measured by a previously reported protocol by courtesy of Kazuo Fujihara at Tohoku University (Sendai, Japan) (33). All CMS patients had relapsing-remitting MS and fulfilled McDonald diagnostic criteria (43).

At the time of blood sampling, 21 seropositive patients were receiving corticosteroids (prednisolone 5–25 mg/d). Seven of these patients were also being treated with azathioprine (12.5–100 mg/d) or tacrolimus (3 mg/d). Six CMS patients were receiving low-dose corticosteroids without immunosuppressants. None of the seropositive or CMS patients had received IFN- $\beta$ , *i.v.* corticosteroids, plasma exchange, or *i.v.* immunoglobulins for at least 1 mo

before blood sampling. Blood sampling during relapse was performed in six seropositive NMO patients before they received intensive therapy starting with *i.v.* corticosteroids. Five of these patients were followed up further and blood was collected again after they entered remission. Anti-nuclear and/or anti-SS-A Abs were detected in some of the seropositive patients, but none met the diagnostic criteria for SLE or Sjögren syndrome. Demographic features of the patients are presented in Table 1. The study was approved by the Ethics Committee of the NCNP.

**Reagents.** The following Abs were used in this study: mAbs against CD38, CD19, CD27, CD20, and PE-streptavidin (Beckman Coulter); mAbs against CD180 and BAFF-R (BD Biosciences); mAbs against IL-6R and TACI as well as Abs against BCMA and CD40 (R&D Systems); rabbit anti-human AQP4 antibody (Santa Cruz Biotechnology); FITC-anti-rabbit IgG (Jackson ImmunoResearch Laboratories); and FITC-anti-human IgG antibody (MP Biomedicals). Recombinant proteins of BAFF (ProSpec), APRIL (Abnova), IL-6 (PeproTech), and IL-21 (Invitrogen) were purchased. Propidium iodide (PI) was obtained from Sigma-Aldrich. RPMI 1640 supplemented with 10% FBS, 2 mM L-glutamine, 100 U/mL penicillin, and 100 mg/mL streptomycin (Life Technologies) was used for cell culture.

**Flow Cytometry, Cytology, and Cell Culture.** PBMC were separated using density centrifugation on Ficoll-Paque PLUS (GE Healthcare Biosciences). B cells were analyzed and sorted by FACSaria (BD Biosciences). Each B-cell subset was stained with May-Grünwald-Giemsa. To evaluate AQP4-Ab production *in vitro*, each B-cell subset ( $1$  or  $2 \times 10^4$ ) was cultured for 6 d in the medium alone, in the presence of IL-6 (1 ng/mL) or in the presence of IL-6 (1 ng/mL), IL-21 (50 ng/mL), and anti-CD40 (1  $\mu$ g/mL). Culture supernatants were harvested and analyzed for AQP4-Ab production as described below. To examine the effect of cytokines on the survival of PB, the cells ( $4 \times 10^3$ ) were cultured in the medium alone or in the presence of BAFF (100 ng/mL), APRIL (300 ng/mL), or IL-6 (1 ng/mL) in 96-well U-bottom plates for 2 d and stained with PI to assess cell survival. In parallel, the cells were cultured for 1 d and harvested to evaluate mRNA expression by qRT-PCR. To assess the effect of IL-6 signaling blockade, PBMC ( $5 \times 10^5$ ) were preincubated with anti-IL-6R Abs (1  $\mu$ g/mL) at 4 °C for 20 min, cultured in AIM-V medium (Invitrogen) containing 20% of heat-inactivated serum obtained from each patient in 96-well flat-bottom plates for 2 d, and analyzed by flow cytometry.

**Quantitative RT-PCR Analysis.** mRNA from each cell subset was isolated according to the manufacturer's instructions using an RNeasy Kit (Qiagen). RNA was further treated with DNase using the RNase-Free DNase Set (Qiagen) and reverse-transcribed to cDNA using a cDNA synthesis kit (Takara Bio). PCR was performed using iQ SYBR Green Supermix (Takara Bio) on a LightCycler (Roche). RNA levels were normalized to endogenous  $\beta$ -actin (ACTB) for each sample. Primers used are listed in Table S1.

**Measurement of Ig Isotypes and Serum IL-6.** Secreted IgG in the culture supernatant was quantitated by sandwich ELISA using affinity-purified goat anti-human IgG-Fc (Bethyl Laboratories). Bound IgG was measured according to the manufacturer's instructions. Serum IL-6 was measured by ELISA (R&D Systems) according to the manufacturer's instructions.

**AQP4-Ab Detection Assay.** Human AQP4-expressing cells were established to detect AQP4-Abs by flow cytometry. A human AQP4 (hAQP4) M23 splice variant from a clone collection (Invitrogen) was amplified by PCR and subcloned into a pIRES-DsRed-Express vector (Clontech). CHO cells (American Type Culture Collection) were transfected with this hAQP4 M23 vector (CHO<sup>AQP4</sup>) or vector

Table 1. Demographic features

	HS	Seropositive patients	CMS patients
Number	20	24	17
Age	44.7 $\pm$ 2.8	47.9 $\pm$ 3.2	41.3 $\pm$ 3.0
Male:female	5:15	1:23	5:12
Disease duration		12.0 $\pm$ 1.6	9.4 $\pm$ 2.4
Age of symptom onset		36.1 $\pm$ 3.0	31.9 $\pm$ 3.4
Relapses in last 2 y		1.4 $\pm$ 0.3	0.7 $\pm$ 0.2
EDSS score in disease remission		5.0 $\pm$ 0.5	2.1 $\pm$ 0.6
Other autoantibodies		ANA 13, SS-A 5	ND

Demographic features for HS, AQP4-Ab seropositive patients, and CMS patients. Values are expressed as number or mean  $\pm$  SEM. ANA, anti-nuclear antibody; ND, not detected; SS-A, anti-SS-A antibody; EDSS, expanded disability status scale.

control (CHO<sup>VC</sup>) using FuGENE 6 Transfection Reagent (Roche). After 2 wk of geneticin (Invitrogen) selection, stable clones were established by single-cell sorting. The expression of hAQP4 in the established clones was confirmed using anti-human AQP4 antibody and FITC-anti-rabbit IgG antibody. Reactivity of AQP4-Abs to CHO<sup>AQP4</sup> was confirmed using seropositive NMO patients' sera diluted at 1:1,000 and FITC-anti-human IgG antibody. To measure the AQP4-Ab activity in culture supernatants, these supernatants were concentrated up to 10 times using an Amicon Ultra 0.5 mL 100K device (Millipore), and 10  $\mu$ L of the solution was added to  $3 \times 10^4$  CHO<sup>AQP4</sup> and CHO<sup>VC</sup> cells. After incubation on ice for 20 min, cells were washed with sterile PBS containing 1% BSA and stained with FITC-anti-human IgG antibody. After a 10-min incubation on ice, the cells were washed and fixed for 15 min in 2% paraformaldehyde. Then the cells were washed and analyzed by flow cytometry.

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