Characterization and quantification of alveolar monocyte-like cells in human chronic inflammatory lung disease

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ABSTRACT: This flow cytometric study was designed to identify, characterize and quantify alveolar monocyte-like cells in healthy volunteers and in patients with chronic inflammatory lung disease.

Cells were obtained by bronchoalveolar lavage (BAL) from 19 patients with sarcoidosis, 29 with idiopathic pulmonary fibrosis, 10 with extrinsic allergic alveolitis, 19 with collagen vascular disease, and from 10 healthy volunteers. By taking advantage of the distinct electro-optical features of alveolar macrophages (AMs) and monocyte-like cells, the numbers of alveolar monocyte-like cells were counted, the cell dimensions calculated, and the densities of antigens on the surface of alveolar monocyte-like cells and AMs were compared. By using a panel of monoclonal antibodies detecting CD11b, CD14, CD16, and human leucocyte antigen-DR (HLA-DR), the immunophenotypes of these cells were selectively characterized.

In the BAL fluid of patients with chronic inflammatory lung disease, significantly increased numbers of alveolar monocyte-like cells were detected that exhibited an immunophenotype intermediate between blood monocytes and mature AMs. Positive correlations were found between numbers of monocyte-like cells and expression of the monocyte-associated surface antigens CD11b and CD14 on total AMs; in contrast, an inverse relationship existed between monocyte numbers and expression of the macrophage-associated surface antigens CD16 and HLA-DR. When the patients were assigned to two groups according to the percentage of BAL monocyte-like cells being lower or higher than 13% (=mean value of the controls +2sd), it could be demonstrated that a high percentage of BAL monocyte-like cells was associated with significantly reduced lung function parameters.

In summary, our flow cytometric data strongly support the view that considerable numbers of blood monocytes are recruited to the bronchoalveolar space in patients with chronic inflammatory lung disease.

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Chronic inflammatory lung diseases are characterized by the presence of an alveolitis that causes a derangement of alveolar structures and ultimately leads to loss of functional gas units. This progressive respiratory failure is thought to result from a disease process that is characterized by the infiltration of inflammatory cells, the increased proliferation of interstitial fibroblasts, and the excessive accumulation of extracellular matrix compounds [1]. Studies on the molecular events leading to pulmonary fibrosis have generally shown that monocytes/macrophages contribute significantly to the disease process [2]. Alveolar macrophages (AMs) are the predominant population of free lung cells in the healthy lung. They represent a heterogeneous group of cells with respect to morphology, immunophenotype and function Under noninflammatory conditions, AMs are thought to be maintained both by local proliferation of resident pulmonary macrophages and extravasation of circulating, bone marrow-derived monocytes [6, 7]. In

lung inflammation, however, it has been suggested that increased numbers of monocytes extravasate from the pulmonary microvasculature to the bronchoalveolar space [8, 9].

The alveolitis in interstitial lung diseases appears to be maintained by the local production of specific cell-derived chemotactic factors that recruit inflammatory cells from the pulmonary micovasculature into the alveolar interstitium [10]. An array of cytokines and chemokines has been identified that exhibit chemotactic activity for monocytes, such as monocyte chemoattractant protein-1 (MCP-1). Recently, MCP-1 was found to be intensely expressed in lung tissue [11] and bronchoalveolar lavage (BAL) fluid of patients with interstitial lung disease [12]. Several studies described a monocyte-like phenotype of AMs in chronic inflammatory lung diseases, possibly reflecting the influx of blood monocytes and/or the emergence of a proinflammatory macrophage phenotype [13–17].

In this study, we used a flow cytometric approach to identify monocytic cells in the BAL fluid of control subjects and of patients with chronic inflammatory lung disease. By using a panel of monoclonal antibodies directed against the β_2 -integrin CD11b, the lipopolysaccharide (LPS) receptor CD14, the low-affinity receptor for immunoglobulin G (IgG) (Fc γ RIII, CD16), and the major histocompatability complex (MHC) class II molecule human leucocyte antigen-DR (HLA-DR), we selectively characterized the immunophenotype of these cells. Based on the distinct electro-optical features of AMs and monocytes, we quantified the numbers of alveolar monocyte-like cells and correlated the results with lung function parameters.

Methods

Study population

A total of 77 nonsmoking patients with chronic inflammatory lung disease were studied. BAL fluid was obtained from 19 patients with sarcoidosis (7 males and 12 females; mean age 43±3 yrs), 29 patients with idiopathic pulmonary fibrosis (IPF) (13 males and 16 females; mean age 53±2 yrs), 10 patients suffering from extrinsic allergic alveolitis (EAA) (7 males and 3 females; mean age 43±3 yrs), and 19 patients (9 males and 10 females; mean age 51±2 yrs) with pulmonary involvement by collagen vascular diseases (CVD). The diagnosis of sarcoidosis,

IPF, EAA and CVD was based upon clinical and radiological findings, as well as on pulmonary function tests and positive findings in histological examinations of transbronchial biopsy specimens. A total of 16 patients (three with sarcoidosis, eleven with IPF, one with EAA, and one with CVD) were receiving maintenance prednisone therapy (7.5–15 mg·day-1) for a period of at least 4 weeks prior to BAL.

The control group consisted of 10 nonsmoking volunteers (10 males; mean age 26±1 yrs) without pulmonary disease, who underwent fibreoptic bronchoscopy and BAL for the purpose of this study. Results of lung function tests (table 1) and differential cell counts in the BAL fluid (table 2) were within the normal range in all control subjects. They were not under therapy at the time of the study.

The study protocol was approved by the institutional Ethical Committee. Informed consent was obtained from all control subjects and patients.

Lung function tests

Vital capacity (VC) was determined using a standard spirometer (Jaeger, Würzburg, Germany). The carbon monoxide transfer factor of the lung (TL,CO) was measured by a single-breath method using a gas mixture containing 0.15% CO and 4% helium. Blood gas analyses, including arterial oxygen tension (Pa,O₂), arterial carbon dioxide tension (Pa,CO₂), and pH, were performed in arterialized capillary blood from the ear, at rest and during

Table 1. - Lung function characteristics of control subjects and patients

	Control subjects (n=10)	Patients with sarcoidosis (n=19)	Patients with IPF (n=29)	Patients with EAA (n=10)	Patients with CVD (n=19)
VC % pred	109±2	107±5	79±5*	91±9*	73±5*
TL,CO % pred	109±5	96±6	62±4*	73±4*	69±5*
Resting P_{a,O_2} kPa	11.8±0.2	11.1±0.3	9.2±0.2*	10.5±0.5	10.5±0.2*
mmHg	88.7±1.7	83.6±2.0	69.0±1.7*	79.1±3.9	79.1±1.3*
Exercise Pa,O2 kPa	ND	10.8±0.4	8.6±0.4	9.1±0.4	10.0±0.5
mmHg	ND	80.8±2.7	64.5±3.3	68.0±2.9	75.1±3.6

Data are presented as mean \pm sem. IPF: idiopathic pulmonary fibrosis; EAA: extrinsic allergic alveolitis; CVD: collagen vascular disease; VC: vital capacity; $T_{L,CO}$: transfer factor of the lung for carbon monoxide; % pred: percentage of predicted value; P_{a,O_2} : arterial oxygen tension; ND: not determined. *: p<0.05 *versus* control value.

Table 2. - BAL fluid cellular constituents in control subjects and patients

		Control subjects (n=10)	Patients with sarcoidosis (n=19)	Patients with IPF (n=29)	Patients with EAA (n=10)	Patients with CVD (n=19)
Total cells ×	106	6.5±1.5	16.0±3.6*	15.5±2.2*	33.7±13.3*	10.5±1.2*
Cells·mL-1 BA	AL fluid ×10 ⁴	12.1±2.3	33.5±6.5*	36.1±4.9*	58.5±19.5*	21.3±2.0*
Macrophages	%	93±2	80±4*	70±4*	50±5*	85±3*
1 0	cells·mL-1 ×104	11.3±2.8	27.6±6.3*	22.5±2.8*	27.4±8.6*	17.8±1.7*
Lymphocyes	%	5±1	19±4*	21±4*	25±5*	6±1
3 1 3	cells·mL-1 ×104	0.5 ± 0.1	5.5±1.4*	10.2±3.7*	15.9±6.2*	1.2±0.2*
Granulocytes	%	2±1	2±0	8±1	24±6*	10±2*
,	cells·mL-1 ×104	0.2 ± 0.1	0.4 ± 0.1	2.9±0.7*	14.9±6.8*	2.2±0.5*

Data are presented as mean±sem. BAL: bronchoalveolar lavage. For further definitions see legend to table 1. *: p<0.05 versus control value.

steady-state exercise. Lung volumes were compared with the European Coal and Steel Community normal values [18]. The CO transfer factor was expressed as a percentage of our laboratory reference values [19].

Bronchoalveolar lavage

Fibreoptic bronchoscopy with BAL was performed as described previously [20]. A flexible fibreoptic bronchoscope was wedged into a subsegmental bronchus of the middle lobe or lingula, and five 20 mL aliquots of sterile saline solution (0.9% NaCl) were infused and immediately aspirated. The recovered BAL fluid was pooled, filtered through two layers of sterile cotton gauze, and centrifuged at 300 ×g for 10 min. The cell pellet was washed twice, resuspended in phosphate-buffered saline (PBS), and the total cell counts were measured using a standard haemacytometer (Coulter Electronics, Krefeld, Germany). Cell viability was assessed by the trypan blue exclusion technique. Cytocentrifuge smears (500 rpm ×5 min) served to identify the cellular populations after staining with May-Grünwald-Giemsa. A total of 300 cells were counted to determine the percentages of alveolar macrophages, lymphocytes, neutrophils, eosinophils and mast cells.

Processing of BAL cells for flow cytometry

Surface antigen expression was determined by direct immunofluorescence applying the following panel of fluorescein isothiocyanate (FITC) or phycoerythrin (PE) conjugated monoclonal antibodies (Becton Dickinson, Heidelberg, Germany): anti-CD11b (Leu-15), anti-CD14 (Leu-M3), anti-CD16 (Leu-11c), and anti-HLA-DR. To estimate the amount of nonspecific binding of mouse monoclonal antibodies to human cell surface antigens, in particular that caused by Fc receptors, the cells were also incubated with irrelevant mouse monoclonal antibodies of IgG_1 or $IgG_{2\alpha}$ isotype. For removal of contaminating erythrocytes and fixation of cells, a commercially available lysing solution (FACS Lysing Solution, Becton Dickinson) was used. Incubation with monoclonal antibodies and flow cytometry was performed as described previously [21, 22]. Briefly, cell density was adjusted to 1×10⁶ cells·mL⁻¹, and 100 μL of the suspension was incubated with saturating amounts of conjugated monoclonal antibodies for 30 min at 4°C, in the dark. Cells were fixed and contaminating erythrocytes were removed by addition of 1 mL of the lysing solution for 10 min. The samples were centrifuged, washed and resuspended in PBS before flow cytometric analysis was performed.

Flow cytometry

All data were obtained using a fluorescence-activated cell sorter (FACS) Analyzer flow cytometer (Becton Dickinson, Heidelberg, Germany) equipped with a HgCd arc lamp, and a nozzle orifice of 75 µm. FITC and PE fluorescence signals were collected after excitation at 485±10 nm using standard band-pass filters (530/30 and 595/60). In general, a minimum of 10⁴ events was

collected for each sample, and data were recorded in list mode on a Consort 30 data handling system (Becton Dickinson) and stored for later analysis. To ensure the comparability of the flow cytometric data over a prolonged period of time, the instrument setting of the flow cytometer was calibrated daily by using standard fluorescent microspheres.

The flow cytometer used in this study generates volume signals that are based on the Coulter-type measurement of electrical resistance. These signals are directly proportional to the volume of a particle/cell (as well as to its electrical resistance), in contrast to the forward scatter signal of other flow cytometers that is proportional to the cross-sectional area of a particle/cell (as well as to its refractive signal). This enabled us to calculate morphometric data, such as volume, diameter and surface area. To convert the arbitrary units of the Coultertype volume signals into geometric units, a calibration curve was generated by plotting the known diameters of a set of microsphere samples of five different particle sizes (Flow Cytometry Standards Corp., Research Triangle Park, NC, USA) against the corresponding volume signals.

The AM subpopulations were selectively analysed by electronic gating of volume side-scatter dot plots. Fluore-scence histograms were used to calculate mean fluorescence intensities and to assess the percentages of positively stained cells. To overcome the problem of autofluorescence, the relative fluorescence intensity of a given sample was calculated by subtracting the signal obtained from cells incubated with the corresponding isotype control from the signal generated by cells that were incubated with the test antibody.

Statistics

Data were expressed as arithmetic means±standard error of the mean (sem), except for the normal range of the percentage of BAL monocyte-like cells that was defined as the mean percentage of the control value +2sd. Multiple group comparisons were performed using Kruskal-Wallis analysis of variance (ANOVA) and Mann-Whitney rank sum test corrected by the Bonferroni method. Correlation coefficients were determined by Pearson's test. For differences between groups and correlation coefficients between variables, a two-tailed p-value of less than 0.05 was considered significant. The statistical computations were performed with the aid of the statistical package for the social sciences (SPSS)/PC+ statistical software.

Results

Functional characteristics of control subjects and patients with interstitial lung disease

The results of lung function tests in the control, sarcoidosis, IPF, EAA, and CVD groups are shown in table 1. There was a significant reduction of VC and TL,CO in patients with IPF, EAA and CVD, when compared with the control group. Patients with sarcoidosis did not differ significantly from the control group. P_{a,O_2} values at rest were significantly reduced in patients with IPF and CVD.

Total and differential cell counts from BAL fluid

BAL fluid recovery and BAL cell viability were within the normal range and comparable between the five groups

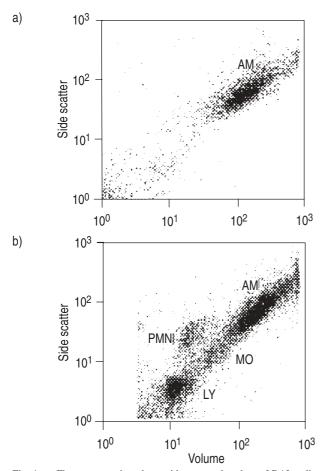


Fig. 1. – Flow cytometric volume side-scatter dot plots of BAL cells characteristic for: a) healthy control subject; and b) a patient with idiopathic pulmonary fibrosis. AM: alveolar macrophage; MO: monocytelike cells; PMN: granulocytes; LY: lymphocytes; BAL: bronchoalveolar lavage.

(data not shown). The cell counts in the various disease groups and healthy volunteers are summarized in table 2. The total number of BAL cells recovered was significantly increased in all disease groups as compared with healthy volunteers. The percentages of lymphocytes were elevated in the sarcoidosis, IPF and EAA groups, whilst the percentages of granulocytes were higher in the EAA and CVD groups. In all disease groups, the percentage of AMs was significantly lower than in the control group. However, because of the increased total BAL cell numbers, the absolute number of AMs was increased in all disease groups. Mast cells were found only in small numbers that did not differ significantly between the study groups.

Flow cytometric data

Figure 1 shows two characteristic volume side-scatter dot plots derived from flow cytometric analyses of BAL cells. Figure 1a demonstrates the normal picture of BAL cells from a healthy control subject featuring a homogeneous population of predominantly large cells with considerable granularity (as defined by light-scattering properties), corresponding to the normal "large" AM population. Figure 1b, in contrast, shows the characteristic finding in a patient with chronic inflammatory lung disease. In addition to increased lymphocyte and granulocyte numbers, two distinct subpopulations of macrophages can be observed, i.e. a subpopulation of normal, "large" AM, along with a well-defined subpopulation of smaller monocyte-like cells. The electro-optical characteristics of the latter proved to be almost identical to those of blood monocytes (data not shown).

To further analyse these cell populations, firstly a gate that included the entire AM population ("large" AM and monocyte-like cells) was applied. As shown in table 3, a tendency to lower expression of HLA-DR and CD16, and to increased expression of CD11b and CD14 on the surface of AMs was observed in most of the disease groups compared to the control group. Due to the large number of group comparisons and also because of high interindividual variations, however, statistical significance (p<0.05) was only reached for the increased CD11b intensity and the elevated percentage of CD14 positive AM in EAA patients, for the decreased percentage of CD16 positive cells in IPF and CVD patients, for the reduced

Table 3. - Surface antigen expression on the total alveolar macrophage population

		Control subjects (n=10)	Patients with sarcoidosis (n=19)	Patients with IPF (n=29)	Patients with EAA (n=10)	Patients with CVD (n=19)
CD11b	%	52±5	72±9	70±5	76±5	48±4
	FI	117±14	201±32	219±31	267±26*	128±19
CD14	%	23±3	32±6	37±4	48±8*	24±3
	FI	56±5	69±8	97±18	101±15	62±7
CD16	%	83±3	63±7	41±5*	55±9	50±7*
	FI	267±26	154±25*	107±18*	173±40	112±16*
HLA-DR	%	92±1	89±3	84±2	70±8*	80±5
	FI	587±17	552±42	455±27	380±53	434±49

Data are presented as mean±sem. %: percentage of positively stained cells; FI: relative fluorescence intensity; HLA-DR: human leucocyte antigen-DR. For further definitions see legend to table 1. *: p<0.05 versus control value.

Table 4. – Comparison of morphometric and immunophenotypic features of alveolar monocyte-like cells and "large" AMs

		Monocyte- like cells (n=87)	"Large" AMs (n=87)	p-value
Diameter	•	12.3±0.1	19.5±0.3	<0.001
Surface ar		476±9	1213±31	<0.001
CD11b	%	76±2	66±4	<0.02
	FI	118±11	184±17	<0.001
	FD	0.23±0.02	0.14±0.01	<0.001
CD14	%	64±2	34±3	<0.001
	FI	78±6	79±8	>0.05
	FD	0.17±0.01	0.07±0.01	<0.001
CD16	%	63±2	65±3	>0.05
	FI	43±5	174±14	<0.001
	FD	0.09±0.01	0.15±0.01	<0.001
HLA-DR	%	86±1	92±2	<0.001
	FI	190±11	550±21	<0.001
	FD	0.41±0.02	0.49±0.04	<0.02

Data are presented as mean±sem. %: percentage of positively stained cells; AMs: alveolar macrophages; FI: relative fluorescence intensity (mean channel fluorescence); FD: fluorescence density (relative FI·μm-² cell surface); HLA-DR: human leucocyte antigen-DR.

CD16 staining intensity in patients with sarcoidosis, IPF and CVD, and for the decreased expression of HLA-DR molecules on AM from patients with EAA.

Secondly, the two AM populations ("large" AM and monocyte-like cells) were selectively gated and their morphometric and immunophenotypic features were compared (table 4). Monocyte-like cells and "large" AM had mean diameters of 12.3±0.1 and 19.5±0.3 μm, respectively, and a mean surface area of 476±9 and 1213±31 μm², respectively. Whilst expression of HLA-DR was found to be significantly higher on "large" AMs, expression of CD14 and CD11b proved to be significantly higher on monocyte-like cells. No significant difference was found for the percentage of CD16 positive cells. When comparing the relative fluorescence intensities as a measure of the overall number of binding sites on the cell surface, "large" AMs expressed a threefold higher number of HLA-DR antigens, a fourfold higher number of CD16 molecules, and 1.5 fold more CD11b surface epitopes, as compared to monocyte-like cells. However, no statistical difference was found for the fluorescence intensity of CD14-labelled cells (table 4).

To evaluate whether the observed differences in the expression of surface molecules were simply due to differences in the size of the surface area of the two macrophage populations, the relative fluorescence density of surface molecules, *i.e.* the relative fluorescence intensity (FI)·μm-² cell surface, was calculated. These calculations revealed that HLA-DR (1.2 fold) and CD16 (1.7 fold) molecules were expressed with a higher density on the surface of "large" AMs, while the density of CD11b (1.6 fold) and CD14 (2.4 fold) molecules was higher on the surface of monocyte-like cells.

Next, the percentage and the total numbers of BAL monocyte-like cells·mL-1 BAL fluid were assessed by counting the numbers of cells included in the two macrophage gates, calculating the percentage of monocyte-like

Table 5. — Correlations of numbers of alveolar monocyte-like cells with lung function parameters and expression of surface antigens on total AMs

Parameter	Correlation coefficient	p-value
TL,CO % pred	-0.42	< 0.001
Resting P_{a,O_2} mmHg	-0.43	< 0.001
CD11b+ AM %	0.42	< 0.01
CD14+ AM %	0.63	< 0.001
CD16+ AM %	-0.49	< 0.001
HLA-DR+ AM %	-0.65	< 0.001

For definitions see legends to tables 1 and 3.

cells within the entire AM population, and transferring these data to the conventional BAL differential and total cell counts. According to these calculations, the percentage of BAL monocyte-like cells was 9±1% in control subjects, 13±2% in patients with sarcoidosis, 16±1% in patients with IPF (p<0.05 versus control subjects), 15± 3% in patients with EAA, and 16±2% in patients with CVD (p<0.05 versus control subjects). The total number of monocyte-like cells·mL-1 BAL fluid was significantly (p<0.05) increased in all disease groups, with 3.5 ± 0.6 ×10⁴ cells⋅mL⁻¹ in patients with sarcoidosis, 5.4±0.6 ×10⁴ cells·mL⁻¹ in patients with IPF, 8.3±2.5 ×10⁴ cells·mL⁻¹ in patients with EAA, and 3.5±0.6 ×10⁴ cells·mL⁻¹ in patients with CVD, as compared to only $1.0\pm0.1 \times 10^4$ cells·mL-1 in control subjects. Whilst negative correlations were found between numbers of monocyte-like cells and expression of the macrophage-associated surface receptors CD16 and HLA-DR on the total AM population, positive correlations were calculated between monocyte numbers and total expression of the monocyteassociated surface antigens CD11b and CD14 (table 5). Most interestingly, a close relationship was also detected between monocyte numbers and lung function parameters (table 5).

Finally, all patients, irrespective of their diagnosis, were assigned to two groups according to whether the calculated percentage of BAL fluid monocyte-like cells was lower or higher than 13% (=mean value of the control group +2sD). As shown in table 6, high percentage of these cells in the BAL fluid (>13%) was associated with reduced VC, and impaired gas exchange, as assessed by measurements of TL,CO, resting Pa,O2, and exercise Pa,O2.

Table 6. – Comparison of lung function findings in patients with high or low percentage of alveolar monocyte-like cells

		Group A <13 % monocytes in BALF	Group B >13 % monocytes in BALF	p-value
VC % pred		96±4	80±4	< 0.05
TL,CO % pred		84±4	64±3	< 0.005
Resting P_{a,O_2}	kPa	11.0 ± 0.2	9.6 ± 0.2	< 0.001
C . 2	mmHg	82.7±1.2	71.8±1.7	
Exercise Pa,O ₂	kPa	10.2 ± 0.4	9.1 ± 0.3	< 0.01
-	mmHg	76.8±2.7	67.9±2.6	

Data are presented as mean±sem. BALF: bronchoalveolar lavage fluid. For further definitions see legend to table 1.

Discussion

Alveolar macrophages have been described as a heterogeneous population with respect to morphology, immunophenotype and function. Subpopulations of AMs were identified in the lungs of normal rats [23-25], guineapigs [26], rabbits [27], and humans [3, 5, 28, 29]. Several experimental studies have suggested that resident AM populations in the normal lung are maintained by local division of cells and that the influx of peripheral monocytes is not needed to sustain the AM population [6, 30, 31]. In contrast, Bowden and Adamson [7] postulated that under normal circumstances the predominant mechanism of macrophage production is direct passage of monocytes across the interstitium to the alveoli, and that only a smaller proportion of cells appears to arise after division of resident interstitial cells with subsequent migration to the alveoli.

Kinetic studies after carbon loading in mice have suggested an acceleration of the normal dual mechanism, whereby most cells are derived from monocytes crossing the interstitium without division and a smaller proportion arising by division of interstitial cells with migration to the alveoli [8]. Further animal studies have demonstrated that the increase of the pulmonary macrophage population during an acute inflammation is brought about mainly by monocyte influx and to a minor extent by temporarily increased local production of macrophages [9]. Consequently, the appearance of small monocyte-like AMs was described in further experimental studies both of acute [32] and chronic [22] lung inflammation. However, local macrophage replication may also play a role in the expansion of the AM population in chronic lung inflammation [33, 34].

There is a growing body of evidence that an increased recruitment of monocytes also takes place in human interstitial and other lung diseases. An elevated expression of monocyte-lineage surface antigens was found on AMs from patients with sarcoidosis [13, 14, 17, 35–38], idiopathic pulmonary fibrosis [15, 16, 36–38], extrinsic allergic alveolitis [36, 37], bronchogenic carcinoma [35], asthma [39], human immunodeficiency virus (HIV) infection [40], as well as in smoking individuals [41, 42]. The increased expression of binding sites for several lectins by AMs from patients with interstitial lung disease may reflect the influx of immature blood monocytes and/or the emergence of a proinflammatory macrophage phenotype [43]. In a recently published study, however, the authors postulated that the majority of AMs recovered by BAL in sarcoidosis are mature, activated cells [44].

To characterize the AM subsets in the present study, we used a panel of monoclonal antibodies directed against CD11b, CD14, CD16, and HLA-DR surface molecules on AMs. CD11b is the α -chain of a β_2 -integrin that is expressed at high levels on blood monocytes [36], quantitatively upregulated on the surface of fresh extravasated monocytes [45], and expressed at only low levels on mature AMs [35]. CD14 is a receptor for LPS that is expressed at high levels in normal blood monocytes, and at low levels in the subset of CD14+/CD16+ monocytes, as well as in mature AMs [46, 47]. CD16, the low-affinity receptor for IgG (Fc γ RIII), in contrast, is expressed at only low levels on blood monocytes, but is

highly expressed on end-stage differentiated macrophages [47]. The MHC class II molecule HLA-DR has been shown to be expressed both on blood monocytes and AMs, with a considerably higher staining intensity in AMs [48].

The results presented here confirm the findings reported by others in patients with chronic inflammatory lung disease, i.e. a monocyte-like immunophenotype of the total AM population, displaying increased expression of CD14 [13, 14, 17, 35] and CD11b [35, 38], and reduced expression of CD16. Our quantitative flow cytometric analyses, however, extend these findings in that they demonstrate that in the BAL fluid of patients with chronic inflammatory lung disease increased numbers of monocyte-like cells can be identified that exhibit the electrooptical characteristics of blood monocytes and an immunophenotype intermediate between blood monocytes and AMs. The close positive correlations between numbers of monocyte-like cells and expression of monocyteassociated (CD11b, CD14) antigens, as well as the inverse correlations between monocyte numbers and macrophageassociated antigens (CD16, HLA-DR) on the surface of the total AM population provide further evidence for an exaggerated monocyte influx. When calculating the densities of antibody binding sites, we found that alveolar monocyte-like cells express significantly higher numbers of CD11b and CD14 receptors per unit, and significantly lower numbers of CD16 and HLA-DR molecules per unit compared to mature AMs. Taken together, these data strongly suggest that blood monocytes recruited from the pulmonary microvasculature to the bronchoalveolar space are likely to be responsible for the observed alterations in surface molecule expression on the total alveolar monocyte/macrophage population.

Interestingly, we detected a considerable percentage of monocyte-like cells (9±1%) in the BAL fluid of control subjects. The monocyte chemoattractant MCP-1 has been shown to be present in the BAL fluid of healthy control subjects [12]. Thus, MCP-1 produced by normal AMs or alveolar type II epithelial cells may, in part, be responsible for the maintenance of the AM population under noninflammatory conditions. In chronic inflammatory lung disease, however, high numbers of monocyte-like cells are closely associated with impaired lung function, as indicated by the results presented here.

In summary, this study extends previous findings regarding the immature phenotype of the alveolar macrophage in chronic inflammatory lung disease. Our quantitative flow cytometric data confirm the appearance of increased numbers of alveolar monocyte-like cells in the patients' bronchoalveolar lavage fluid, suggesting that in human chronic inflammatory lung disease considerable numbers of monocytes are recruited into the bronchoalveolar space. The sequence of events leading to a "monocytic alveolitis", such as monocyte-endothelium cell interactions, transmigration through the endothelial layer, migration through the pulmonary interstitium, including various cell-matrix interactions, and, finally, emigration into the alveolar space, may play a key role in the pathogenesis of interstitial lung diseases. Accumulation of these inflammatory cells in the lungs of patients with chronic inflammatory lung disease may, in turn, contribute to the disease process by virtue of exaggerated production of oxidants, cytokines and other mediators.

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