The influence of different culture microenvironments on the generation of dendritic cells from non-small-cell lung cancer patients

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Abstract

Introduction: Monocyte-derived dendritic cells (DCs) are currently under extensive evaluation as cell vaccines for cancer treatment. Many protocols regarding DCs generation *in vitro* with different protein components, especially autologous proteins, have been described. On the other hand, active tumor-derived factors in patients' serum could impair monocytes, which might result in their abrogated differentiation into DCs *in vitro*.

Materials and Methods: Autologous DCs from non-small-cell lung cancer (NSCLC)-bearing patients were generated in different culture microenvironments. Peripheral blood mononuclear cells (PBMCs) were cultured in the presence of interleukin-4 and granulocyte-monocyte-stimulating factor with supplementation of 10% autologous serum, 10% allogenic serum, or 2% human albumin. The course of apoptosis, phagocytic ability, and the immunophenotype of the generated DCs were analyzed using flow cytometric methods.

Results: After 48 h of culture, we found a lower percentage of $CD1a^+/CD14^+$ and a higher percentage of $CD1a^+/CD14^-$ cells in the culture supplemented with human albumin than in the cultures supplemented with serums. The lowest CD14 antigen expression was found in the human albumin-supplemented 48-h cultures. After 48 h in the cultures carried out with human albumin we found significantly higher percentages of AV^+/PI^+ cells and AV^-/PI^+ cells than in cultures supplemented with autologous or allogenic serum. We also noted that the expression of FITC-dextran after 4 and 24 h of incubation was significantly higher in the cultures supplemented with both serums than in the HA-SC. The percentage of semi-mature DCs and of CD83 expression was lowest in the culture supplemented with 2% human albumin.

Conclusions: The kind of culture supplementation had a great impact on the apoptosis of cultured PBMCs. It could also influence the yield of monocyte-derived DCs. It was also confirmed that autologous and allogenic serums provide suitable microenvironments for the generation of autologous DCs from NSCLC patients. The choice of culture supplementation for DC generation is still unsolved and further studies should be undertaken.

Key words: dendritic cells, non-small-cell lung cancer, immunophenotyping, cell culture.

Abbreviations: AlloS-SC – allogenic serum-supplemented culture, APC – antigen-presenting cell, AutoS-SC – autologous serum-supplemented culture, AV – annexin V, DCs – dendritic cells, FCS – fetal calf serum, HA-SC – human albumin-supplemented culture, NSCLC – non-small-cell lung cancer, PBMCs – peripheral blood mononuclear cells, PI – propidium iodide, IL-4 – interleukin-4, PBS – phosphate-buffered saline.

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INTRODUCTION

Lung cancer has the highest mortality rate amongst men and women in western countries [4, 47]. Non-smallcell lung cancer (NSCLC), which includes adenocarcinoma, squamous cell carcinoma, and giant cell carcinoma, comprises the majority of cases and is refractory to conventional therapeutic modalities [4, 19, 47]. About 50% of NSCLC patients have stage IV disease at the time of diagnosis and their median survival rate seldom exceeds 6–9 months [19]. What is more, stages IIIB and IV, which preclude surgical therapy, account for 70% of all cases. Chemotherapy and radiotherapy applied to that type of cancer are of low efficacy [4, 19, 24, 48]. The

lack of a standard therapy has prompted study of the development of novel therapies for the treatment of NSCLC. One of them is the administration of drugs that suppress neoplastic cell proliferation by inhibition of the epidermal growth factor receptor; another consists of immunotherapy using unspecific factors that stimulate the immune system, such as BCG vaccine, cytokines which modulate the immunological response, neoplastic antigens, and immune system cells [4, 19, 47]. There have been many attempts to treat NSCLC patients with cellular therapies, such as lymphokine-activated killer cells or tumor-infiltrating lymphocytes, although results have not been encouraging [4, 47, 50].

The discovery of dendritic cells (DCs) as professional antigen-presenting cells (APCs) opened up new possibilities for their use in the development of cancer vaccines [10, 17, 20, 28, 38, 41, 49]. The principal purpose of DC-based immunotherapy is to induce an antigen-specific immune response [20, 38]. The effector arm of anti--tumor immunity are CD4⁺ and CD8⁺ cells, which can only recognize antigen when it is presented by APCs [17, 26, 38]. Owing to their unique properties, DCs, which are the most potent APCs, are considered attractive adjuvants in such immunotherapy [30, 38]. The preparation of a DCs vaccine comprises of the generation of DCs from CD34⁺ precursor cells or monocytes, loading the generated DCs with specific tumor antigens and, finally, administration of the vaccine to the patient [10, 26, 30, 41]. Upon administration, the DCs should migrate to secondary lymphoid organs and induce an antigen-specific immune response [17, 30]. DC-based cancer vaccines are used in patients with melanomas, myelomas, prostate cancers, renal cancers, breast cancers, ovarian cancers, and gastrointestinal cancers [10, 20, 26, 28, 38].

It is well known that in tumor-bearing hosts the immune system frequently does not react to the tumors [8, 15, 32, 45]. The tumoral microenvironment is usually referred to as a "black hole", where the mechanisms which impair immune responses are not clearly identified [6, 18, 26, 32]. Clinical observations showed that DCs possess the ability to migrate directly to a neoplastic site, but it is still not clear why they are unable to induce an efficient immune response [37, 42]. A growing body of evidence suggests that autologous DCs in a cancer-bearing host are not fully competent, which could account for the decreasing, or even obstructive, efficiency of vaccination [10, 18, 22, 23, 26, 31]. How strongly could tumor-derived factors inhibit the host immune system?

With the above problems in mind, the aim of the present study was to verify the ability to generate *in vitro* DCs from NSCLC patients. The cultures were prepared with an addition of 10% autologous serum from the patients to create a microenvironment resembling the *in vivo* appearance [12]. DCs were also generated in medium supplemented with 2% human albumin to avoid the possible inhibitory effect of the tumor microenvironment [2]. Additionally, a generation with healthy human serum was cultured to establish the influence of physio-

logical serum on DCs cultures. These methodologies could be conducive to improving the methods of generating autologous DCs from NSCLC patients and to adopting them in future immunotherapy.

MATERIALS AND METHODS

Clinical characteristic of the lung cancer patients

The study population comprised of 14 NSCLC subjects (mean age: 61.75 ± 9.92 years). Confirmation of the histopathologic type of lung cancer was obtained for all patients. Laboratory tests, chest X-rays, computed tomography, ultrasonography, and bronchoscopy were used for the evaluation of the lung cancer stage and type of disease manifestation (Table 1). Stages from I to IIIA were specified as local disease, whereas stages IIIB and IV were specified as generalized lung cancer. Owing to the small number of examined patients, this division was applied in the statistical analysis [1].

Preparation of serum

Autologous serum was obtained by centrifugation (3000 rpm/10 min.) of peripheral blood samples taken from each patient, which were then frozen in aliquots (10 aliquots of 1 ml, 4 aliquots of 200 µl) at 80°C and successively thawed when examined. Allogenic serum obtained from 14, only AB healthy volunteers served as controls. The serums were not mixed. The serum samples was used to supplement the culture medium and for cytokine measurement using ELISA.

Generation of DCs

Peripheral blood was drawn from NSCLC patients into heparinized tubes according to the protocols accepted by the local ethics committee. Peripheral blood mononuclear cells (PBMCs) were isolated by density gradient centrifugation (Lymphoprep, Nycomed, Norway). Total mononuclear cell counts oscillated between 33 and 54 millions cells. After isolation, the PBMCs were resuspended in RPMI 1640 medium (BioWhittaker, Belgium) supplemented with penicillin--streptomycin (1% v/v, Sigma, Germany), seeded in 6--well tissue culture plates in an appropriate culture microenvironment, and left to adhere for 1.5 h at 37°C in a humidified 5% CO₂ atmosphere. The following culture systems were carried out: 1) human albuminsupplemented cultures (HA-SC): PBMCs from patients in medium containing 2% human albumin [2]; 2) autologous serum-supplemented culture (AutoS-SC): PBMCs from patients in medium containing 10% autologous serum; 3) allogenic serum-supplemented culture (AlloS-SC): PBMCs from patients in medium containing 10% allogenic serum from a healthy donor. After the adherence time, non-adherent cells were removed and the culture plates were washed with cold phosphate-

Table 1. Clinical characte	istic of examined patients
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No.	Age (years)	Gender	Disease	TNM	Stage	Localization	Prospects	Leukocyte count
1	51	М	SCC	$T_4N_3M_1$	IV	right, central	death	above normal
2	50	М	SCC	$T_2N_0M_0$	Ι	right, central	chemotherapy	normal
3	65	М	SCC	$T_3N_2M_0$	IIIA	left, central	chemotherapy	normal
4	53	Μ	AC	$T_4N_2M_1$	IV	left, central	death	normal
5	58	Μ	SCC	$T_2N_1M_0$	II	left, central	surgery	above normal
6	75	М	SCC	$T_3N_2M_0$	IIIA	left, central	surgery	above normal
7	53	F	other*	$T_3N_3M_0$	IIIB	right, peripheral	death	normal
8	70	М	AC	$T_4N_2M_0$	IIIB	left, central	chemotherapy	normal
9	76	Μ	other**	$T_4N_2M_0$	IIIB	right, central	chemotherapy	above normal
10	78	М	other**	$T_4 N_3 M_0$	IIIB	right, peripheral	chemotherapy	normal
11	62	М	SCC	$T_4 N_1 M_0$	IIIB	right, central	chemotherapy	above normal
12	70	М	GCC	$T_2N_1M_0$	II	left, peripheral	surgery	normal
13	63	М	SCC	$T_4N_3M_0$	IIIB	left, central	chemotherapy	above normal
14	67	М	SCC	$T_{3}N_{1}M_{0}$	IIIA	left, central	surgery	normal

SCC – squamous cell carcinoma, AC – adenocarcinoma, GCC – giant cell carcinoma, other – including *low-differentiated NSCLC and **unclassified NSCLC, TNM – classification of tumors [4].

-buffered saline (PBS) without Ca2+ and Mg2+ (Biochrom AG, Germany) to obtain a pure fraction of adherent CD14⁺ monocytes. The adherent cells were then cultured in an appropriate medium supplemented with recombinant human interleukin (IL)-4 (500 IU/ml, Strathmann, Germany) and recombinant human granulocyte-monocyte-stimulating factor (1000 IU/ml, Gentaur, Belgium). The cells were fed every three days. On the 5th day, recombinant human tumor necrosis factor α (50 ng/ml, Strathmann, Germany) was added to induce terminal maturation of the generated DCs. On day 7, at the end of the culture, the supernatants were collected and the adherent cells were harvested by incubation with a trypsin/EDTA solution (0.25%/0.02%, Biochrom AG, Germany). The obtained mature DCs were washed in PBS without Ca2+ and Mg2+ and resuspended prior to immunophenotyping. The yield of cultured DCs was 3-4% of the initial PBMC counts.

Phagocytosis assay

To measure the phagocytic activity of the generated DCs, 5-day cultures in the aforementioned systems were also carried out. At the end of the culture, the generated immature DCs were resuspended in RPMI 1640 and incubated with FITC-dextran (ICN, Germany) at a concentration of 1 mg/ml of medium at 37°C for 4 and 24 h and at 4°C for 4 h. The incubations were stopped by adding cold RPMI 1640. The cells were washed twice with cold RPMI 1640 and immediately analyzed by flow cytometry.

Apoptosis analysis in the course of the cell cultures

For quantification of the cells undergoing early programmed cell death, an annexin V (AV) and propidium iodide (PI) apoptosis detection kit (R&D, USA) was utilized according to the manufacture's protocol. The cells were cultured in the above systems and then removed from the culture plates after 48 and 96 h of incubation. Freshly isolated PBMCs were also examined. The cells were washed with PBS without Ca^{2+} and Mg^{2+} and resuspended in the binding buffer provided. AV (FITC-conjugated) and specific monoclonal antibodies (anti-CD14 TC-conjugated and anti-CD1a PE--conjugated, Caltag, USA) were added. The cells were incubated for 15 min at room temperature in the dark. PI was added in the last 5 min of incubation to the tubes containing only AV; then the cells were analyzed by flow cytometry.

Immunophenotyping of generated DCs

To characterize the phenotype of the generated DCs, the following panel of FITC-, PE-, and TC-conjugated monoclonal antibodies (Caltag, USA) was used: isotype control mouse anti-human IgG1/IgG2/IgG2a, mouse anti-human CD45/CD14, mouse anti-human CD83/CD1a/HLA-DR, and mouse anti-human CD80/ /CD86/HLA-DR.

The cells were incubated for 20 min in the dark and then washed in PBS without Ca^{2+} and Mg^{2+} . The cells were analyzed using a FACSCalibur flow cytometer (Becton Dickinson, USA) equipped with a 488-nm argon laser. An acquisition gate was established based on forward scatter (FSC) and side scatter (SSC) that included only the monocyte population (mononuclear cells) and excluded lymphocytes. After acquisition, the cells were analyzed with CellQuest Software.

ELISA analysis of IL-10 level in the patients' serum

The level of IL-10 was measured in the patients' serum using the Quantikine HS Human IL-10 ELISA Kit (R&D Systems, USA) and $EL_X 800$ equipment (Bio-Tek Instruments, USA).

Statistical analysis

The data are presented as the mean value \pm SD. To evaluate the statistical significance, the Mann-Whitney U-test was used to compare unpaired group data. The Wilcoxon matched pairs test was applied for comparing two variables. We also applied the non-parametric Kolmogorow-Smirnow test to compare the two groups. The Spearman correlation test was used for correlating observations. A level of p<0.05 was considered significant.

RESULTS

The influence of culture supplementation on the expressions of CD1a and CD14 antigens as well as cell apoptosis

In this investigation the cell population was analyzed using an assay for phosphatidylserine expression as originally developed by Vermes et al. [44]. We defined early apoptotic cells as AV⁺/PI⁻, late apoptotic cells and necrotic cells as AV⁺/PI⁺, and dead cells as AV⁻/PI⁺. A representative cytometric analysis is shown in Fig. 1.

After 48 h of culture, we found a significantly (p<0.05) lower percentage of CD1a⁺/CD14⁺ cells in the culture supplemented with human albumin than in the cultures supplemented with autologous or allogenic serum. Moreover, in the culture with human albumin we

found a significantly (p < 0.05) higher percentage of CD1a⁺/CD14⁻ cells than in the cultures carried out with the addition of autologous or allogenic serum (Table 2).

The expression of CD14 antigen was significantly higher on freshly isolated cells than on cells cultured in all types of medium. In the 96-h cultures a small population of cells, resembling macrophages, with a high expression of CD14 antigen and high values of the FSC and SSC parameters, appeared. These macrophage-like cells were also found at the end of the DC generation (Fig. 2). The lowest CD14 antigen expression was found in the human albumin-supplemented 48-h cultures. A significant (p<0.05) increase in CD1a antigen expression was observed between 48 and 96 h in all cultures.

After 48 h of culture, the percentage of AV^+ cells among the populations of $CD1a^+/CD14^+$ cells and $CD1a^+/CD14^-$ cells was significantly higher in the HA--SCs than in the autologous or AlloS-SC. More interestingly, in the cultures carried out with human albumin we found significantly higher percentages of AV^+/PI^+ cells and AV^-/PI^+ cells than in cultures supplemented with autologous or allogenic serum (Table 2, Fig. 3).

In HA-SC after 96 h of incubation we found a significantly higher percentage of AV^+ cells amongst $CD1a^+/CD14^+$ and $CD1a^+/CD14^-$ cell populations in comparison with AutoS-SC and AlloS-SC. We also noted a significantly higher percentage of AV^+/PI^+ cells in culture supplemented with human albumin compared with the two other culture systems.



Fig. 1. Representative cytometry analysis of AV and PI expression on generated mononuclear cells after 48 and 96 h of incubation. A – culture supplemented with autologous serum, B – culture supplemented with allogenic serum, C – culture supplemented with human albumin. Dot plots correspond to the gated cells from region R1.

Percentage of	Refore culture	Autologous serum		Allogenic serum		Human albumin	
generated cells	Delore culture	48 h	96 h	48 h	96 h	48 h	96 h
CD1a ⁺ /CD14 ⁺	79.72 ±15.33	87.67 ±8.83	84.66 ±13.69	88.84 ±9.15	82.48 ±15.2	79.45 ±8.38	76.02 ± 22.07
CD1a ⁺ /CD14 ⁻	7.07 ± 7.13	7.57 ± 7.44	8.22 ±8.37	6.7 ± 4.43	10.74 ± 10.54	13.46 ± 6.5	15.01 ± 17.29
AV+/CD1a+/CD14	+ 3.59 ±2.13	25.55 ±8.35	33.95 ±23.36	25.68 ±13.93	32.06 ± 22.16	43.6 ± 16.36	49.9 ± 18.84
AV+/CD1a+/CD14	- 2.5 ±1.54	28.54 ± 12.99	18.82 ± 13.56	27.95 ± 14.1	20.02 ± 7.5	49.8 ± 14.8	39.8 ± 18.4
AV ⁺ /PI ⁻	0.64 ± 1.0	8.49 ± 5.93	6.58 ±5.43	10.13 ± 9.24	7.93 ±4.73	6.35 ±4.31	9.14 ± 5.03
AV ⁺ /PI ⁺	2.16 ± 1.72	12.33 ±7.79	24.04 ± 12.07	12.79 ±8.96	22.8 ± 9.94	40.62 ± 15.8	35.34 ±12.9
AV ⁻ /PI ⁺	2.19 ± 0.78	2.97 ± 2.86	6.88 ±3.74	2.42 ± 2.26	11.19 ± 8.56	6.5 ±3.77	9.68 ± 9.31
MFI CD14	681 ±454	151.1 ±64.27	206.14 ± 181.1	152.18 ±66.8	146.03 ±82.9	73.63 ±39.3	132.05 ±91.46
MFI CD1a	98.8 ± 70.65	64.9 ± 13.98	118.41 ±64.6	68.8 ± 16.17	110.7 ± 42.4	68.32 ±33.84	139.98 ±98.6

MFI - mean fluorescence intensity.



Fig. 2. Macrophage-like cells and DCs generated after 7-day culture supplemented with 10% autologous serum. The cells were stained by the May-Grünwald Giemsa method.

We did not find any significant differences in the examined parameters between the cultures supplemented with autologous serum and allogenic serum after 48 and 96 h of incubation. The percentages of cells generated in all the cultures are shown in Table 2 and Fig. 3.

Relationship between cancer stage and the expressions of CD1a and CD14 antigens as well as cell apoptosis in differently supplemented cultures

After 96 h of incubation in AutoS-SC and AlloS-SC, a significantly higher percentage of double-positive CD1a⁺/CD14⁺ cells and lower percentage of CD1a⁺//CD14⁻ cells were found in patients with advanced disease than in subjects suffering from earlier stages of lung cancer. Interestingly, a significantly lower percentage of AV⁺/PI⁺ and AV⁺/PI⁻ cells were detected in patients with advanced stages of lung cancer than in



Fig. 3. Percentages of apoptotic, late apoptotic, and necrotic cells before and after 48 h of culture in different microenvironments.

patients with local tumor growth. The percentages of examined cells were similar in both groups of patients in the cultures performed with human albumin.

There were significant negative correlations between the percentages of CD1a⁺/CD14⁻ cells and AV⁺/PI⁻ cells in AutoS-SC and AlloS-SC after 48 and 96 h of incubation in all the patients (R=–0.70, p<0.05). Also, after 96 h of incubation in AutoS-SC the percentages of CD1a⁺/CD14⁻ cells and of AV⁺/PI⁻ cells correlated significantly negatively with patient leukocytosis.

Serum IL-10 level and its influence on the expressions of CD1a and CD14 antigens and cell apoptosis

The total mean level of IL-10 in the patients' serum was 6.57 ± 3.51 pg/ml. Serum IL-10 level was significantly higher (p=0.01) in NSCLC patients than in control subjects (2.957±2.177 pg/ml). The IL-10 concentration was similar in patients with different clinical stages of the disease.

After 48 h of culture with autologous serum supplementation we found a significant negative correlation between the percentage of CD1a⁺/CD14⁺ cells and the level of IL-10 (R=-0.69, p<0.05). We also noted positive correlation between the IL-10 level and the percentage of CD1a⁺/CD14⁻ cells (R=0.69, p<0.05). The

 Table 3. Expression of FITC-dextran phagocytozed by immature DCs cultured with autologous serum, allogenic serum, and human albumin supplemented cultures

Culture microenvironment	Cells incubated at 4°C	Cells incubated at 37°C for 4 h	Cells incubated at 37°C for 24 h
10% Autologous serum	15.1±10 MFI	28.29±11.95 MFI	37.39±17.4 MFI
10% Allogenic serum	15.55±12.92 MFI	25.47±11.7 MFI	38.02±18.72 MFI
2% Human albumin	18.78±22.56 MFI	20.69±9.13 MFI	24.9±11.42 MFI

MFI - mean fluorescence intensity.



Fig. 4. Phagocytosis of FITC-dextran particles by immature DCs generated from representative patients in: A - AutoS-SC, B - AlloS-SC and C - HA-SC, after 4 and 24 h of incubation. Dotted line: control cells incubated at 4°C, continuous line: cells incubated at 37°C. Histograms correspond to the gated cells from region R1 and presented mean fluorescent intensity of FITC-dextran and cell counts.

IL-10 level correlated slightly negatively with the percentage of AV^+/PI^- cells (R=-0.60, p=0.11) and non--significantly positively with the percentage of AV^+/PI^+ cells (R=0.57, p=0.14). There were no significant correlations after 96 h of culture with autologous serum supplementation.

Engulfment of FITC-dextran by generated immature DCs

To test the phagocytic function of immature DCs generated in the above three culture systems, we evaluated the uptake of FITC-dextran by DCs using flow cytometry. The results are presented as the mean fluorescence intensity of dextran-positive cells and are shown in Table 3. The flow cytometric analysis of FITC-dextran engulfment by generated cells is shown in Fig. 4. In all the cultures we found a significantly (p<0.001) higher expression of FITC-dextran in cultured cells after 4 and 24 h of incubation than in control cells incu-

bated at 4°C. We also noted that the expression of FITC-dextran after 4 and 24 h of incubation was significantly (p < 0.05) higher in the cultures supplemented with autologous and allogenic serum than in the HA-SC. There were no significant differences between cells derived from autologous and AlloS-SCs in FITC-dextran expression.

Immunophenotype of generated DCs

The generated DCs were divided into two groups according to the expressions of leukocyte-monocyte-specific markers: double-positive cells (CD45⁺/CD14⁺), which did not lose their monocytic attribute during the cultures, and CD45⁺/CD14⁻ cells, which appeared to lose their monocytic feature and turn into DCs. In all the cultures we observed low percentages of CD45⁺/CD14⁺ cells and high percentages of CD45⁺/CD14⁻ cells. We did not find any significant differences



Fig. 5. Morphology of generated mature DCs in culture: A – supplemented with 10% autologous serum; B – 10% allogenic serum; C – 2% human albumin. The cells were stained by the May-Grünwald Giemsa method.

in the examined cell populations and in CD14 antigen expression in all the cultures.

In all the cultures we observed cells with typical DC morphology (Fig. 5). According to the expression of CD1a antigen, we recognized semi-mature DCs as CD83⁺/CD1a⁺/HLA-DR⁺ cells and mature DCs as CD83⁺/CD1a⁻/HLA-DR⁺ cells [36, 52]. The percentage of semi-mature DCs and of CD83 expression was lowest

in the culture supplemented with 2% human albumin. Moreover, we found a significantly higher percentage of CD83⁺/CD1a⁺/HLA-DR⁺ cells and a significantly higher expression of CD83 antigen in the culture supplemented with autologous serum than in the culture with human albumin.

We also divided the generated DCs according to the expressions of co-stimulatory molecules into CD80⁺/(CD86⁺/HLA-DR⁺ cells and CD80⁺/CD86⁻/HLA-DR⁺ cells [11]. There were no differences among all the cultures in the percentages of these cells. A small population of CD80⁻/CD86⁺/HLA-DR⁺ cells in all the examined microenvironments was observed in only one patient (no. 3). A significantly (p<0.001) higher expression of CD86 antigen was observed on cells from AutoS-SC and from AlloS-SC compared with cells from HA-SC. The immunophenotype analyses of the generated DCs are presented in Table 4.

Comparison of patients' clinical stage and immunophenotype of generated DCs

In the culture supplemented with autologous serum we observed a significantly (p<0.001) lower expression of CD14 antigen on DCs generated from the subjects in earlier stages than from those with advanced stages of disease. In the patients with localized lung cancer, the expressions of the HLA-DR and CD86 antigens were non-significantly higher than in the patients with advanced disease. CD86 antigen expression significantly positively (R=0.67, p<0.05) correlated with IL-10 level.

Dendritic cells generated in AlloS-SC from patients with earlier stages of lung cancer were characterized by a significantly (p<0.05) higher expression of HLA-DR and a significantly (p<0.05) lower expression of CD80 antigens than DCs generated from patients with advanced disease. We also found a non-significantly higher expression of CD14 antigen on DCs from patients with advanced stages than from those with localized lung cancer. Interestingly, in the culture supplemented with human albumin, the expression of CD14 antigen was also non-significantly higher in patients with advanced stages than in patients with earlier stages of the disease.

DISCUSSION

Currently, monocyte-derived DCs are becoming widely used as cell vaccines for cancer treatment [10, 26, 38, 49]. The requirement for optimized and standardized protocols for the generation of DCs on a large scale has led to the description of several protocols [2, 17, 21, 33, 34, 49]. In many clinical studies, autologous plasma, fetal calf serum (FCS), allogenic AB plasma, and human albumin were used as serum or protein components to support the generation of DCs [2, 12, 21, 26]. The use of FCS in cultures to prepare DCs precludes their practical application, because FCS contains xenogenic proteins

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Immunophenotype of DCs (%)	10% Autologous serum	10% Allogenic serum	2% Human albumin
CD45 ⁺ /CD14 ⁻	81.9±18.13	84.82±13.04	78.74±17.73
CD45 ⁺ /CD14 ⁺	13.84±15.7	12.33±11.9	17.8±16.8
CD83 ⁺ /CD1a ⁺	52.69±41.01	46.27±38.58	64.03±32.86
HLA-DR ⁺ cells among CD83 ⁺ /CD1a ⁺ DCs	98.53±2.51	97.46±4.26	90.25±25.01
CD83 ⁺ /CD1a ⁻	35.81±42.64	37.19±43.1	29.17±35.12
HLA-DR ⁺ cells among CD83 ⁺ /CD1a ⁻	87.56±18.3	91.51±10.78	83.29±21.11
CD80 ⁺ /CD86 ⁺	82.82±16.66	87.76±13.21	80.66±23.14
HLA-DR ⁺ cells among CD80 ⁺ /CD86 ⁺	86.73±17.37	81.31±22.04	86.62±21.73
CD80 ⁺ /CD86 ⁻	10.28±14.31	5.29±8.04	15.88±23.66
HLA-DR ⁺ cells among CD80 ⁺ /CD86 ⁻	72.25±28.79	71.79 ± 8.04	64.41±35.96

Table 4. Immunophenotype analysis of DCs generated in different culture microenvironments

and other substances which may affect the function of DCs [26]. It is possible to use allogenic human plasma, but this raises the possibility of the transmission of infectious pathogens [21]. It has been proposed that human albumin is a kind of non-antigen culture substrate and can provide the "clearest" culture microenvironment, but one that is destitute of natural growth factors [2]. These limitations suggest that autologous plasma or serum would be a less hazardous choice for DCs generation [12, 34]. Of course, this situation is suitable for healthy volunteers. Unfortunately, there is still insufficient information about the influence of serum from cancer patients on DCs generation from CD14⁺ monocytes. The active tumor-derived factors in patients' serum could impair monocytes, which might result in their abrogated differentiation into DCs in vitro [1, 22, 31, 35, 40]. The questions that then arise are whether tumors affect DCs generation and whether it is possible to reverse the restraining effect of the tumor microenvironment under in vitro conditions. To answer the questions, the aim of this study was to verify the ability to generate DCs from NSCLC-bearing patients in cultures supplemented with autologous serum, allogenic serum, and human albumin.

In this study we first tried to evaluate the fate of cultured monocytes and the induction of apoptosis on early differentiated cells. It is known that the induction of apoptosis does not occur only by direct contact with tumor cells [23, 46]. It has been demonstrated that murine DCs treated with supernatants from tumor cell lines undergo apoptosis [46]. By combining AV, which was shown [43] to have a high affinity for phosphatidylserine, with PI we discriminated between early apoptotic cells (AV+/PI-), late apoptotic cells and necrotic cells (AV⁺/PI⁺), and death cells (AV⁻/PI⁺) among the mononuclear cells in different culture microenvironments. We found that the highest percentage of AV⁺ cells as well as the highest percentages of AV⁺/PI⁺ and AV⁻/PI⁺ cells were in those cultured for 48 h in human albumin. The highest percentages of apoptotic and necrotic cells were also detected after 96 h in the aforementioned cultures' microenvironments. Moreover, the highest percentage of CD1a⁺/CD14⁻ cells was found in the HA-SC cells. Unexpectedly, this pattern also appeared when we compared the subtypes of cells from AutoS-SC from patients in different stages of lung cancer. The percentages of AV⁺ cells as well as the percentage of CD1a⁺/CD14⁻ cells were highest in patients with early stages of lung cancer. We noted negative correlation between the percentages of CD1a⁺/CD14⁻ and AV⁺/PI⁻ cells as well as negative correlation between patients' leukocytosis and the percentages of the above cells.

It is known that monocytes lose their CD14 antigen expression during their differentiation into DCs and turn into a CD1a⁺/CD14⁻ cell population [5, 26, 34]. In our experiment schedule we found a significantly decreasing expression of CD14 antigen in all the performed cultures, but most of all in the HA-SCs. In connection with the apoptosis levels in the human albumin culture after 48 and 96 h, we can suggest that the decrease in CD14 antigen expression is caused by the cells dying rather than their differentiation into immature DCs. CD14 antigen could be peeled off by apoptotic cells. This could be confirmed by results obtained in cultures performed with autologous or allogenic serum supplementation. In the aforementioned culture conditions we found lower percentages of apoptotic and necrotic cells as well as a higher expression of CD14 antigen compared with the HA-SCs. On the other hand, Avila-Moreno et al. [3] found that soluble factors from lung cancer cell lines caused an increase in the expression of myelomonocytic markers and altered the differentiation of monocytes in vitro from DCs to macrophages. In our study we also observed macrophage-like cells in AutoS-SC and AlloS-SC, but the percentage of these cells was very small. The results presented here let us speculate that autologous serum, in the case of lung cancer, provides a better culture microenvironment for the generation of autologous cells, which seem to be prevented from early apoptosis.

The cause of this phenomenon remains unclear. It has been shown repeatedly that factors which possess a strong inhibitory capacity can be detected in the serum of cancer-bearing patients [22, 23, 29]. Some trials have been carried with tumor culture supernatants (TSN). Kiertscher et al. [23] have shown a reduced capacity to produce IL-12 by DCs generated with TSN supplementation. However, Marten et al. [27] have shown the strong stimulatory activity of DCs generated with patient-derived CA-19-9-antigen-containing serum. The HA-SCs are a clear environment for DCs generation, but it seems to be devoid of important cell viability factors. Interestingly, Araki et al. [2] generated efficient DCs from CD14⁺ monocytes of healthy volunteers in the presence of human albumin. It is widely known that many cytokines or immunosuppressive cancer-derived factors detected in the serum of cancer patients interfered in many ways with each stage of the generation of DCs [7, 14, 16, 22, 29, 35, 39]. It has been suggested that IL-10 is the most important inhibitory factor responsible for impaired immune responses in the peripheral blood of cancer-bearing patients [7, 13, 14]. A variety of human tumors produce IL-10. The mRNA for IL-10 was detected in metastatic melanoma and renal, ovarian, and basal cell carcinomas [25, 32]. We have also shown that the level of IL-10 detected in patients' serum positively correlated with the percentage of CD1a⁺/CD14⁻ cells and AV⁺/PI⁺ cells in AutoS-SC.

It should be noted that in most cancer patients the inflammation process coexists with the development of neoplasm, which could result in a greater release of immunomodulatory factors [9]. Our results suggest that the progression of lung cancer had a beneficial impact on the vitality of mononuclear cells in short-terminated cultures. In light of the present results, the restraining effect of tumor-derived factors in autologous serum from NSCLC patients appears to be overestimated. We propose that generalized lung cancer does not exert as strong an inhibitory pressure on autologous DCs generation as suspected up to now.

Some literature data concerned the generation of DCs from patients with advanced cancer [26, 29, 33, 51]. Menetrier-Caux et al. [29] demonstrated that tumor cells produce factors which could significantly inhibit the differentiation of hematopoietic precursors and, in turn, DCs generation. It has been shown repeatedly that the expressions of CD83 and CD86, human DC-related markers, decreased in DCs cultures co-incubated with supernatant from human tumor cells [7, 13]. In this study we estimated the presence of mature CD83⁺/CD1a⁻ DCs with co-expression of costimulatory molecules in all the cultures. Interestingly, the highest expression of CD83 antigen was found in AutoS-SC. These findings are contrary to those of Kiertscher et al. [23], who had demonstrated that tumor culture supernatants effectively suppress the function of monocyteand stem cell-derived DCs. Onishi et al. [31] showed that monocyte-derived DCs from cancer patients possessed weaker antigen-presenting abilities than those generated from healthy volunteers. In our study we found that immature DCs generated in all the cultures show phagocytic activity measured by dextran particle engulfment. When we compared the culture microenvironments we found the highest level of FITC-dextran fluorescence in cells generated with autologous and allogenic serum supplementation. This confirms the above thesis about the beneficial influence of autologous and allogenic serum on DCs generation.

To sum up, we have documented the course of the apoptosis, phagocytic ability, and the immunophenotype of DCs generated in different culture microenvironments. First of all we can conclude that the kind of culture supplementation had a great impact on the apoptosis of generated DCs. It could also influence the yield of monocyte-derived DCs, which should be examined in a further study. Secondly, we can confirm that autologous and allogenic serums provide a suitable microenvironment for the generation of autologous DCs from NSCLC patients. Of course, further studies should expand upon lymphocytic stimulation by generated DCs, which could provide an important insight into the stimulatory activity of DCs. Also, the efficacy of DCs generation from PBMCs of lung cancer patients should be tested using serum-free media available commercially, such as CellGro DC. The choice of culture supplementation for DCs generation is still unsolved and further studies should be undertaken.

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