Phenotypic characteristics of tumour-infiltrating lymphocytes in human oesophageal cancer tissues defined by quantitative two-colour analysis with flow-cytometry

Hideaki Tahara¹, Hitoshi Shiozaki¹, Kenji Kobayashi¹, Tokiharu Yano¹, Hiroshi Yano¹, Shigeyuki Tamura¹, Kunihiko Oku¹, Mikiyo Miyata¹, Ken'ichi Wakasa², Masami Sakurai², and Takesada Mori¹

Summary. Phenotypic subpopulations of tumourinfiltrating lymphocytes (TIL) separated from human oesophageal cancer tissues were defined by quantitative two-colour analysis with flow-cytometry (FACScan), and their characteristics were investigated by comparison with peripheral blood lymphocytes (PBL) and intra-oesophageal lymphocytes from noncancerous tissue (IEL) as the controls. Fifteen patients (13 males and 2 females) with squamous cell carcinoma of the oesophagus were entered into the study. Lymphocytes were analyzed by FACScan and the frequencies of the subpopulations were determined using monoclonal antibodies for surface markers. Single colour analysis revealed a predominance of T cells among TIL and a significant reduction of natural killer (NK) cells compared with the controls. Two-colour analysis showed that CD4⁺Leu8⁻ (helper T cells) and CD8⁺CD11b⁻ (cytotoxic T cells) were significantly increased among TIL when compared with the controls. This significant increase of both helper and cytotoxic T cells, which are indispensable components of cellular immunity, strongly implies that TIL are performing a role in the expression of antigen-specific cellular immunity against the tumours. This is the first report of a phenotypic study of human oesophageal cancer that clearly indicates the significance of the TIL and suggests their potential for use as a source of adoptive immunotherapy.

Key words: Oesophageal cancer – Tumour infiltrating lymphocyte – Phenotypic subpopulation – Cell-mediated immunity – Flow-cytometry

Tumour-infiltrating lymphocytes (TIL) have been suggested as expressions of a tumour-specific host

Offprint requests to: H. Tahara at the above address

response in various solid tumours including oesophageal cancer, judging from their correlations with the length of survival after surgery (Moore and Foote 1949; Takahashi 1961; Martin and Bechwith 1968; Bennett et al. 1971; Berg 1971; Black et al. 1972; Shiozaki et al. 1983). Experimental evidence has supported this hypothesis and implies a tight correlation between TIL and cell-mediated immunity (Kikuchi et al. 1972; Shimokawa et al. 1982; Vose 1982; Hiratsuka et al. 1984).

It has also recently been reported that TIL cultured with interleukin-2 (IL-2) have much more tumour-specific cytotoxicity than lymphokine-activated killer (LAK) cells both in vitro and in vivo (Rosenberg et al. 1986; Rabinowich et al. 1987; Kuppner et al. 1988; Takagi et al. 1989). Thus, TIL have been seen as possible specific effectors for autologous tumour cells and as an important source for adoptive immunotherapy. However, it still remains unclear whether or not the TIL in human oesophageal cancer are really the effector cell population of a specific immune reaction recognizing tumour cells as antigens in vivo.

In order to elucidate their functions, we applied flow-cytometric quantitative two-colour analysis to TIL separated from surgically-resected oesophageal cancer sepcimens. Double staining of the surface markers relevant to function was performed and the phenotypic make-up was compared with that of peripheral blood lymphocytes (PBL) and intra-oesophageal lymphocytes from non-cancerous tissue (IEL) obtained from the same patients.

Materials and methods

Fifteen patients with squamous cell carcinoma of the oesophagus, who underwent subtotal oesophageal resection, were examined in this study. Each patient gave informed voluntary consent. None of the patients had received any therapy prior to

¹ Second Department of Surgery, Osaka University Medical School, and ² Department of Pathology, Osaka University Hospital, 1-1-50, Fukushima, Fukushima-ku, Osaka City, 553, Japan

surgery. The patients ranged in age from 44 to 76 years (mean age = 56) and the male to female ratio was 13:2.

The histological features of the specimens for flow-cytometric analysis were assessed using H &E stained sections taken from immediately adjacent to the specimens used for flow-cytometry. The terminology used in this report for location, staging, and histological type is derived from the Tumour-Node-Metastasis staging system (UICC 1987).

Samples of 15 ml of heparinized venous blood were obtained from all patients on the morning of surgery before the administration of any narcotic agent, and the mononuclear cells were separated by standard Ficoll-Hypaque gradient centrifugation (Boeyum 1968) using LSM (Organon Teknika Corporation, North Carolina, USA) as the separating medium. In brief, blood samples were diluted twice to half their original concentration with Hanks' balanced salt solution (HBSS; Biken, Japan), floated onto LSM, and centrifuged at $400 \times g$ for 30 min. The mononuclear cells were collected at the gradient interface, washed twice with HBSS, and resuspended in HBSS containing 0.1% bovine serum albumin (BSA) and 0.02% NaN₃ (medium A) at a concentration of 5.0×10^6 cells/ml. Cell viability was confirmed to be greater than 98% by trypan blue dye exclusion.

For the isolation of TIL and IEL rich aliquots, tumour tissue and non-cancerous oesophageal tissue samples were obtained from the surgically-resected specimens. Processing was initiated within 2 h of resection. The weights of the specimens used to obtain TIL and IEL ranged from 1 to 2.6 g and 2 to 4 g, respectively. Necrotic tissue and connective tissue was removed. Then the specimen was minced with scissors and suspended in RPMI 1640 (Biken, Japan) containing collagenase (type IV, 200 units/ml, Sigma Chemical Co., St. Louis, MO, USA) and 10% fetal bovine serum (FBS). The suspension was stirred for 2 h at 37° C, passed through a fine nylon mesh filter, harvested by centrifugation, washed twice with HBSS, and suspended in HBSS containing 10% FBS. The cell suspension was then floated on LSM and centrifuged at $400 \times g$ for 20 min. Cells at the interface consisted of lymphocytes, monocytes, macrophages, and tumour cells. They were collected and adjusted to concentrations of from 5×10^5 to 5×10^6 cells/ml in medium A. The viability of the TIL and IEL was confirmed to be greater than 98% by trypan blue dye exclusion, and the numbers of TIL and IEL per weight of wet tissue sample were calculated.

Table 1 shows the monoclonal antibody combinations which were used in this study and also shows the phenotypes they defined. All of them were conjugated with either fluorescein (FITC) or phycoerythrin (PE) (Beckton Dickinson Monoclonal Antibody Center, Mountain View, CA). Leu7 (Si and Whiteside 1983; Lanier et al. 1983), Leu8 (Gatenby et al. 1982; Kansas et al. 1985), and Leu15 (Landay et al. 1983) are the monoclonal antibodies which can detect most of the specific functional capabilities of lymphocytes when paired with the major appropriate antibodies. The PBL, TIL, and IEL (1× 10⁵ cells each) were incubated with 10 μl of the combined antibodies shown in Table I for 20 min at 4° C. The cells were then washed once with HBSS and resuspended in a further 0.4 ml of HBSS. Two-colour analysis of the immunofluorescent staining was performed using a flow-cytometer (FACScan; Beckton Dickinson, Mountain View, CA, USA) according to the method stated below.

Fluorescence-conjugated antibody staining was detected using a FACScan for flow-cytometric analysis. Signal acquisition was triggered on forward light scatter and there was simultaneous acquisition of forward and side light scatter (FSC and SSC) and of optimally compensated two-colour log-fluorescence signals. Data were acquired by a computer module (Hewlett-Packard, series 9000, type 310, USA) using data acquisition

Table 1. Monoclonal antibody combinations and the phenotypes they defined

FITC conjugated	PE conjugated	Defined phenotypes
Leu4 (CD3)	Leu12 (CD19)	pan T cell pan B cell
Leu2a (CD8)	Leu3a (CD4)	cytotoxic/ suppressor T cell helper/inducer T cell
Leu3a (CD4)	Leu8	CD4 ⁺ Leu8 ⁺ : inducer T cell CD4 ⁺ Leu8 ⁻ : helper T cell
Leu2a (CD8)	Leu15 (CD11b)	CD8 ⁺ CD11b ⁺ : suppressor T cell CD8 ⁺ CD11b ⁻ : cytotoxic T cell
Leu7 (HNK-1)	Leu11c (CD16)	CD16 ⁺ : Natural Killer CD16 ⁺ Leu7 ⁻ : most potent NK
HLe-1 (CD45)		CD45 ⁺ : pan leukocyte

FITC=Fluorescein, PE=Phycoerythrin, CD=Cluster differentiation. Leu series monoclonal antibodies were purchased from Becton-Dickinson. Mountain View, CA, USA

software (Consort 30, Beckton Dickinson), and were stored on 3.5-inch floppy disks. For each experiment, a minimum of 10000 and a maximum of 30000 cells was counted.

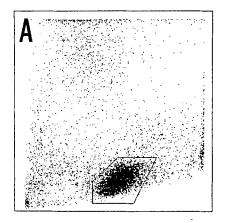
Acquired data files were analyzed by the program for analysis in the Consort 30 software which simultaneously determined two-colour fluorescence data on samples gated by FSC and SSC. For PBL analysis, stored data from the floppy disks were gated through the window established for lymphocytes by their FSC and SSC characteristics (Fig. 1A). For TIL and IEL analysis, in order to separate lymphocytes from contaminating cells such as monocytes, macrophages, and viable tumour cells, the same window was used as for PBL (Fig. 1B), and more than 2000 cells were gated from each specimen. The two-colour fluorescence data of the gated cells were indicated by contour lines and dots on a figure divided into four contiguous areas which were designated as "quadrants" (Fig. 2), and the percentage of cells in each quadrant was calculated simultaneously.

The frequency of lymphocytes among the gated cells was confirmed by the percentage of CD45⁺ cells present. If the gated population showed more than 5% contamination, the frequency of each phenotypic subpopulation was compensated according to the percentage of CD45⁺ cells.

The data were analyzed for statistical significance by the non-parametric Wilcoxon test, and the result was considered significant if the P value was less than 0.05.

Results

All of the primary tumours were located in the intrathoracic oesophagus. Tumour size ranged



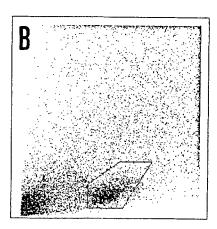


Fig. 1. Gated windows for lymphocytes on FSC and SSC characteristics. The window established for lymphocytes was used for PBL (A). The same window was used for TIL (B) and IEL (not shown). FITC conjugated HLe-1 (CD45, pan leukocyte) monoclonal antibody was used as the positive control. FSC=forward light scatter, SSC=side light scatter

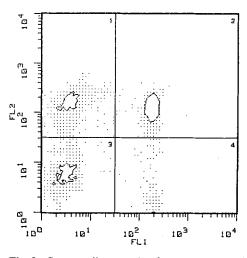


Fig. 2. Contour line graph of two-colour analysis. The dots and the contour lines indicate the numbers of cells determined by dual-fluorescence within the gated populations. This is a graph of the analysis of CD4⁺ and Leu8⁺ cells in case 7. FL 1 indicates green fluorescence (*CD4-FITC*), and FL2 indicates orange-red fluorescence (*Leu8-PE*). Four contiguous areas, which were designated as "quadrants", were defined by the marked lines. The percentage of cells in each quadrant was calculated simultaneously. *dot* = single cell, —= 10 cells

from 2.6 to 8.3 cm in maximum diameter. Most of the tumours (11/15) were classified as T3, 2 were T2, and 2 were T4. Twelve of the 15 subjects had regional lymph node metastases (N1) and the other 3 had no lymph node metastases (N0). Distant metastasis was found intraoperatively in one case. From these findings, 3 of the cases were classified into stage IIA, 1 into stage IIB, 10 into stage III, and 1 into stage IV. Histological examination revealed that all of the tumours were squamous cell carcinoma and they showed a normal distribution of differentiation (well differentiated: 3, moderately differentiated: 8, poorly differentiated: 4).

The number of TIL per gram of wet tissue was $(2.2\pm1.8)\times10^6$ (mean \pm SD), which was significantly increased compared with the number of IEL $((2.4\pm2.4)\times10^5; P<0.005)$.

The means and standard deviations of the frequencies of the subpopulations of TIL, PBL, and IEL detected by the major antibodies are shown in Fig. 3. Anti-CD3 and anti-CD19 antibodies were used to determine the T/B ratio in the 3 populations. The frequency of CD3⁺ cells in the TIL

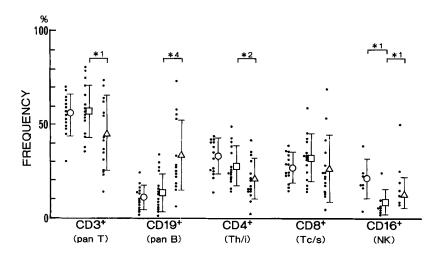


Fig. 3. Phenotyping of the TIL, PBL, and IEL populations as defined by the major antibodies. Symbols and vertical bars represent the mean \pm SD, and a dot indicates the frequency of each specimen. Single-colour analysis of the TIL population revealed a predominance of T cells and a significant reduction of the frequency of NK cells compared with the controls. 0 = PBL, D = TIL, $\Delta = IEL$. Th/i = helper/inducer T cell, Tc/s = cytotoxic/suppressor T cell, NK = Natural Killer cells. Significance of difference; * = P < 0.05, ** = P < 0.025, *** = P < 0.01, **** = P < 0.005

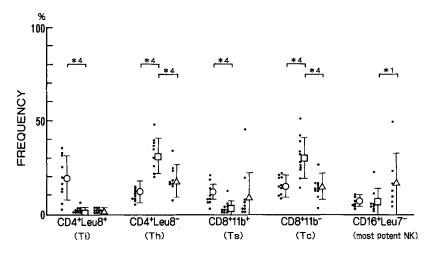


Fig. 4. Phenotypes of TIL, PBL, and IEL populations defined by two-colour analysis (mean \pm SD). Two-colour analysis clearly revealed a significant increase of the helper and cytotoxic T cell subpopulations. $\circ = PBL$, $\square = TIL$, $\triangle = IEL$. Significance of difference; *=P < 0.05, **** = P < 0.005

population was $58.2\pm14.1\%$ (mean \pm SD), which was similar to that for PBL ($55.3\pm10.8\%$), but higher than that for IEL ($45.6\pm19.8\%$; P<0.05). In contrast, CD19⁺ cells comprised $14.0\pm10.5\%$ of the TIL population, which was similar to the proportion in PBL ($10.9\pm6.5\%$), but was much lower than in the IEL population ($33.8\pm18.9\%$, P<0.005). The T/B ratio of TIL (28.6 ± 84.7) was hence significantly higher than that of IEL (2.0 ± 1.4 , P<0.005) but was similar to that for PBL.

Anti-CD4 and anti-CD8 antibodies were used to determine major T cell subsets such as helper/ inducer T cells (Th/i) and cytotoxic/suppressor T cells (Tc/s). The CD8⁺ cells had a similar frequency of occurrence in all 3 cell populations $(TIL:31.8\pm13.1\%;$ PBL: $27.3 \pm 7.7\%$ and IEL: 27.2 ± 17.6%). In contrast, CD4⁺ cells were found at a similar frequency among TIL $(28.2 \pm 10.6\%)$ and PBL $(32.8 \pm 10.1\%)$, but were common among IEL significantly less $(10.9 \pm 10.8\%)$.

The frequency of CD16⁺ lymphocytes among TIL was only $8.2\pm7.4\%$, which was significantly lower than for both PBL and IEL (PBL:20.5±11.3%, P<0.05, IEL:13.5±7.6%, P<0.05).

The CD4⁺, CD8⁺, and CD16⁺ cells of 10 patients (cases No. 6 to 15) were examined in detail by analyzing the coexpression of their second antibodies (Leu8, CD11b, and Leu7, respectively; Table 1, Fig. 4).

The average frequency of Leu8⁺ cells among the CD4⁺ cells in the TIL population was only $5.0\pm4.3\%$, thus the frequency of CD4⁺Leu-8⁺ cells was $2.0\pm2.3\%$ and that of CD4⁺Leu-

 8^- cells was $31.5\pm9.9\%$. Thus, the percentage of CD4⁺Leu8⁻ cells in the TIL population was significantly higher than in both the PBL and IEL populations (PBL:12.1±6.3%, P<0.005; IEL:17.5±8.8%, P<0.005), while the frequency of CD8⁺Leu8⁺ cells was much lower than in the PBL population (19.1±11.8, P<0.005).

The average frequency of CD11b+ cells among CD8+ TIL was $9.7\pm8.0\%$, so the frequency of CD8+CD11b+ cells was $3.2\pm3.7\%$ which was significantly lower than in the PBL population ($12.2\pm7.4\%$, P<0.005). The frequency of CD8+CD11b- cells in the TIL population was $29.6\pm11.3\%$, which was significantly higher than those in both the PBL and IEL populations (PBL: 15.4 ± 6.3 , P<0.005; IEL: 14.6 ± 6.6 , P<0.005).

Cells which are CD16⁺Leu7⁻ are considered to be the most potent NK cells. The frequency of CD16⁺Leu7⁻ cells in the TIL population was $6.7\pm7.8\%$, which was similar to that in the PBL population $(6.6\pm2.9\%)$, and was significantly lower than that in the IEL population $(17.1\pm15.8\%)$, P<0.05).

Discussion

We attempted to define the functions of TIL in human oesophageal cancer by their phenotypic characteristics. The present study is unique in two ways compared with previous reports on the phenotyping of TIL (Shimokawa et al. 1982; Hiratsuka et al. 1984; Goettlinger et al. 1985; Hurlimann and Saraga 1985). One is the application of two-colour analysis with flow-cytometry, and the other is the

comparison made using IEL and PBL as controls in the same patients. The use of two-colour analysis with flow-cytometry was designed to bring us quantitative and detailed information about the phenotypes in the TIL population. The comparison with controls was performed to clarify shifts of the phenotypic make-up of the TIL from the original phenotypes which might be closer to the PBL and/or IEL phenotypes.

We were clearly able to show the characteristics of the phenotypes in the TIL population. With single-colour analysis, we showed a predominance of T cells and a significant reduction of the frequency of natural killer (NK) cells among TIL compared with those in the controls. With twocolour analysis using the appropriate pairs of antibodies, we also showed that the frequencies of (helper CD4⁺Leu8⁻ T cells; Th) CD8⁺CD11b⁻ cells (cytotoxic T cells; Tc) in the TIL population were significantly greater than in the control populations. These results strongly suggest that T cells have a greater role in TIL functioning than do NK cells, and that Th and Tc might play a leading role in TIL activities. Since both Th and Tc are indispensable components of cellular immunity (Cantor and Boyse 1977; Cantor and Gershon 1979), these results indicate that TIL in human oesophageal cancer may be an expression of antigen-specific cellular immunity against the tumours. The potential of TIL for use in adoptive immunotherapy is also suggested. The increased numbers of TIL per weight of wet tissue compared to IEL also support this proposition.

We have shown phenotypic evidence for the first time in human oesophageal cancer that TIL might be expressing a specific anti-tumour response of the cancer-bearing host and might have a potential value for use in adoptive immunotherapy. The controversy caused by previous studies using the immunohistochemical staining of surface markers may be clarified to some degree by the present study.

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