

at is yet to be clarified whether CD16⁺ monocytes are differentiated from CD16⁻ monocytes or not.

Multiple factors are reported to enhance CD16 expression on monocytes, such as

macrophage-colony-stimulating factor (M-CSF), IL-10 and transforming growth factor (TGF)-\(\begin{align*} 1 \)

23). In addition, some studies disclosed that hypoxia and MIF contributed to TIE2 expression on

monocytes in vitro (14, 24). Cumulative data have been published showing that cancer cells,

including HCC, are dichotomously capable of releasing various inflammatory (TNF-α, IL-1β) and

anti-inflammatory (TGF-β and IL-10) cytokines, as well as hematopoietic factors (M-CSF and MIF)

(7,25-29). However, in this study, no correlation was found between the frequency of TEMs and any

of serum angiogenesis factors (Supplementary figure 3). It is thus plausible that, not a sole but a

mixture of such HCC-derived factors contributes to the generation of TEMs from CD16⁺ monocytes

with the aid of a hypoxic microenvironment. In our preliminary study, a combination of cytokines

and growth factors were able to induce TIE2 on CD16⁺ monocytes (manuscript in preparation).

Although the presence of TEMs has been reported in cancer tissues from patients with colorectal,

pancreatic or renal cancer (15), the actual roles of TEMs in the angiogenesis process of HCC have

yet to be elucidated. In this study, we showed that the peripheral and intrahepatic frequency of TEMs

is positively correlated with the density of micro-vessels in the liver. Additionally, we found that

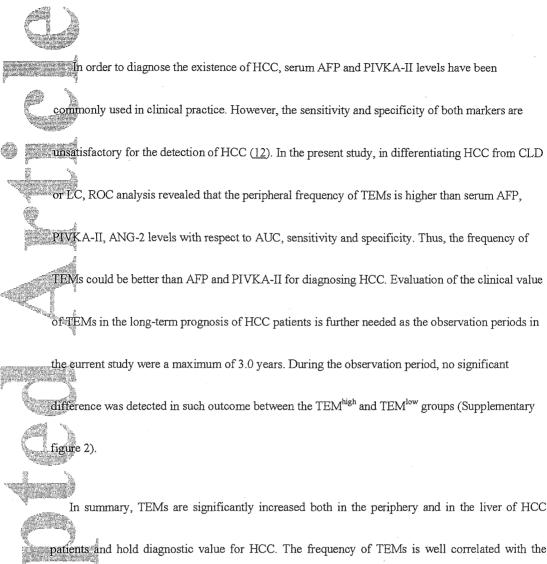
TEMs accumulate in the liver and are located in the perivascular area of HCC tissues. In support for

our observations, Venneri et al. reported that TEMs preferentially localize in the vicinity of tumor



blood vessels and avascular viable areas in human cancer specimens but are not found in non-neoplastic tissues adjacent to tumors (15). These results suggest that TEMs may take some part in HCC-related vascular formation. Further investigation is still needed to disclose the molecular mechanisms of TEMs accumulating in the liver. In analogy of CD14⁺CD16⁺ monocytes, some interactions between chemokine and its receptors, such as CCR5 or CX3CR1, may be involved in such process (22), all of which are shown to be expressed in TEMs as well (Fig. 1D).

Thus far, many studies have reported that the histological degree of angiogenesis in the liver is closely correlated with the prognosis or survival of patients with HCC. In agreement with these results, several serum angiogenesis factors, such as VEGF, ANG-2, sVEGFR-1 and MIF, are reported to be feasible as markers of prognosis, invasiveness or post-therapeutic recurrence in HCC patients (7-11). Multivariate analysis disclosed that the degree of angiogenesis in the liver, as assessed by MVD, is an independent factor significantly involved in the disease-free survival rate in patients with resectable HCC (17) Therefore, the positive correlation between peripheral and intra-hepatic TEM frequency and MVD in the liver observed in this study offers the possibility of TEMs being as a prognostic marker of HCC patients. In support for this, the patients with higher TEM frequency were at a more advanced Child-Pugh stage, had poorer liver function and showed higher post-therapeutic HCC recurrence in this cohort.



patients and hold diagnostic value for HCC. The frequency of TEMs is well correlated with the degrees of neo-vascularization in HCC tissue. Thus, TEMs would be feasible as a diagnostic cellular marker of HCC potentially reflecting angiogenesis in the liver.





Figure Legends



Fig. 1: Identification and phenotypic analyses of TEMs as CD14⁺CD16⁺TIE2⁺ cells in the

peripheral blood.

A. PBMC obtained from HCC patients were stained and analyzed by flow cytometry. CD14⁺

monocytes were divided into two distinct subsets, CD14⁺⁺CD16⁻ and CD14⁺CD16⁺ cells. These cells

were examined for TIE2 expression. The numbers in the histograms depict the percentages of gated

cells. Representative plots from three patients are shown.

B Frequencies of TIE2 in CD14⁺ monocytes were compared among two distinct subsets; CD16⁺

monocytes and CD16 monocytes, See Fig.1A. The bars indicate mean ± SE of 89 patients. *:

p≤0.0001 by Mann-Whitney nonparametric U test.

C. Western-blot analysis of TIE2 expression in FACS-sorted CD14⁺CD16⁺ and CD14⁺⁺CD16⁻ cells

from HCC patients. As shown, the bands were TIE2 (140 kDa molecular weight) (the top panels)

and those are β-actin (45 kDa molecular weight) (the bottom panels), respectively. The

representative results of three series of experiments from 7 HCC patients are shown.

DIFEMs and TIE2 monocytes in the periphery were gated and analyzed for the expression of

warious molecules, as indicated in the histogram plots. The filled light gray line, black line and gray

line depict the negative control, the expression of relevant markers in TEMs and TIE2 monocytes,



respectively. The percentage of the marker-positive cells is shown in the histograms. The upper

numbers are for TEMs and the lower ones are for TIE2 monocytes. The representative plots of six

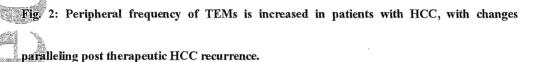
series of experiments are shown.

E. Comparative analyses of the expression of CCR4, CCR5, CX3CR1, CD40 and CD86 between

TEMs and TIE2 monocytes, assessed by FACS as described above. The bars indicate mean \pm SE of

six series of experiments. *: p<0.05, **: p<0.005, ***: p<0.001 by Mann-Whitney non-parametric U

test



A. Frequencies of peripheral TEMs in CD14⁺ monocytes are shown in four groups of HCV-positive

patients; HS, CH, LC and HCC, see Table.1. *: p<0.0001 by Kruscal-Wallis test with Dunn's

multiple comparison test.

B. Frequencies of peripheral TEMs in HCC patients at different clinical TNM stages (early stage;

IHI; n=70, advanced stage; III+IV; n=19). n.s., not significant by Mann-Whitney nonparametric U

test

C. In patients who underwent RFA therapy or resection of HCC, the frequencies of TEMs in CD14⁺

monocytes were examined serially after confirmation of complete ablation or operation to remove



HEC lesions. The bold arrows depict the time point of the RFA therapy or the operation. Both panels

depict the frequencies of TEMs in patients before and after the treatment. The left panel shows the

results of TEMs in patients without HCC recurrence (n=12) as assessed by CT/MRI examinations,

while the right panel shows the results in patients with HCC recurrence (n=5), respectively. *:

p≤0.05 by Paired t-test.

D. In patients with HCC who underwent RFA or the operation, the recurrence-free survival rate after

the treatment was compared between those with TEM^{high} (frequency of TEMs ≥ 2.75; n=45) and

TEM of requency of TEMs < 2.75; n=44) using the Kaplan-Meier method, with the log-rank test

for comparison. TEM^{high} and TEM^{low}, see Table 2. p = 0.047.



in HCC tissue than in the periphery.

A. Immunofluorescence staining was performed as described in Materials and Methods. The staining

for CD14 (green), CD16 (red) and TIE2 (red) identifies CD14⁺TIE2⁻, CD14⁺TIE2⁺, CD14⁺CD16⁻

eells and CD14⁺CD16⁺ monocytes in human liver tissue (blue: nuclei counterstained with Dapi).

Representative results of the resected samples obtained from 12 HCC patients are shown. The panels

show CD14⁺TIE2⁺ cells (A, B, D) and CD14⁺CD16⁺ cells (C) in the perivascular area of HCC tissue

magnification, 400×), Bold arrows depict CD14 TIE2 cells, thin arrows CD14 CD16 cells, bold



arrowheads vascular endothelial cells (TIE2⁺CD14⁻) and thin arrowheads CD14⁺TIE2⁻ cells.

B. The frequencies of TEMs in 9 patients with HCC are compared for those in the peripheral blood or in the liver. Liver-infiltrating leukocytes are divided into two distinct groups: leukocytes infiltrating non-tumor tissue (NIL) and tumor-infiltrating leukocytes (TIL). These cells were stained using anti-human CD14, CD16 and TIE2 mAbs. The analyses were performed as described in Materials and Methods. The samples were obtained from nine patients who underwent tumor resection. *: p<0.05, **: p<0.0005, by Paired t-test.

C. The correlation is analyzed between the frequencies of TEMs in PBMC and those in tumor-infiltrated lymphocytes. Analysis (n=14) was based on Pearson's correlation coefficient. P = 0.003, $R^2 = 0.53$

Fig_4: The degree of microvessels in HCC is correlated with the frequency of TEMs

A. Assessment of neo-vascularization in HCC tissue was performed by staining CD34⁺ cells in the resected samples obtained from patients with HCC. Immunohistochemical staining for CD34 was done as described in Materials and Methods. The CD34⁺ cells were mainly confined to the cytoplasm of vascular endothelial cells as brownish yellow granules. Microvessels were represented by brownish yellow capillaries or small cell clusters. Representative results are shown. The panel shows a turnor cell area of a high grade.



B. The correlation is analyzed between TEM frequencies and the counts of CD34⁺ cells (MVD

values) in relevant patients. The panels show the correlation of peripheral (left: n=11) or

intra-tumoral TEM frequencies (right: n=9) and MVD (p = 0.0009, R^2 = 0.72 and p = 0.04, R^2 = 0.44,

respectively). Analysis was based on Pearson's correlation coefficient.

Fig. 5: Frequency of TEMs is superior as a diagnostic for HCC to common tumor markers or angiogenesis factor.

A. Correlation between TEM frequency and AFP (n=87) or PIVKA-II (n=81) was analyzed using

Pearson's correlation coefficient. P = 0.45, $R^2 = 0.007$ or P = 0.27, $R^2 = 0.02$, respectively.

BROC analyses were performed in order to assess the diagnostic value of TEM frequency for

differentiating HCC (n=89) from chronic liver disease (CLD, n=79) or liver cirrhosis (LC, n=30).

The left panel shows the diagnostic value of TEMs, PIVKA-II, AFP and angiopoietin-2 (ANG-2) for

HCC from CLD and the right panel shows those for HCC from LC, respectively.

Supplementary figure 1: Frequency of TEMs is comparable in HCC patients regardless of

étiology.

The frequencies of peripheral TEMs are shown for HCC patients with or without HCV infection

(n=89 and n=26, respectively). The group without HCV infection included patients with HBV



infection or those without HBV nor HCV (HBV-HCC and Non-B, Non-C [NBNC] HCC patients).

n.s., not significant by Mann-Whitney non-parametric U test.

Supplementary figure 2: Peripheral frequency of TEMs is not related to the overall survival in

patients with HCC.

In patients with HCC who underwent RFA or the operation, the overall survival rate after the treatment was compared between those with TEM^{high} (frequency of TEMs \geq 2.75; n=45) and

TEM^{low} (frequency of TEMs < 2.75; n=44) using the Kaplan-Meier method, with the log-rank test

for comparison. TEM^{high} and TEM^{low}, see Table 2. P=0.36

Supplementary figure 3: Correlation between the frequency of TEMs and serum levels of

angiogenesis factors

All plots indicate the correlation between TEM frequency and VEGF, ANG-2, sVEGFR-1 or MIF.

The numbers of patients examined were 32, 37, 23 and 31, respectively. Analyses were based on

Pearson's correlation coefficient.





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Table.1: The clinical backgrounds of the subjects

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	Clinicopathologic characteristics	СН	LC	HCC
	Gender: Male/female	21/28	13/17	63/26
	Age: mean±SD	63.4±7.8	67.5±8.8	70.0±7.0
The state of the s	Alanine aminotransferase (IU/L)	56.1±43.5	54.0±28.6	46.3±29.5
	Prothrombin time (%)	91.5±14.8	73.4±9.7	77.7±15.4
The state of the s	Platelet (×10 ⁴ /mm ³)	15.8±5.0	8.6±3.7	11.5±5.7
	Albumin (g/dl)	4.0±0.3	3.5±0.4	3.5±0.5
	Total bilirubin (mg/dl)	0.7±0.3	0.9±0.4	0.9±0.6
	- Child-Pugh grade: A/B	49/0	21/9	59/30
	a-fetoprotein (ng/ml)	8.7±10.7	42.6±80.6	264.0±1281.7
	TNM stage: I/II/III/IV			33/37/15/4
	BCLC stage: A/B/C/D			44/13/26/6

TNM, Tumor, Lymph Node and Metastasis; BCLC, Barcelona-Clinic Liver Cancer; CH, chronic

hepatisis; LC, liver cirrhosis; HCC, hepatocellular carcinoma





Table.2: The comparison of clinical parameters of HCC patients between those with higher

frequency of TEMs and those with lower frequency

——————————————————————————————————————	istics	Peripher		
		High (n=45)	Low (n=44)	Р
Gender: Male/female		32/13	31/13	1.000 [†]
Age: mean±SD		71.4±6.6	68.7±7.3	0.065 [‡]
Child-Pugh grade: A/B	NOW THE PROPERTY OF THE PROPER			0.043^{\dagger}
MELD score		9.1±2.3	8.2±1.8	0.010 [§]
α-fetoprotein (ng/ml)	α-fetoprotein (ng/ml)			0.1018
TNM stage: I+II/III+IV		35/10	35/9	1.000 [†]
BCLC stage: A+B/C+D		25/20	32/12	0.123 [†]
Vascular invasion: present/a	bsent	4/41	2/42	0.677 [†]
Tumor size: <3/>3/cm		36/9	34/10	0.800 [†]
Tumor number: single/multi	ple	21/24	25/19	0.399 [†]
Alanine aminotransferase (I	U/L)	51.5±34.5	41.0±22.6	0.251§
Prothrombin time (%)		73.3±14.5	82.6±15.0	0.004\$
Platelet (×10 ⁴ /mm ³)	Platelet (×10 ⁴ /mm ³)		12.2±6.2	0.319 [§]
Albumin (g/dl)	Albumin (g/dl)			0.045 [§]
Total bilirubin (mg/dl)		0.9±0.6	0.9±0.5	0.820 [§]

P value of less than 0.05 was expressed by thick one.

 $^{^{\}ddagger}$ Student's ttest, $^{\dagger}\chi 2$ test or Fisher exact test, $^{\$}$ Mann-Whitney Utest.



TEMs, TIE2-expressing monocytes; MELD, model for endstage liver disease

Table.3: The assessment of diagnostic value of the TEMs, AFP and PIVKA-II by ROC analyses

Discrimination between	Parameter	AUC	Cut off value	Sensitivity (%)	Specificity (%)	Positive predictive values (%)	Negative predictive values (%)
HCC and CLD	TEMs	0.85	1.46	86.1	71	76.3	82.4
	AFP	0.69	10	65.9	43.9	65.9	56.1
	PIVKA-II	0.77	47	56.3	96.5	97.8	44.5
	ANG-2	0.62	2681	51.4	67.3	52.8	64
HCC and LC	TEMs	0.93	2.75	81.3	90	89.7	81.8
	AFP	0.61	10	71.9	32.1	54.8	50
	PIVKA-II	0.79	39	64.5	93.3	95.2	56
	ANG-2	0.57	2440	68.8	52	47.8	. 72.2

The optimal cut-off point was determined as those yielding the minimal value for (1-sensitivity)² +

(1 specificity)². Such point with those sensitivity and specificity values is the closest to (0, 1) point

on ROC curve. .

AUC, Area under the curve; TEMs, \underline{T} IE2-expressing \underline{m} onocytes; CLD, chronic liver disease; LC,

liver cirrhosis; ANG-2, angiopoietin-2





Supplementary Table.1: The inclusion and exclusion criteria for the enrollment of patients

Eligible criteria for patients with hepatocellular carcinoma (HCC)

- 1) PS: 0~2 (ECOG PS)
- 2) Age: 40 80 years
- 3) Viable HCC was diagnosed by imaging analyses, such as CT and/or MRI.
 - Histological diagnosis is optional depending on the availability for tissue samples.
- 4) De novo as well as recurrent HCC are included
- 5) Patients who were tolerable for trans-catheter arterial chemo-embolization therapy,

local ablation therapy and/or surgical resection of the tumor

Eligible criteria for patients with chronic liver disease

- 1) PS: 0~2 (ECOG PS)
- 2) Age: 40 80 years
- 3) The presence of HCC was excluded by imaging analyses
- 4) Chronic hepatitis or liver cirrhosis was clinically diagnosed by biochemical and imaging examinations

Exclusion criteria for patients

- 1) Pregnant women
- 2) Patients with bleeding tendency (%prothrombin time<50%, platelet counts<30000/mm³)
- 3) Patients with autoimmune hepatitis, primary biliary cirrhosis or primary sclerosing cholangititis
- 4) Patients with autoimmune diseases in any organs
- 5) Patients whom the attending doctors determined as ineligible
- PS, Performance status; ECOG, Eastern Cooperative Oncology Group

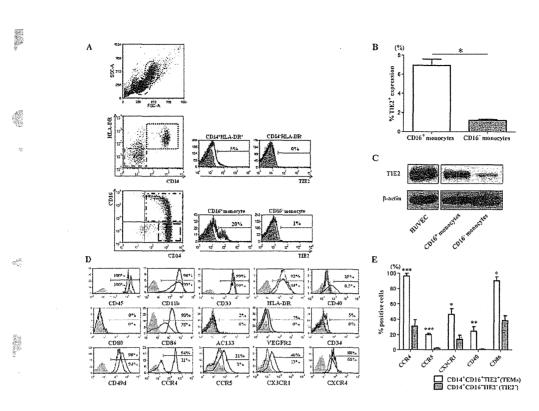


Fig. 1: Identification and phenotypic analyses of TEMs as CD14+CD16+TIE2+ cells in the peripheral blood.

A. PBMC obtained from HCC patients were stained and analyzed by flow cytometry. CD14+ monocytes were divided into two distinct subsets, CD14++CD16- and CD14+CD16+ cells. These cells were examined for TIE2 expression. The numbers in the histograms depict the percentages of gated cells. Representative plots from three patients are shown.

B. Frequencies of TIE2 in CD14+ monocytes were compared among two distinct subsets; CD16+ monocytes and CD16- monocytes, See Fig.1A. The bars indicate mean \pm SE of 89 patients. *: p<0.0001 by Mann-Whitney nonparametric U test.

C. Western-blot analysis of TIE2 expression in FACS-sorted CD14+CD16+ and CD14++CD16- cells from HCC patients. As shown, the bands were TIE2 (140 kDa molecular weight) (the top panels) and those are β -actin (45 kDa molecular weight) (the bottom panels), respectively. The representative results of three series of experiments from 7 HCC patients are shown.

D. TEMs and TIE2- monocytes in the periphery were gated and analyzed for the expression of various molecules, as indicated in the histogram plots. The filled light gray line, black line and gray line depict the negative control, the expression of relevant markers in TEMs and TIE2- monocytes, respectively. The percentage of the marker-positive cells is shown in the histograms. The upper numbers are for TEMs and the lower ones are for TIE2- monocytes. The representative plots of six series of experiments are shown.

E. Comparative analyses of the expression of CCR4, CCR5, CX3CR1, CD40 and CD86 between TEMs and TIE2- monocytes, assessed by FACS as described above. The bars indicate mean ± SE of six series of experiments. *: p<0.05, **: p<0.005, **: p<0.001 by Mann-Whitney non-parametric U test. 254x190mm (300 x 300 DPI)



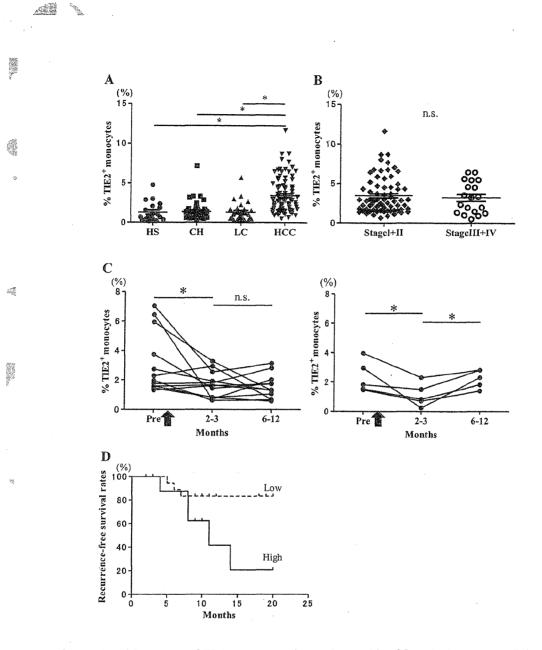


Fig. 2: Peripheral frequency of TEMs is increased in patients with HCC, with changes paralleling post therapeutic HCC recurrence.

A. Frequencies of peripheral TEMs in CD14+ monocytes are shown in four groups of HCV-positive patients; HS, CH, LC and HCC, see Table.1. *: p<0.0001 by Kruscal-Wallis test with Dunn's multiple comparison test. B. Frequencies of peripheral TEMs in HCC patients at different clinical TNM stages (early stage; I+II; n=70, advanced stage; III+IV; n=19). n.s., not significant by Mann-Whitney nonparametric U test. C. In patients who underwent RFA therapy or resection of HCC, the frequencies of TEMs in CD14+ monocytes were examined serially after confirmation of complete ablation or operation to remove HCC lesions. The bold arrows depict the time point of the RFA therapy or the operation. Both panels depict the frequencies of TEMs in patients before and after the treatment. The left panel shows the results of TEMs in patients without HCC recurrence (n=12) as assessed by CT/MRI examinations, while the right panel shows the results in patients with HCC recurrence (n=5), respectively. *: p<0.05 by Paired t-test.

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190×254mm (300 x 300 DPI)

