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Relation	



Lymphocyte subset characterization associated with persistent hepatitis C virus infection and subsequent progression of liver fibrosis

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Abbreviations:

AHS, Adult Health Study; ALT, alanine aminotransferase; AST, aspartate aminotransferase; γ -GTP, γ -glutamyltransferase; Gy, gray; HCV, hepatitis C virus; RERF, Radiation Effects Research Foundation

Abstract

This study aims to deepen understanding of lymphocyte phenotypes related to the course of hepatitis C virus (HCV) infection and progression of liver fibrosis, in a cohort of atomic-bomb survivors. The study subjects comprise three groups: 162 HCV persistently infected, 145 spontaneously cleared, and 3511 uninfected individuals. We found increased percentages of peripheral blood T_{H1} and total CD8 T cells and decreased percentages of NK cells in the HCV persistence group, compared with the other two groups, after adjustment for age, gender, and radiation exposure dose. Subsequently, we found that increased T_{H1} cell percentages in the HCV persistence group were significantly associated with an accelerated time-course reduction in platelet counts—accelerated progression of liver fibrosis—while T_{C1} and NK cell percentages were inversely associated with the progression. This study suggests that T_{H1} immunity is enhanced by persistent HCV infection, and that percentages of peripheral T_{H1} , T_{C1} , and NK cells may help predict progression of liver fibrosis.

Keywords:

Cohort study; hepatitis C virus; liver fibrosis; lymphocyte subset

Abbreviated title:

Lymphocyte subsets and HCV-infected persons

1. Introduction

HCV infects some 120-170 million people worldwide, and persistent HCV infection is a major cause of liver diseases including chronic hepatitis, cirrhosis, and hepatocellular carcinomas [1, 2]. It is now widely recognized that both innate and adaptive arms of the host immune system are closely involved in persistent infection, liver injury, and virus clearance [3, 4]. For instance, cytotoxic granule release and cytokine production of NK cells are inhibited by direct binding of HCV envelope protein E2 to CD81 on NK cells, or stabilizing the HLA-E expressions on hepatocytes in HCV-infected patients [5, 6]. However, comprehensive understanding of interactions between HCV and the immune system remains incomplete [3, 4]. Moreover, aging, gender, and several environmental factors such as alcohol drinking, smoking, and ionizing radiation have been reported to influence host immune functions as well as HCV spontaneous clearance [7-9], which may increase the complexity of virus-host interactions. Therefore, a comprehensive characterization of host immunological phenotypes in HCV infection is needed, especially with a cohort-based study design without conceivable selection bias [10]. Nevertheless, few studies along those lines have been carried out. One prospective cohort study (Adult Health Study) of atomic-bomb survivors-a longevity cohort with biennial health examinations-has been conducted at the Radiation Effects Research Foundation (RERF), provides and the study clinicoepidemiological data related to HCV infection and immunological status [11, 12].

Within the cohort study, we conducted a cross-sectional analysis for peripheral blood lymphocyte subsets among HCV persistently-infected, spontaneously-cleared, and uninfected groups, aiming to delineate immunological distinctions among these three groups. We also aimed to identify the lymphocyte subsets that can predict hepatitis progression in HCV-persistent individuals, on the basis of a longitudinal analysis of time-course changes of platelet counts.

2. Materials and methods

2.1. Study population

The Atomic Bomb Casualty Commission, subsequently the RERF, established the Adult Health Study (AHS) cohort in 1958. This cohort study enrolled a total of 23,000 atomic-bomb survivors in Hiroshima and Nagasaki, who biennially received health examinations in outpatient clinics [11]. Hepatitis screening (HBsAg, anti-HBc Ab, anti-HBs Ab, and anti-HCV Ab tests, as well as HCV RNA test if anti-HCV Ab was positive) was conducted among 6,121 AHS participants in 1993 – 1995 [12]. Anti-HCV Ab negative subjects were categorized as the HCV-uninfected group in this study, whereas a "persistence" group was identified by anti-HCV Ab positive with detected HCV RNA, and a "spontaneous clearance" group was identified by anti-HCV Ab positive and undetectable HCV RNA. Subjects with hepatitis B virus surface antigen-positive were excluded from this study. From the 6,121 AHS subjects, lymphocyte subsets in the peripheral blood were then examined in 162 HCV persistence, 145 virus clearance, and 3511 uninfected subjects in 2000 Most subjects (N = 120, 74%) in the persistence group (N = 162, including those -2002.with cancer history) were confirmed by a second RNA test at least two years after the first RNA test performed in 1993 – 1995. Although the remaining 42 subjects in the group did not undergo the second RNA test, these subjects were confirmed to have developed type C chronic liver disease, based on medical chart review (e.g., treatment history, abdominal

sonographic observation, changes in platelet count, zinc sulfate turbidity, AST, and ALT, between 1993 – 1995 and 2000 – 2002) by a hepatologist (one of the authors, WO). Subsequent treatment data of hepatitis C from attending physicians were also taken into account. No subjects in the persistence group underwent IFN therapy in 2000 – 2002.

This study was approved by the RERF Human Investigation Committee, and all subjects gave written informed consent before each examination.

2.2. Assays in hepatitis screening and clinical examinations

In 1993 – 1995, anti-HCV Ab and hepatitis B virus surface antigen were examined using a second-generation passive hemagglutination kit and a reverse passive hemagglutination kit (Dynabott, Tokyo), as described previously [12]. Subjects were diagnosed as having Ab when agglutination was found in a serum diluted 2⁵. Qualitative and quantitative detection of HCV RNA was carried out using the Amplicor HCV ver. 2.0 and the Amplicor HCV monitor test ver. 1.0 and/or ver. 2.0 (Roche Diagnostics Systems, Tokyo, Japan).

Platelet count decreases with progression of liver fibrosis, and this marker has widely been used as a reliable diagnostic tool for liver fibrosis/cirrhosis in patients with chronic HCV infection [13-16]. Postulated mechanisms for such platelet reduction include decreased secretion of the hematopoietic growth factor thrombopoietin from the liver and increased destruction of platelets by antiplatelet antibodies [17, 18]. Platelet count was routinely measured in the AHS health examination, and an automatic blood cell counter (Coulter MAXM, Beckman Coulter, Inc. Tokyo, Japan) was used in 2000 – 2002. Aspartate aminotransferase (AST), alanine aminotransferase (ALT), γ -glutamyltransferase (γ -GTP), and total cholesterol were also routinely measured, and an autoanalyzer (Hitachi 7180, Hitachi, Ltd., Tokyo, Japan) was used in 2000 – 2002.

2.3. Information on lifestyle/environmental factors and clinical data

Information on alcohol drinking and smoking was obtained from questionnaires at the time of the AHS health examination in 1993 – 1995 and 2000 – 2002, respectively. Body mass index (BMI) was measured at the AHS health examination in 2000 – 2002. Radiation dose was estimated by the DS02 dosimetry system [19], based on the weighted skin dose computed as the gamma dose plus 10 times the neutron dose. No subjects were diagnosed with HIV infection. No subjects underwent organ transplantation or immunosuppressive therapy. Clinical information was obtained at the AHS examination in 2000 – 2002 as well as medical chart review, and classified according to the International Classification of Diseases (ICD) code.

2.4. Lymphocyte subset analysis

Circulating T_H1 and T_H2 cells can be straightforwardly enumerated by flow cytometry,

using cell surface markers for chemokine receptor, CXCR3, and prostaglandin D receptor, CRTH2, respectively [20, 21]. CD8 T cells expressing CXCR3, known as $T_{C}1$, are also involved in the viral control during HCV infection [22]. We thus focused on $T_{H}1$, $T_{H}2$, $T_{C}1$ and $T_{C}2$ cell subsets, as well as total CD4 T and CD8 T, NK, and B cell subsets in relation to HCV infection status.

Analytical flow cytometry was conducted in a FACScan machine (BD Biosciences, San Jose, CA), as described previously [23]. Monoclonal antibodies as specific cell surface markers were purchased from BD Pharmingen (San Diego, CA), unless otherwise noted. CD4 or CD8 T cells were enumerated as PerCP-labeled CD3 positive and PE-CD4 or FITC-CD8 positive cells; CD16 or CD20 cells were enumerated as PerCP-CD3 negative and FITC-CD16 (Beckman Coulter, Brea, CA) or PE-CD20 positive cells. We used CXCR3 as a marker for T_H1 and T_C1 cells [20, 22] and CRTH2 for T_H2 and T_C2 cells [21]. Namely, T_H1 and T_H2 cells were identified with PerCP-CD4, FITC-CXCR3 (R&D Systems, Minneapolis, MN), and biotinylated CRTH2 (kindly provided by Dr. K. Nagata, BML, Kawagoe, Japan) plus PE-streptavidin; T_C1 and T_C2 cells were identified with PerCP-CD8, FITC-CXCR3, and biotinylated CRTH2 plus PE-streptavidin. In every measurement, approximately 20,000 cells were analyzed.

2.5. Statistical analysis

Two sample Wilcoxon or Pearson chi-square tests were performed to compare distributions of age, gender, city, radiation dose (Gy), smoking (packs/day), alcohol drinking (converted to grams of ethanol/day), BMI (kg/m²), AST (IU/L), ALT (IU/L), γ -GTP (U/L), total cholesterol (mg/dL), and platelet count (×10⁴/µL) between all combinations of the three groups.

Since aging and past radiation exposure likely influenced various immunological markers [23], these events were also evaluated in this study. In each study group, the associations of lymphocyte subsets with age (at the time of examination), gender, radiation dose, and city were evaluated based on the multiple regression model [24]:

log (subset percentages or ratios) = $\alpha + \beta_1 \times age + \beta_2 \times gender + \beta_3 \times dose + \beta_4 \times city + \beta_5 \times alcohol + \beta_6 \times smoking + \beta_7 \times BMI + \beta_8 \times autoimmune disease + \beta_9 \times allergic disease + \beta_{10} \times cancer + \beta_{11} \times other non-cancer diseases$

where log is the logarithm at base 10, gender = 0 for male and 1 for female, city = 1 for Hiroshima and 2 for Nagasaki. Smoking, alcohol drinking, BMI, autoimmune disease (1 if diagnosed, otherwise 0), allergic disease (1 or 0), cancer (1 or 0), and other non-cancer diseases (i.e., hypergammaglobulinemia and sarcoidosis, 1 or 0) were also used as additional explanatory variables.

We compared lymphocyte subset percentages or ratios between all combinations of the three groups in normal regression analysis with adjustment for age, gender, radiation dose, city, alcohol, smoking, BMI, autoimmune diseases, allergic diseases, and other non-cancer diseases: In the regression analysis, an explanatory variable regarding a group (one group = 0, another group = 1) was used.

Regression analysis was also performed to investigate if there was any association between subset percentages or ratios and time-course changes in platelet counts through the period from 2000 through 2006. Changes in platelet counts were calculated by:

(platelet counts at the last examination – platelet counts at the first examination) / follow-up years.

In a regression analysis, a forward step-wise procedure was used for 8 immunological variables, %CD4, T_H1 , T_H2 , CD8, T_C1 , T_C2 , CD16, and CD20. Four variables (% T_H1 , T_C1 , CD16, and CD20) were consequently selected (significance level to select, P < 0.2) to construct a statistical model. All analyses were conducted using Stata software (Stata/SE 9.2 for Windows, StataCorp LP, College Station, TX).

3. Results

3.1. Basic characteristics of study subjects

Table 1 compares characteristics of study subjects in the HCV persistence, clearance, and uninfected groups. The persistence group showed increased levels of blood γ -GTP, and decreased levels of total cholesterol and platelet counts, compared with the other two groups, indicating enhanced liver injury by persistent HCV infection. The proportion of Hiroshima subjects in the persistence or clearance group was higher than that in the uninfected. This is in accordance with the previous study that showed a higher anti-HCV Ab prevalence in Hiroshima atomic-bomb survivors than in Nagasaki survivors [12]. There were no significant differences in radiation dose by HCV infection status.

This study primarily aimed to evaluate immunological alterations associated with HCV infection that might be modulated by age, gender, or past radiation exposure. Therefore, the effects of those factors on lymphocyte subsets were first analyzed and are summarized in Supplemental Tables. In the uninfected group, we found: i) age- and dose-dependent decreases in total CD4 T cell percentages, ii) higher total CD4 T cell percentages in females than in males, iii) no significant effects of age, gender, or radiation dose on total CD8 T cell percentages, iv) increased CD16 (NK) cell percentages with increasing age and higher percentages in males than in females, v) increased T_H1 and T_H2 cell percentages with increasing age and dose, and vi) both $T_H 1/T_H 2$ and $T_C 1/T_C 2$ cell ratios negatively or positively

associated with age and the female gender, respectively, but not with radiation dose (Supplemental Table 1). The persistence and clearance groups also showed similar associations, although most associations were not statistically significant, probably due to smaller numbers of subjects in these groups (Supplemental Tables 2 and 3). In addition to the three factors (age, gender, and radiation), other selected factors such as city, alcohol, smoking, and BMI also influenced various lymphocyte subsets (data not shown), and they were used as confounding variables in adjustments.

3.2. Comparison of lymphocyte subsets among the HCV persistence, clearance, and uninfected groups

The study subjects having cancer history numbered 60, 32, and 698 in the persistence, clearance, and uninfected groups, respectively. We then analyzed the lymphocyte subset alterations associated with HCV infection among subjects who have no history of cancer shown in Table 2, so as to eliminate potential effects of cancer development and/or cancer therapy (Table 3). In addition, basic characteristics were not largely changed by excluding subjects with a history of cancer, but the radiation effects—specifically on T_H1 and T_H2 cells—were no longer seen among subjects with no cancer history (data not shown).

In the persistence group, T_H1 and total CD8 T cell percentages and T_H1/T_H2 ratios were significantly higher than those in the HCV-uninfected group, while total CD4 T and CD16

cell percentages were lower. Similar differences were also seen between the persistence and clearance groups. However, except for total CD8 T cell percentages, no significant differences were observed between the clearance and uninfected groups.

3.3. Relationship between lymphocyte subsets and progression of liver fibrosis in the HCV persistence group

Next, we analyzed the relationship between lymphocyte subsets and platelet counts that had been longitudinally examined at the biennial AHS examination during 2000 – 2006 in the HCV persistence group, excluding subjects with cancer history (Table 4). Average follow-up period was 4.7 years, and average decrement of platelet counts per year was -0.75 (×10⁴/µL). We found that increased percentages of T_H1 cells were associated with accelerated time-course reduction in platelet counts—accelerated progression of liver fibrosis (P = 0.027)—while T_C1 and NK cell percentages were inversely associated with the progression (P = 0.027 and 0.058, respectively).

4. Discussion

We investigated immunological alterations associated with HCV infection in a longevity study cohort of atomic-bomb survivors. First, the effects of age, gender, and radiation on total CD4 T, CD8 T, and NK cells were studied in the uninfected group (Supplemental Table 1) and found to be in close agreement with our previous studies [9, 23]. A new finding related to the uninfected group is that percentages of both T_{H1} and T_{H2} cells increased with increasing radiation dose and age. That result is consistent with our previous findings in an expanded cohort of atomic-bomb survivors, which showed age- and radiation dose-dependent elevations of cytokine levels for both T_{H1} -related cytokines (IFN- γ and TNF- α) and a T_{H2} -related cytokine (IL-6) [25].

Second, we found that persistent HCV infection was associated with increases in T_H1/T_H2 cell ratios and CD8 T cell percentages, and a decrease in NK cell percentages (Table 3). Regarding cytokine responses to persistent HCV infection, past reports in diversified patient groups were rather inconsistent: enhanced T_H1 responses [26-29], T_H2 responses [30-32], or both types [33, 34]. Although differing degrees of pathogenesis and/or inflammation among study patient groups may be in part responsible for this discrepancy [34, 35], some potential methodological drawbacks in studies demonstrating T_H2 cytokine predominance were indicated [27]. The present study on lymphocyte subsets suggested enhanced T_H1 immunity alone is not

sufficient to regulate the virus in many cases of HCV infection.

As observed in this study, both decreased and increased percentages of peripheral NK cells and total CD8 T cells, respectively, have been reported in HCV persistence individuals [36-39]. A reduction in NK cells is assumed to be linked to ongoing viremia that may induce continuous proliferation of CD8 T cells. A recent study on murine cytomegalovirus infection showed that NK cells negatively regulated the number and activity of virus-specific CD8 T cells as well as CD4 T cells that played a critical role in limiting viral persistence; lack of NK cell activation resulted in increased numbers of CD8 T and CD4 T cells along with enhanced effector functions through antigen-presentation by viral infected APCs [40]. Such functional interplay among NK cells, APCs, and CD8 T cells may be common features in virally infected hosts. It is also plausible that reduced NK cells of individuals may in part reflect their weakened natural immunity upon HCV infection, preferentially leading to failure of HCV-infected cell clearance [41].

Finally, our follow-up survey of the HCV persistence group showed that increased T_{H1} cell percentages were associated with accelerated progression of liver fibrosis, while T_{C1} and NK cell percentages were inversely associated with the progression (Table 4). In accordance with preceding studies [26, 28, 42], this study demonstrates that T_{H1} -immunity plays a vital role in HCV-related fibrosis progression. In the liver as well as peripheral blood of individuals with chronic HCV infection, T_{H1} cells may enhance CTL response and macrophage activation by producing cytokines such as IL-2, IFN- γ , and TNF- α , thereby facilitating the necroinflammatory process of hepatitis C [26, 42]. On the other hand, increased T_C1 cell percentages in total CD8 T cells were associated with slower progression of fibrosis—new findings in this study. T_C1 cells express a chemokine receptor, CXCR3, which is required for migration to the HCV-infected liver [22], and we inferred that the increased T_C1 fraction includes HCV-specific CD8 T cells. Several studies have shown relationships between higher numbers of circulating as well as intrahepatic HCV-specific CD8 T cells and lesser degrees of liver fibrosis during chronic HCV infection [43-46]: One plausible explanation is that HCV-specific CD8 T cells might control the virus without exerting cytotoxic effects on hepatocytes [46]. Alternatively, a population of CD8 T cells secreting an anti-fibrotic cytokine, IL-10, may be implicated in attenuation of hepatocyte killing and protection against liver injury [45, 47].

A limitation of this study is that we examined clinical and immunological data only from peripheral blood. Intrahepatic lymphocytes are assumed to have features discrete from those in peripheral blood [48]. Also, our comparison of lymphocyte subsets among study groups was cross-sectional, making it difficult to identify immunological factors responsible for persistent HCV infection.

In conclusion, this study identified immunological characteristics associated with HCV infection in a Japanese population and also indicated that peripheral T_H1 , T_C1 , and NK cell

subsets will be useful for predicting progression of hepatitis in persistently HCV-infected patients, and consequent development of hepatocellular carcinomas.

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Characteristics of the study subjects

	Persistence	Clearance	Uninfected			
	(anti-HCV +	(anti-HCV +	(anti-HCV -)	Persistence vs. Clearance	Persistence vs. Uninfected	Clearance vs. Uninfected
	/HCV RNA +)	/HCV RNA -)				
—	N=162	N = 145	N = 3511	P value ^a	P value ^a	P value ^a
Age ^b	71.7 (60.8-83.0)	72.3 (64.8-88.0)	72.2 (58.4-87.2)	0.041	0.79	0.029
Gender ^c						
Male	57 (35.2)	48 (33.1)	1078 (30.7)	0.70	0.23	0.54
Female	105 (64.8)	97 (66.9)	2433 (69.3)			
City ^c						
Hiroshima	112 (69.1)	96 (66.2)	2034 (57.9)	0.58	0.005	0.048
Nagasaki	50 (30.9)	49 (33.8)	1477 (42.1)			
Radiation dose (Gy) ^b	0.147 (0-2.658)	0.071 (0-1.890)	0.096 (0-2.032)	0.26	0.47	0.39
Smoking (pack/day) ^b	0 (0-1.0)	0 (0-1.0)	0 (0-1.0)	0.80	0.087	0.18
Alcohol drinking (gram/day) ^b	$0(0-85.0)^{d}$	$0(0-103.0)^{d}$	$0(0-69.8)^{d}$	0.35	0.57	0.062
BMI $(kg/m^2)^b$	22.4 (16.7-28.5)	22.6 (17.3-29.6)	22.7 (17.6-28.7)	0.46	0.084	0.58
AST (IU/L) ^b	21 (15-51)	21 (14-35)	22 (15-43)	0.15	0.85	0.081
ALT (IU/L) ^b	17 (9-46)	18 (9-42)	17 (9-44)	0.87	0.99	0.84
γ-GTP (U/L) ^b	28.5 (12.5-131.5)	25 (12-127)	24 (11-111)	0.10	< 0.001	0.29
Total cholesterol (mg/dL) ^b	168.5 (117-234)	206 (142-264)	208 (154-266)	< 0.001	< 0.001	0.41
Platelet count $(\times 10^4/\mu L)^b$	17.3 (7.2-28.2)	22.0 (14.3-30.8)	22.9 (14.5-33.7)	< 0.001	< 0.001	0.031

^aTwo-sample Wilcoxon rank-sum test, or Pearson chi-square test for gender and city.

^bMedian (5-95% percentiles).

^cNumber (%).

^dThe percentages of never-drinkers were 59.3, 53.8, and 60.6 in the 3 groups, respectively.

	Persistence	Clearance	Uninfected			
	(anti-HCV +	(anti-HCV +	(anti-HCV -)	Persistence vs. Clearance	Persistence vs. Uninfected	Clearance vs. Uninfected
	/HCV RNA +)	/HCV RNA -)				
	N =102	N = 113	N = 2813	P value ^a	P value ^a	P value ^a
Age ^b	71.5 (61.3-82.7)	72.3 (62.4-90.8)	72.0 (58.1-87.3)	0.026	0.64	0.021
Gender ^c						
Male	36 (35.3)	38 (33.6)	832 (29.6)	0.80	0.22	0.36
Female	66 (64.7)	75 (66.4)	1981 (70.4)			
City ^c						
Hiroshima	70 (68.6)	77 (68.1)	1613 (57.3)	0.94	0.023	0.023
Nagasaki	32 (31.4)	36 (31.9)	1200 (42.7)			
Radiation dose (Gy) ^b	0.031 (0-1.862)	0.056 (0-1.890)	0.072 (0-1.878)	0.84	0.68	0.49
Smoking (pack/day) ^b	0 (0-1.0)	0 (0-1.0)	0 (0-1.0)	0.67	0.070	0.18
Alcohol drinking (gram/day) ^b	$0(0-90)^{d}$	$0(0-105.8)^{d}$	$0(0-69.8)^{d}$	0.067	0.76	0.014
BMI (kg/m ²) ^b	22.6 (16.7-28.5)	22.6 (17.3-29.6)	22.9 (17.7-28.8)	0.49	0.097	0.55
AST (IU/L) ^b	22 (15-50)	21 (13-37)	22 (15-42)	0.12	0.34	0.26
ALT (IU/L) ^b	18 (9-39)	17 (9-48)	17 (9-44)	0.57	0.69	0.69
γ -GTP (U/L) ^b	26 (12-100)	27 (12-127)	23 (11-106)	0.99	0.19	0.18
Total cholesterol (mg/dL) ^b	175 (124-243)	211 (136-272)	209 (157-266)	< 0.001	< 0.001	0.46
Platelet count $(\times 10^4/\mu L)^b$	19.3 (10.3-32.0)	21.7 (14.2-30.8)	23.0 (14.8-33.7)	< 0.001	< 0.001	0.006

^aTwo-sample Wilcoxon rank-sum test, or Pearson chi-square test for gender and city.

^bMedian (5-95% percentiles).

^cNumber (%).

^dThe percentages of never-drinkers were 64.7, 50.4, and 61.1 in the 3 groups, respectively.

	Persistence	Clearance	Uninfected	Persistence vs. Clearance	Clearance vs. Uninfected	
	N = 102	N = 113	N = 2813	P value ^a	P value ^a	P value ^a
CD4 (%) ^b	40.8 (8.9)	42.2 (9.0)	43.0 (8.9)	0.32	0.007	0.46
$T_{H}1$ (%) ^b	35.2 (9.6)	27.1 (8.7)	26.0 (8.9)	< 0.001	< 0.001	0.23
$T_{\rm H}^{2} (\%)^{\rm b}$	1.55 (0.88)	1.74 (1.03)	1.79 (1.10)	0.54	0.11	0.65
$T_H 1/T_H 2^b$	30.7 (20.6)	20.8 (12.8)	20.8 (23.0)	< 0.001	< 0.001	0.27
CD8 (%) ^b	23.4 (9.6)	20.9 (7.9)	19.0 (7.8)	0.20	< 0.001	0.040
$T_{C}1(\%)^{b}$	42.3 (14.7)	38.8 (15.2)	39.5 (14.6)	0.25	0.17	0.68
$T_{C}^{2}(\%)^{b}$	2.78 (3.72)	2.87 (4.03)	3.35 (5.09)	0.34	0.68	0.40
$T_C 1/T_C 2^b$	42.8 (45.8)	50.9 (78.2)	51.1 (91.4)	0.66	0.38	0.51
CD4/CD8 ^b	2.08 (1.02)	2.42 (1.35)	2.75 (1.57)	0.15	< 0.001	0.054
CD16 (%) ^b	14.0 (9.4)	17.0 (8.8)	17.1 (9.4)	0.035	< 0.001	0.82
CD20 (%) ^b	14.5 (7.6)	13.5 (5.3)	13.9 (6.1)	0.95	0.44	0.60

Comparisons of peripheral lymphocyte subsets among subjects with no cancer history

^aTest of difference of logarithmic values between two groups using normal regression analysis with adjustment for age, gender, city, radiation dose, alcohol, smoking, BMI, autoimmune disease, allergic disease, and other non-cancer diseases.

^bMean (SD).

Regresson analysis of decrement	(per year)	in platelet counts among HCV	persistence subjects with no cancer histo	ry (N = 96)
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_	Unadjusted		Adjı	usted ^a	Adjusted ^b	
Explanatory variables	coefficient	P value	coefficient	P value	coefficient	P value
Age (+10 yrs)	-0.26	0.33	_	_	_	_
Gender (female vs. male)	0.43	0.21	_	_	_	_
Radiation dose (Gy)	0.28	0.26	_	_	_	_
City (Nagasaki vs. Hiroshima)	0.00	0.99	_	_	_	_
Log CD4	-0.25	0.86	1.11	0.50	_	_
Log T _H 1	-1.90	0.13	-2.38	0.086	-3.19	0.027
Log T _H 2	-0.53	0.41	-0.34	0.65	_	_
$Log T_H 1/T_H 2$	0.03	0.97	-0.31	0.67	_	_
Log CD8	-1.72	0.060	-2.63	0.011	_	-
Log T _C 1	1.66	0.068	2.05	0.041	2.36	0.027
Log T _C 2	-0.30	0.37	0.23	0.57	_	_
$Log T_C 1/T_C 2$	0.51	0.12	0.10	0.81	_	_
Log CD4/CD8	1.04	0.16	2.03	0.015	_	_
Log CD16	0.73	0.21	1.13	0.062	1.18	0.058
Log CD20	1.10	0.11	1.04	0.15	1.03	0.17

^aRegression model: *decrements in platelet counts* = $\alpha + \beta_1 \times log$ (*lymphocyte subset*) + $\beta_2 \times age + \beta_3 \times gender + \beta_4 \times dose + \beta_5 \times city$ + $\beta_6 \times alcohol + \beta_7 \times smoking + \beta_8 \times BMI + \beta_9 \times autoimmune disease + \beta_{10} \times allergic disease + \beta_{11} \times other non-cancer diseases.$ ^bForward step-wise procedure (P < 0.2) was used for 8 lymphocyte variables (CD4, T_H1, T_H2, CD8, T_C1, T_C2, CD16, and CD20). Selected 4 variables (T_H1, T_C1, CD16, and CD20) were used in the regression analysis with 10 explanatory variables in footnote a.