



# Impact of Epstein-Barr Virus on Peripheral T-Cell Lymphoma Not Otherwise Specified and Angioimmunoblastic T-Cell Lymphoma

Tong-Yoon Kim<sup>1</sup>, Gi-June Min<sup>1</sup>, Young-Woo Jeon<sup>2</sup>, Sung-Soo Park<sup>1</sup>, Silvia Park<sup>1</sup>, Seung-Hawn Shin<sup>3</sup>, Seung-Ah Yahng<sup>4</sup>, Jae-Ho Yoon<sup>1</sup>, Sung-Eun Lee<sup>1</sup>, Byung-Sik Cho<sup>1</sup>, Ki-Seong Eom<sup>1</sup>, Yoo-Jin Kim<sup>1</sup>, Seok Lee<sup>1</sup>, Hee-Je Kim<sup>1</sup>, Chang-Ki Min<sup>1</sup>, Jong-Wook Lee<sup>1</sup> and Seok-Goo Cho<sup>1\*</sup>

## OPEN ACCESS

### Edited by:

Swami P. Iyer,  
University of Texas MD Anderson  
Cancer Center, United States

### Reviewed by:

Praveen Ramakrishnan Geethakumari,  
University of Texas Southwestern  
Medical Center, United States  
Ranjit Nair,  
University of Texas MD Anderson  
Cancer Center, United States

### \*Correspondence:

Seok-Goo Cho  
chosg@catholic.ac.kr

### Specialty section:

This article was submitted to  
Hematologic Malignancies,  
a section of the journal  
Frontiers in Oncology

**Received:** 18 October 2021

**Accepted:** 20 December 2021

**Published:** 11 January 2022

### Citation:

Kim T-Y, Min G-J, Jeon Y-W,  
Park S-S, Park S, Shin S-H,  
Yahng S-A, Yoon J-H, Lee S-E,  
Cho B-S, Eom K-S, Kim Y-J, Lee S,  
Kim H-J, Min C-K, Lee J-W and  
Cho S-G (2022) Impact of Epstein-  
Barr Virus on Peripheral T-Cell  
Lymphoma Not Otherwise  
Specified and Angioimmunoblastic  
T-Cell Lymphoma.  
*Front. Oncol.* 11:797028.  
doi: 10.3389/fonc.2021.797028

<sup>1</sup> Department of Hematology, Seoul St. Mary's Hematology Hospital, College of Medicine, The Catholic University of Korea, Seoul, South Korea, <sup>2</sup> Department of Hematology, Yeouido St. Mary's Hematology Hospital, College of Medicine, The Catholic University of Korea, Seoul, South Korea, <sup>3</sup> Department of Hematology, Eunpyeong St. Mary's Hematology Hospital, College of Medicine, The Catholic University of Korea, Seoul, South Korea, <sup>4</sup> Department of Hematology, Incheon St. Mary's Hospital, College of Medicine, The Catholic University of Korea, Seoul, South Korea

**Purpose:** The significance of Epstein-Barr virus (EBV) infections for the prognosis of patients with peripheral T-cell lymphomas (PTCLs), specifically angioimmunoblastic T-cell lymphoma (AITL) and PTCL not otherwise specified (PTCL-NOS), remains unclear. The Epstein-Barr encoding region can be used to detect EBV in tissue sections by *in situ* hybridization (ISH) and by polymerase chain reaction (PCR) assays of peripheral blood samples from patients with PTCLs. This study compared the outcomes patients with AITL or PTCL-NOS for whom the presence of EBV infection was assessed by these two methods.

**Patients and Methods:** This was a retrospective study of patients newly diagnosed with AITL or PTCL-NOS. All patients were selected from a single transplantation center. EBV-positive lymphomas were detected at the time of diagnosis in tissue sections by ISH or in the blood by PCR.

**Results:** Out of a cohort of 140 patients with histologically confirmed AITL or PTCL-NOS, 105 were EBV-positive. The 3-year overall survival of patients with EBV-positive TCL was 43.3% compared to 68.6% in patients with EBV-negative TCL ( $p = .01$ ). Patients who were treated with autologous or allogeneic hematopoietic stem cell transplantation ( $n = 28$  and  $n = 11$ , respectively) or chemotherapy alone ( $n = 66$ ) had 3-year survival rates of 67.0%, 62.3%, and 30.2%, respectively ( $p < .02$ ). Patients with EBV-positive TCL had a better prognosis after treatment with hematopoietic stem cell transplantation compared to chemotherapy alone, but no difference was seen among patients with EBV-negative TCL.

**Conclusions:** EBV infection was shown to negatively affect the clinical outcomes of patients with TCL. Stem cell transplantation has been found to be an effective treatment

for EBV-associated lymphomas. Further investigations are warranted to determine the optimal treatment for these patients.

**Keywords:** Epstein-Barr virus, angioimmunoblastic T-cell lymphoma, peripheral T-cell lymphoma, T-cell lymphoma, transplantation

## INTRODUCTION

Peripheral T-cell lymphomas (PTCLs) are neoplasms derived from the post-thymic lymphocytes and account for approximately 15% of non-Hodgkin lymphoma in the Western hemisphere (1). Angioimmunoblastic T-cell lymphoma (AITL) and PTCL not otherwise specified (PTCL-NOS) are the second and third most common PTCLs in Asia and account for 24.7% and 20.8% of cases, respectively (2).

PTCL-NOS is diagnosed after excluding other subtypes and consists of both histologically and immunohistochemically heterogeneous tumors (3). Epstein-Barr virus (EBV) is detected in approximately 74–91% of all AITL cases and about 21% of PTCL-NOS cases (4, 5).

The effect of EBV on PTCLs has long been debated. According to the literature, some studies have reported a worse prognosis in patients with EBV detected in peripheral blood by a polymerase chain reaction (PCR) (6–8). However, no studies comparing the different EBV detection methods, such as *in situ* hybridization (ISH) in tissue sections and PCR assays using blood specimens, have been conducted on patients with PTCL-NOS nor AITL.

This study aimed to investigate the impact of EBV status on the prognosis of patients with PTCL and to compare the use of the two methods for detecting EBV: EBV-encoded small RNA *in situ* hybridization and PCR of DNA polymerase gene. We also assessed the role of hematopoietic stem cell transplantation (HSCT) in these patients.

## MATERIALS AND METHODS

### Study Population

The study protocol was approved by the Institutional Review Board and Ethics Committee of the Catholic Medical Center in South Korea (KC21RISI0603). The need for informed consent was waived due to the retrospective nature of the study.

This study included patients diagnosed with AITL or PTCL-NOS between May 2009 and May 2019 at the Catholic Hematology Hospital in South Korea. The inclusion criteria were patients with biopsy-proven lymphoma and EBV status analyzed using both ISH and PCR. Patients either without PCR or ISH testing were excluded from the study. All cases were confirmed as PTCL-NOS or AITL using the morphological and immunophenotypic diagnostic criteria of the World Health Organization (3). Patients were evaluated before treatment in terms of performance status, physical examination, complete and differential blood counts, blood chemistry, lactate dehydrogenase levels, chest radiography, and bone marrow biopsy. Computed

tomography (CT) of the chest, abdomen, and pelvis and positron emission tomography (PET) were used to determine cancer stage in the cohort.

### Treatment and Response

ProMACE-CytaBOM, CHOP, or SMILE chemotherapy regimens were used, as mentioned in previous studies. The ProMACE-CytaBOM regimen consisted of doxorubicin, cyclophosphamide, and etoposide on day 1 followed by bleomycin, cytosine arabinoside, and vincristine on day 8 as well as prednisone on days 1–14. The CHOP regimen consisted of doxorubicin, cyclophosphamide, and vincristine on day 1 and prednisone on days 1–5 and was repeated every 21 days (9). The SMILE regimen consisted of methotrexate on day 1, ifosfamide and etoposide on days 2–4, and intravenous L-asparaginase based on the existing protocol (10). The doses were adjusted for all patients based on their performance status in addition to hematological, neurological, and infusion-related adverse effects.

Response evaluation was done quarterly for all patients during the first year, biannually for the next three years, and annually thereafter to detect disease recurrence. The response criteria for malignant lymphoma included complete remission (CR; no evidence of disease), partial remission (PR; disease regression but with a measurable remnant and no new lesions), progressive disease (PD; new lesion  $\geq$  50% of the size of the previous lesion), and stable disease (SD; absence of CR, PR, and PD). Progress free survival (PFS) is defined as the time from diagnosis until lymphoma progression or death as a result of any cause. Overall survival (OS) is defined as the time from diagnosis until death as a result of any cause (11). The cumulative incidence of relapse (CIR) represented the time between disease diagnosis and relapse.

All patients with AITL and PTCL who achieved CR were initially recommended for frontline autologous HSCT. Our department recommended salvage allogeneic HSCT in refractory or relapsed after CR and PR. For conditioning regimen for autologous HSCT, dose reduction strategy of a BU-MEL-TT regimen were used as previous described (12). BU-MEL-TT regimen consist of 2.4 mg/kg busulfan for three consecutive days (total 7.2 mg/kg), 40 mg/m<sup>2</sup> melphalan for 2 days (total 80 mg/m<sup>2</sup>), and 200 mg/m<sup>2</sup>/thiotepa for two consecutive days (total 400 mg/m<sup>2</sup>). The standard regimen for allogeneic HSCT, RIC FMT regimen previously published (13), consisted of 30 mg/m<sup>2</sup> fludarabine for 6 consecutive days (total 180 mg/m<sup>2</sup>) and 70 mg/m<sup>2</sup> melphalan for 1 day with TBI of 800 cGy in 4 fractionated doses for 2 days. Graft versus host disease prophylaxis mainly consisted of cyclosporine for all sibling transplants or tacrolimus for unrelated or haploidentical transplants) with methotrexate (5 mg/m<sup>2</sup> for tacrolimus and 10 mg/m<sup>2</sup> for cyclosporine on days after transplant 1, 3, 6, and 11).

## Real-Time PCR for Plasma EBV

Blood samples were collected in tubes containing ethylenediaminetetraacetic acid, and DNA was extracted using the Real-Q EBV quantification kit (BioSewoom Inc., Seoul, Korea). Real-time quantitative PCR assays were performed for EBV-specific sequences using the ABI Prism 7500 system (Applied Biosystems Inc., Foster City, CA, USA). Copy numbers were analyzed using a standard curve. The lower threshold for test sensitivity was  $5 \times 10^2$  copies/mL, and patients with lower values were considered EBV-DNA-negative.

## Tissue ISH

The presence of EBV-specific small RNAs was investigated by ISH. EBV-encoded small RNA (EBER) oligonucleotides were added to formalin-fixed paraffin-embedded sections in accordance with the Inform EBER Probe Assay Protocol (Ventana Medical Systems Inc., Oro Valley, AZ, USA).

## Statistical Analysis

All categorical variables were analyzed using the chi-squared or Fisher's exact tests whereas the Mann-Whitney *u* test and Student's *t*-test were used for continuous ones.

PFS, OS, and CIR were assessed using Kaplan-Meier curves and compared using the log-rank test. In order to identify the factors that significantly affected survival in the univariate analysis, a Cox proportional-hazards model was used. All predictors with a *p*-value of  $<.1$  were included in the multivariate analysis. The reported *p*-values were two-sided, and statistical significance was set at  $p <.05$ . All statistical analyses were performed using the R software (version 3.4.3).

## RESULTS

### Patient Characteristics

In total, 140 patients with AITL ( $n = 68$ ) or PTCL-NOS ( $n = 72$ ) were included in this study. We defined the EBV-positive group as either detected by PCR or ISH, which was comprised 105 patients. Neither detected by PCR and ISH was 35 patients with EBV-negative group. The median age was 55 years (range, 18–83 years) in the EBV-positive group and 55 years (range, 27–77 years) in the EBV-negative group. Using the International Prognostic Index (IPI), 45.7% of patients in the EBV-positive group scored 0–2, and 54.3% scored 3–5. In the EBV-negative group, these percentages were 48.6% and 51.4%, respectively. There were more EBV-positive cases among patients with AITL (57.1%,  $n = 60$ ) than among patients with PTCL-NOS (42.9%,  $n = 45$ ). Patient characteristics are summarized in **Table 1**.

### Treatment Outcomes

Chemotherapy was administered to all patients, 75 (53.6%) of whom had an IPI score of 3–5. ProMACE-CytaBOM ( $n = 52$ ), CHOP ( $n = 73$ ), and SMILE ( $n = 15$ ) regimens were administered to the patients. Response evaluation was performed after three or four cycles of chemotherapy. CR was found in 81 patients (57.9%), and PR, PD or SD were observed in

59 patients (42.1%). Among the 55 patients with EBV-positive PTCL who achieved CR with the first treatment, 29 relapsed (52.7%). In comparison, among the 26 patients with CR in EBV-negative group, 11 relapsed (42.3%). Salvage chemotherapy resulted in CR in 14 patients with EBV-positive PTCL who failed to achieve CR in the first chemotherapy (28.0%) and in three patients with EBV-negative PTCL (33.3%).

In the subgroup analysis, significant differences were observed in the CR rates between patients with positive or negative PCR results (hazard ratio [HR] = 0.65, 95% confidence interval [CI]: 0.44 – 0.97,  $p = .04$ ). Among patients with negative PCR results, those who were ISH-positive for EBV also relapsed more frequently compared to patients with negative ISH results (HR = 2.17, 95% CI: 1.03 – 4.56,  $p = .035$ ) (**Supplementary Figure 1**).

After a median follow-up of 33 months (ranging from 0.5 – 123 months), the survival rates of patients with AITL or PTCL-NOS were 52.9% and 46.6%, respectively, with no statistically significant difference between these groups ( $p = 1.0$ ) (**Figure 1A**). The 3-year PFS was significantly lower ( $p = .02$ ) in the patients with EBV-positive PTCL (29.1%,  $n = 105$ ) compared to their EBV-negative counterparts (57.1%,  $n = 35$ ). Patients in the EBV-positive group were less likely to survive compared to those in the EBV-negative group (43.3% and 68.6%, respectively,  $p = .01$ ) (**Figure 1B**).

The univariate analysis identified factors associated with increased mortality, which included advanced age ( $>60$  years), high serum  $\beta$ -microglobulin, high and high-intermediate IPI, and the prognostic index for T-cell lymphoma scores as well as EBV positivity. In the multivariate analysis, patients with EBV-positive PTCL were more likely to die (HR = 2.07, 95% CI: 1.19 – 3.65,  $p = .011$ ) as were those with high and high-intermediate IPI scores (HR = 1.71, 95% CI: 1.11 – 2.66,  $p = .016$ ).

### Distribution and Survival of EBV-Positive AITL and PTCL-NOS

In the AITL group, 41 patients (60.3%) were both PCR- and ISH-positive, 15 (22.1%) were only ISH-positive, 4 (5.9%) were only PCR-positive, and 8 (11.8%) were negative for both (**Figure 2A**). In the PTCL-NOS group, 24 patients (33.3%) were both PCR- and ISH-positive, 12 (16.7%) were only ISH-positive, 9 (12.5%) were only PCR-positive, and 27 (37.5%) were negative for both (**Figure 2B**).

Among patients with both PCR and ISH positivity, the 2-year survival rates were 46.3% and 41.3% for those with AITL and PTCL-NOS, respectively. These percentages were 87.5% and 66.6% in the PCR- and ISH-negative group, 73.3% and 58.3% in the ISH-only positive group, and 50.0% and 33.3% in the PCR-only positive group (**Figure 2C** and **Supplementary Figure 2A**).

### Subgroup Survival Analysis Among Patients With EBV-Positive and EBV-Negative PTCL With or Without Treatment by HSCT

In the EBV-positive group, the 3-year OS rate was 67.0% for the 28 patients treated with autologous HSCT, 62.3% for the 11 treated with allogeneic HSCT, and 30.2% for the 66 treated with

**TABLE 1 |** Characteristics of patients with PTCL-NOS or AITL.

Characteristic	EBV-negative (n = 35)	EBV-positive (n = 105)	p-value
<b>Age</b>			.465
Median (years)	55 (27–77)	55 (18–83)	
Age ≤ 60, n (%)	26 (74.3)	69 (65.7)	
Age > 60, n (%)	9 (25.7)	36 (34.3)	
<b>Sex</b>			.999
Female, n (%)	16 (45.7)	46 (43.8)	
Male, n (%)	19 (54.3)	59 (56.2)	
<b>Histological subtype</b>			.001
AITL, n (%)	8 (22.9)	60 (57.1)	
PTCL-NOS, n (%)	27 (77.1)	45 (45.9)	
<b>Poor performance status</b>			.347
ECOG 0 – 1, n (%)	21 (60)	74 (70.5)	
ECOG > 1, n (%)	14 (40)	31 (29.5)	
<b>Extranodal involvement ≥ 2</b>			.238
n (%)	12/35 (34.3)	50/105 (47.6)	
<b>Bone marrow involvement</b>			.764
n (%)	15/35 (42.9)	40/105 (38.1)	
<b>Ann Arbor stage</b>			.999
I – II, n (%)	8 (22.9)	25 (23.8)	
III – IV, n (%)	27 (77.1)	80 (76.2)	
<b>High serum LDH</b>			.919
n (%)	22/35 (62.9)	69/105 (65.7)	
<b>High serum β-microglobulin</b>			.494
n (%)	15/35 (50.0)	43/105 (59.7)	
<b>CD4 positive</b>			.565
n (%)	25/35 (71.4)	82/105 (78.1)	
<b>CD8 positive</b>			.470
n (%)	14/35 (40.0)	33/105 (31.4)	
<b>CD30 positive</b>			.038
n (%)	9/30 (30.0)	41/75 (54.7)	
<b>Ki-67 ≥ 80</b>			.245
n (%)	14/35 (40.0)	29/105 (27.6)	
<b>IPI score</b>			.922
0 – 2 Low/low-intermediate, n (%)	17 (48.6)	48 (45.7)	
3 – 5 High-intermediate/high, n (%)	18 (51.4)	57 (54.3)	
<b>PIT score</b>			.735
1 – 2 Low/low-intermediate, n (%)	25 (71.4)	80 (76.2)	
3 – 4 High-intermediate/high, n (%)	10 (28.4)	25 (23.8)	
<b>mPIT score</b>			.437
1 – 2 Low/intermediate, n (%)	27 (77.1)	89 (84.8)	
3 High, n (%)	8 (22.9)	16 (15.2)	

AITL, angioimmunoblastic T-cell lymphoma; ECOG, Eastern Cooperative Oncology Group; IPI, International Prognostic Index; LDH, lactate dehydrogenase; PIT, Prognostic Index for T-cell lymphoma; mPIT, modified Prognostic Index for T-cell lymphoma; PTCL-NOS, peripheral T-cell lymphoma not otherwise specified.

chemotherapy only ( $p < .02$ ). However, no statistically significant differences were observed in the OS of the EBV-negative group (**Figure 3**).

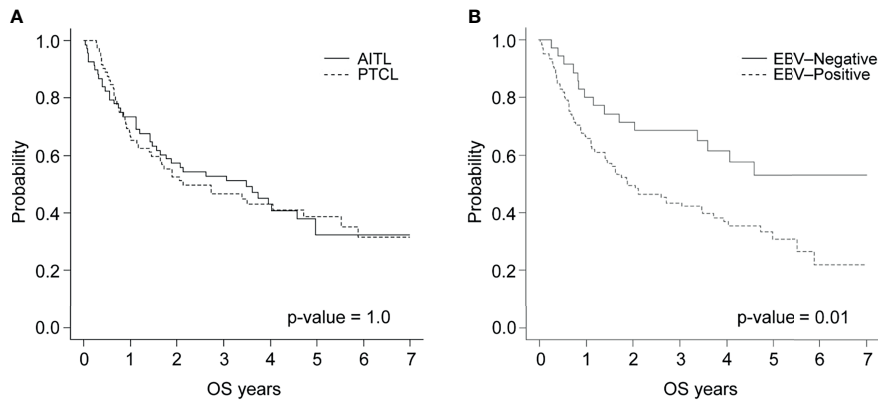
## DISCUSSION

Although there have been several studies on the prognosis of EBV-positive T-cell lymphomas (14), there are few studies that compare the prognosis patients with AITL or PTCL-NOS. EBV infection is rarer in the West than in Asia, and the proportion of T- and NK-cell lymphomas is higher in East Asia than in other regions (15, 16). Because differences in racial and regional genetic factors could influence the immune response to EBV (17), evaluating the prognostic impact of EBV infection is essential in Korea.

Our study was based on the similarity in survival outcomes between patients with AITL or PTCL-NOS. Warnke et al. described the challenges of distinguishing AITL from PTCL-NOS due to their morphological and genetic similarities. Furthermore, their study demonstrated that patients with PTCL-NOS who had a genetic signature similar to AITL were prone to poor outcomes (18).

We hypothesized that EBV status could comparably impact AITL and PTCL-NOS. EBV infection is generally evaluated by two methods, and EBV-ISH is currently used to differentiate lymphoma subtypes and gives clues to distinguish AITL from PTCL-NOS (19, 20). Detection of EBV by PCR is used for prognosis prediction (21) and the early detection (22) of various type of cancers. Therefore, we tried to assess this in our cohort.

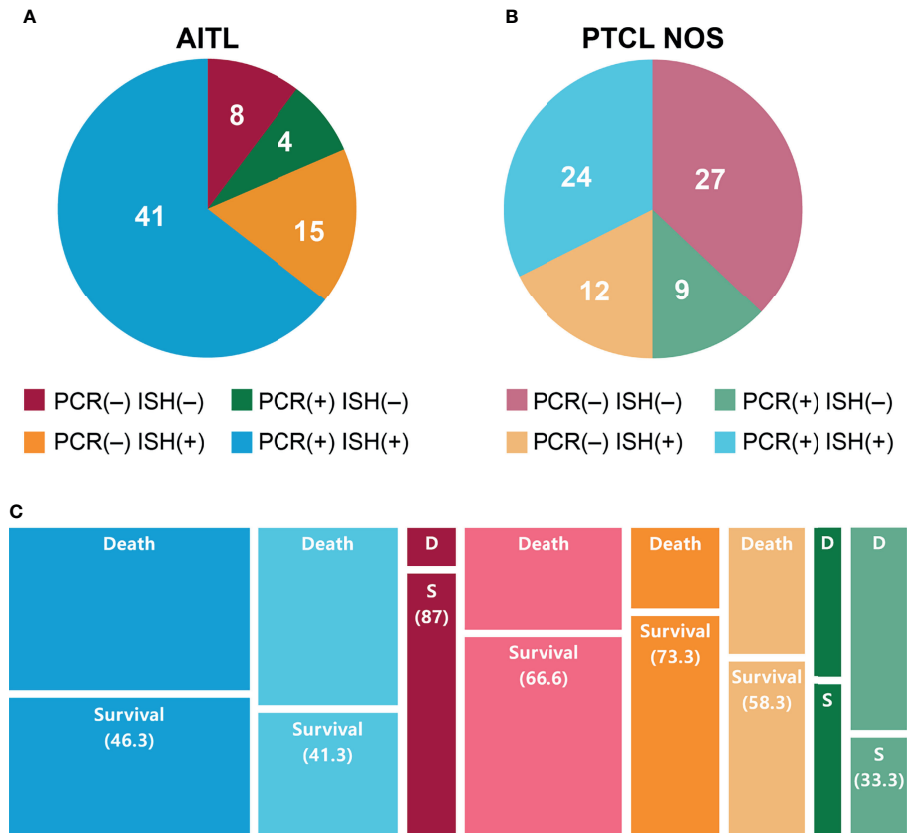
In our study, we observed a poor prognosis of patients with EBV-positive PTCL detected by ISH or PCR. These two



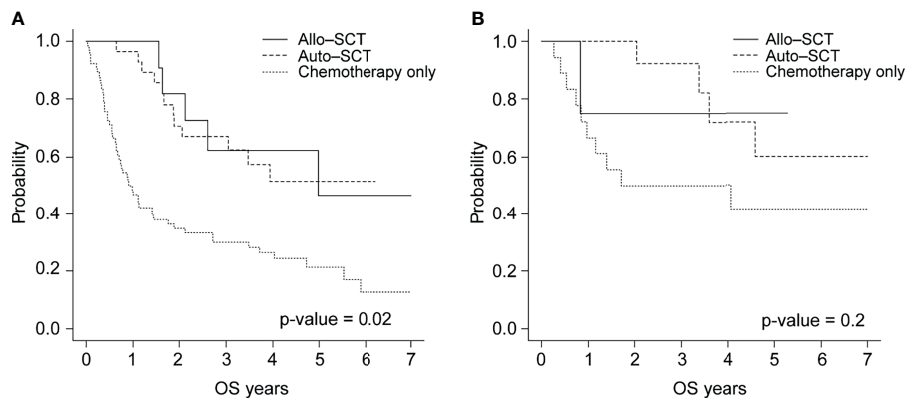
**FIGURE 1** | Overall survival by (A) tumor subtype and (B) EBV status.

detection methods play different roles in predicting poor outcomes. Blood PCR-positive cases showed resistance to the first-line chemotherapy. In the PCR-negative group, patients with ISH-positive PTCL had a higher recurrence rate compared to their ISH-negative counterparts regardless of a good response to chemotherapy; hence, they were all included

in the EBV-positive group. EBV PCR positivity was associated with chemotherapy resistance, which is consistent with published (23). Although a few studies support this trend in patients with PTCL-NOS or AITL, these studies only showed that cases of EBV status detected by PCR and not ISH were associated with poor survival outcomes. However, these studies



**FIGURE 2** | EBV infection in patients with (A) AITL or (B) PTCL-NOS according to PCR and ISH. (C) Survival of patients with AITL or PTCL-NOS after 2 years based on EBV PCR and ISH status.



**FIGURE 3** | Overall survival of patients with (A) EBV-positive or (B) EBV-negative PTCL by treatment type (allogeneic HSCT, autologous HSCT, and chemotherapy).

include cases other types of T-cell lymphoma regardless of different survival outcomes associated with each subtypes, which could obscure the influence of EBV infection (6, 7).

Even though EBV ISH positivity was associated with early relapse in patients with B-cell lymphoma (24, 25) there is a lack of data on TCL. Some studies shown that ISH-positive cases are associated with higher susceptibility of relapse (26) but this has been contradicted by other findings (27). This discordance arises because EBV-ISH and EBV-PCR results should be interpreted together. For example, in the cases of ISH-negative and PCR-positive TCL, which are associated with a poor outcome, the source of the EBV infection could have come from B cells and may need additional treatment such as rituximab administration.

Due to the overlapping characteristics of PTCL-NOS and AITL, similar treatment strategies consisting of autologous HSCT following chemotherapy (28, 29) and salvage allogeneic HSCT have been used in the clinic (30). Despite a few prospective studies that have demonstrated its benefits (31), the role of first-line autologous HSCT is not confirmed (32). However, EBV-associated lymphoma may be amenable to first-line autologous HSCT in TCL. EBV-induced lymphomas have high proliferative signaling and cell death resistance (33). EBV latency in tumor cells increases intracellular ROS activity and the MDR gene expression by activating the STAT1 signaling pathway and phosphorylated STAT1, promoting P-gp expression. P-gp-induced drug efflux resulted in reduced chemo sensitivity (34). Recently, front-line autologous HSCT has been attempted for extranodal NK/T-cell lymphoma (ENKTCL) and has proven useful (35).

As additional treatments, Epstein-Barr nuclear antigen and EBV latent membrane protein-specific cytotoxic T-lymphocytes (CTLs) might be beneficial. CTLs have been used to treat EBV-positive Hodgkin, non-Hodgkin lymphoma and ENKTCL with favorable outcomes (36–38). However, the efficacy of CTLs may be limited in the tumor microenvironment, which impairs the recognition of lymphoma cells by CTLs. For example, the interaction between PD-1 and PD-L1 downregulates T-cell function and leads to CTL exhaustion. As EBV infection

systemically affects the immune system, EBV-positive PTCL-NOS and AITL could be candidates for the use of immune checkpoint inhibitors (39–41). Additional studies focusing on different treatment modalities for EBV-positive PTCL-NOS and AITL are warranted.

In observational studies, there is a potential for bias based on the inclusion criteria. Since we excluded the cases where EBV-related tests were omitted, the EBV distribution may be relatively different from the previously published. This explains the relatively higher EBV status on PTCL NOS in our cohort. We did not perform first-line autologous HSCT as in prospective studies (31). Clinicians recommend autologous HSCT to all patients with susceptible as we described, and some cases were not proceeded due to poor physical status or personal issues. Therefore, interpretation of the effect of first-line autologous HSCT and salvage allogeneic HSCT is limited.

In conclusion, testing for EBV status using both PCR and ISH in patients with AITL or PTCL-NOS may be useful, and HSCT may help improve survival rates in of infected patients. A large-scale study of the effect of EBV infection on the pathological mechanisms of PTCL-NOS and AITL is needed, and prospective randomized clinical trials are required to determine the standard treatments for EBV-associated AITL and PTCL-NOS.

## DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

## ETHICS STATEMENT

The study protocol was approved by the Institutional Review Board and Ethics Committee of the Catholic Medical Center in South Korea (KC21RISI0603). Written informed consent for participation was not required for this study in accordance with the national legislation and the institutional requirements.

## AUTHOR CONTRIBUTIONS

T-YK performed the research, collected and analyzed the data, in addition to writing the manuscript. G-JM, S-HS, S-AY, J-HY, B-SC, and H-JK provided patients and materials and reviewed the manuscript. S-SP, S-EL, K-SE, and C-KM provided the materials and reviewed the manuscript. Y-JK, SL, and J-WL reviewed the manuscript and analyzed the data. S-GC designed and conducted the study, provided patients and materials,

analyzed data, and wrote the manuscript. All authors read and approved the final manuscript.

## SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fonc.2021.797028/full#supplementary-material>

## REFERENCES

- Fiore D, Cappelli LV, Broccoli A, Zinzani PL, Chan WC, Inghirami G, et al. Peripheral T Cell Lymphomas: From the Bench to the Clinic. *Nat Rev Cancer* (2020) 20:323–42. doi: 10.1038/s41568-020-0247-0
- Yoon SE, Song Y, Kim SJ, Yoon DH, Chen TY, Koh Y, et al. Comprehensive Analysis of Peripheral T-Cell and Natural Killer/T-Cell Lymphoma in Asian Patients: A Multinational, Multicenter, Prospective Registry Study in Asia. *Lancet Reg Health West Pac* (2021) 10:100126. doi: 10.1016/j.lanwpc.2021.100126
- Swerdlow SH, Campo E, Pileri SA, Harris NL, Stein H, Siebert R, et al. The 2016 Revision of the World Health Organization Classification of Lymphoid Neoplasms. *Blood* (2016) 127:2375–90. doi: 10.1182/blood-2016-01-643569
- Hsi ED, Horwitz SM, Carson KR, Pinter-Brown LC, Rosen ST, Pro B, et al. Analysis of Peripheral T-Cell Lymphoma Diagnostic Workup in the United States. *Clin Lymphoma Myeloma Leuk* (2017) 17:193–200. doi: 10.1016/j.clml.2016.10.001
- Dobay MP, Lemonnier F, Missaglia E, Bastard C, Vallois D, Jais J-P, et al. Integrative Clinicopathological and Molecular Analyses of Angioimmunoblastic T-Cell Lymphoma and Other Nodal Lymphomas of Follicular Helper T-Cell Origin. *Haematologica* (2017) 102:e148–51. doi: 10.3324/haematol.2016.158428
- Kim YR, Kim SJ, Cheong JW, Chung H, Eun Jang J, Kim Y, et al. Pretreatment Epstein–Barr Virus DNA in Whole Blood Is a Prognostic Marker in Peripheral T-Cell Lymphoma. *Oncotarget* (2017) 8:92312–23. doi: 10.18632/oncotarget.21251
- Liang JH, Lu L, Zhu HY, Li W, Fan L, Li JY, et al. The Prognostic Role of Circulating Epstein–Barr Virus DNA Copy Number in Angioimmunoblastic T-Cell Lymphoma Treated With Dose-Adjusted EPOCH. *Cancer Res Treat* (2019) 51:150–7. doi: 10.4143/crt.2017.476
- Haverkos BM, Huang Y, Gru A, Pancholi P, Freud AG, Mishra A, et al. Frequency and Clinical Correlates of Elevated Plasma Epstein–Barr Virus DNA at Diagnosis in Peripheral T-Cell Lymphomas. *Int J Cancer* (2017) 140:1899–906. doi: 10.1002/ijc.30566
- Montserrat E, García Conde J, Viñolas N, López Guillermo A, Hernández-Nieto L, Zubizarreta A, et al. CHOP vs. ProMACE-CytaBOM in the Treatment of Aggressive Non-Hodgkin's Lymphomas: Long-Term Results of a Multicenter Randomized Trial. (PETHEMA: Spanish Cooperative Group for the Study of Hematological Malignancies Treatment, Spanish Society of Hematology). *Eur J Haematol* (1996) 57:377–83. doi: 10.1111/j.1600-0609.1996.tb01396.x
- Yamaguchi M, Suzuki R, Kwong YL, Kim WS, Hasegawa Y, Izutsu K, et al. Phase I Study of Dexamethasone, Methotrexate, Ifosfamide, L-Asparaginase, and Etoposide (SMILE) Chemotherapy for Advanced-Stage, Relapsed or Refractory Extranodal Natural Killer (NK)/T-Cell Lymphoma and Leukemia. *Cancer Sci* (2008) 99:1016–20. doi: 10.1111/j.1349-7006.2008.00768.x
- Cheson BD, Pfistner B, Juweid ME, Gascoyne RD, Specht L, Horning SJ, et al. Revised Response Criteria for Malignant Lymphoma. *J Clin Oncol* (2007) 25:579–86. doi: 10.1200/JCO.2006.09.2403
- Yoon JH, Min GJ, Park SS, Jeon YW, Lee SE, Cho BS, et al. Autologous Hematopoietic Cell Transplantation Using Dose-Reduced Intravenous Busulfan, Melphalan, and Thiotepa for High-Risk or Relapsed Lymphomas. *Bone Marrow Transplant* (2019) 54:330–3. doi: 10.1038/s41409-018-0289-z
- Jeon YW, Yoon S, Min GJ, Park SS, Park S, Yoon JH, et al. Clinical Outcomes of Fludarabine and Melphalan With an 800 cGy Total Body Irradiation Conditioning Regimen in Patients With Refractory or Relapsed Aggressive Non-Hodgkin Lymphoma Undergoing Allogeneic Hematopoietic Stem Cell Transplantation. *Clin Lymphoma Myeloma Leuk* (2019) 19:345–355.e7. doi: 10.1016/j.clml.2019.03.023
- Harabuchi Y, Takahara M, Kishibe K, Nagato T, Kumai T. Extranodal Natural Killer/T-Cell Lymphoma, Nasal Type: Basic Science and Clinical Progress. *Front Pediatr* (2019) 7:141. doi: 10.3389/fped.2019.00141
- Park S, Ko YH. Peripheral T Cell Lymphoma in Asia. *Int J Hematol* (2014) 99:227–39. doi: 10.1007/s12185-014-1520-3
- Cho EY, Kim KH, Kim WS, Yoo KH, Koo HH, Ko YH. The Spectrum of Epstein–Barr Virus-Associated Lymphoproliferative Disease in Korea: Incidence of Disease Entities by Age Groups. *J Korean Med Sci* (2008) 23:185–92. doi: 10.3346/jkms.2008.23.2.185
- Ko YH. EBV-Associated Lymphoproliferative Disorders. *Clin Pediatr Hematol Oncol* (2021) 28:14–27. doi: 10.15264/cpho.2021.28.1.14
- Iqbal J, Wright G, Wang C, Rosenwald A, Gascoyne RD, Weisenburger DD, et al. Gene Expression Signatures Delineate Biological and Prognostic Subgroups in Peripheral T-Cell Lymphoma. *Blood* (2014) 123:2915–23. doi: 10.1182/blood-2013-11-536359
- Nakatsuka S-i, Homma K, Aozasa K. When to Use *In Situ* Hybridization for the Detection of Epstein–Barr Virus: A Review of Epstein–Barr Virus-Associated Lymphomas. *J Hematopathol* (2015) 8:61–70. doi: 10.1007/s12308-014-0230-3
- Attygalle AD, Chuang SS, Diss TC, Du MQ, Isaacson PG, Dogan A. Distinguishing Angioimmunoblastic T-Cell Lymphoma From Peripheral T-Cell Lymphoma, Unspecified, Using Morphology, Immunophenotype and Molecular Genetics. *Histopathology* (2007) 50:498–508. doi: 10.1111/j.1365-2559.2007.02632.x
- Kanakry JA, Li H, Gellert LL, Lemas MV, Hsieh WS, Hong F, et al. Plasma Epstein–Barr Virus DNA Predicts Outcome in Advanced Hodgkin Lymphoma: Correlative Analysis From a Large North American Cooperative Group Trial. *Blood* (2013) 121:3547–53. doi: 10.1182/blood-2012-09-454694
- Chan KCA, Woo JKS, King A, Zee BCY, Lam WKJ, Chan SL, et al. Analysis of Plasma Epstein–Barr Virus DNA to Screen for Nasopharyngeal Cancer. *N Engl J Med* (2017) 377:513–22. doi: 10.1056/NEJMoa1701717
- Wang L, Wang H, Wang JH, Xia ZJ, Lu Y, Huang HQ, et al. Post-Treatment Plasma EBV-DNA Positivity Predicts Early Relapse and Poor Prognosis for Patients With Extranodal NK/T Cell Lymphoma in the Era of Asparaginase. *Oncotarget* (2015) 6:30317–26. doi: 10.18632/oncotarget.4505
- Park S, Lee J, Ko YH, Han A, Jun HJ, Lee SC, et al. The Impact of Epstein–Barr Virus Status on Clinical Outcome in Diffuse Large B-Cell Lymphoma. *Blood* (2007) 110:972–8. doi: 10.1182/blood-2007-01-067769
- Crombie JL, LaCasce AS. Epstein Barr Virus Associated B-Cell Lymphomas and Latrogenic Lymphoproliferative Disorders. *Front Oncol* (2019) 9:109. doi: 10.3389/fonc.2019.00109
- Dupuis J, Emile JF, Mounier N, Gisselbrecht C, Martin-Garcia N, Petrella T, et al. Prognostic Significance of Epstein–Barr Virus in Nodal Peripheral T-Cell Lymphoma, Unspecified: A Groupe D'Etude Des Lymphomes De L'adulte (GELA) Study. *Blood* (2006) 108:4163–9. doi: 10.1182/blood-2006-04-017632
- Eladl AE, Shimada K, Suzuki Y, Takahara T, Kato S, Kohno K, et al. Status has Prognostic Implication Among Young Patients With Angioimmunoblastic T-Cell Lymphoma. *Cancer Med* (2020) 9:678–88. doi: 10.1002/cam4.2742
- Moskowitz AJ. Practical Treatment Approach for Angioimmunoblastic T-Cell Lymphoma. *J Oncol Pract* (2019) 15:137–43. doi: 10.1200/JOP.18.00511

29. Tang T, Khoo LP, Lim C, Ham JS, Kim SJ, Hong H, et al. Outcomes of Patients With Peripheral T-Cell Lymphoma in First Complete Remission: Data From Three Tertiary Asian Cancer Centers. *Blood Cancer J* (2017) 7:653. doi: 10.1038/s41408-017-0030-y
30. Goldberg JD, Chou JF, Horwitz S, Teruya-Feldstein J, Barker JN, Boulad F, et al. Long-Term Survival in Patients With Peripheral T-Cell Non-Hodgkin Lymphomas After Allogeneic Hematopoietic Stem Cell Transplant. *Leuk Lymphoma* (2012) 53:1124–9. doi: 10.3109/10428194.2011.645818
31. d'Amore F, Relander T, Lauritzsen GF, Jantunen E, Hagberg H, Anderson H, et al. Up-Front Autologous Stem-Cell Transplantation in Peripheral T-Cell Lymphoma: NLG-T-01. *J Clin Oncol* (2012) 30:3093–9. doi: 10.1200/JCO.2011.40.2719
32. Fossard G, Broussais F, Coelho I, Bailly S, Nicolas-Virelizier E, Toussaint E, et al. Role of Up-Front Autologous Stem-Cell Transplantation in Peripheral T-Cell Lymphoma for Patients in Response After Induction: An Analysis of Patients From LYSA Centers. *Ann Oncol* (2018) 29:715–23. doi: 10.1093/annonc/mdx787
33. Mesri EA, Feitelson MA, Munger K. Human Viral Oncogenesis: A Cancer Hallmarks Analysis. *Cell Host Microbe* (2014) 15:266–82. doi: 10.1016/j.chom.2014.02.011
34. Nam YS, Im KI, Kim N, Song Y, Lee JS, Jeon YW, et al. Down-Regulation of Intracellular Reactive Oxygen Species Attenuates P-Glycoprotein-Associated Chemoresistance in Epstein–Barr Virus-Positive NK/T-Cell Lymphoma. *Am J Transl Res* (2019) 11:1359–73.
35. Yhim HY, Kim JS, Mun YC, Moon JH, Chae YS, Park Y, et al. Clinical Outcomes and Prognostic Factors of Up-Front Autologous Stem Cell Transplantation in Patients With Extranodal Natural Killer/T Cell Lymphoma. *Biol Blood Marrow Transplant* (2015) 21:1597–604. doi: 10.1016/j.bbmt.2015.05.003
36. Cho SG, Kim N, Sohn HJ, Lee SK, Oh ST, Lee HJ, et al. Long-Term Outcome of Extranodal NK/T Cell Lymphoma Patients Treated With Postremission Therapy Using EBV LMP1 and LMP2a-Specific CTLs. *Mol Ther* (2015) 23:1401–9. doi: 10.1038/mt.2015.91
37. Bollard CM, Gottschalk S, Torrano V, Diouf O, Ku S, Hazrat Y, et al. Sustained Complete Responses in Patients With Lymphoma Receiving Autologous Cytotoxic T Lymphocytes Targeting Epstein–Barr Virus Latent Membrane Proteins. *J Clin Oncol* (2014) 32:798–808. doi: 10.1200/JCO.2013.51.5304
38. Icheva V, Kayser S, Wolff D, Tuve S, Kyzirakos C, Bethge W, et al. Adoptive Transfer of Epstein–Barr Virus (EBV) Nuclear Antigen 1–Specific T Cells as Treatment for EBV Reactivation and Lymphoproliferative Disorders After Allogeneic Stem-Cell Transplantation. *J Clin Oncol* (2013) 31:39–48. doi: 10.1200/JCO.2011.39.8495
39. Bi XW, Wang H, Zhang WW, Wang JH, Liu WJ, Xia ZJ, et al. PD-L1 Is Upregulated by EBV-Driven LMP1 Through NF- $\kappa$ B Pathway and Correlates With Poor Prognosis in Natural Killer/T-Cell Lymphoma. *J Hematol Oncol* (2016) 9:109. doi: 10.1186/s13045-016-0341-7
40. Nagato T, Ohkuri T, Ohara K, Hirata Y, Kishibe K, Komabayashi Y, et al. Programmed Death-Ligand 1 and Its Soluble Form Are Highly Expressed in Nasal Natural Killer/T-Cell Lymphoma: A Potential Rationale for Immunotherapy. *Cancer Immunol Immunother* (2017) 66:877–90. doi: 10.1007/s00262-017-1987-x
41. Kwong YL, Chan TSY, Tan D, Kim SJ, Poon LM, Mow B, et al. PD1 Blockade With Pembrolizumab is Highly Effective in Relapsed or Refractory NK/T-Cell Lymphoma Failing L-Asparaginase. *Blood* (2017) 129:2437–42. doi: 10.1182/blood-2016-12-756841

**Conflict of Interest:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

**Publisher's Note:** All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

Copyright © 2022 Kim, Min, Jeon, Park, Park, Shin, Yahng, Yoon, Lee, Cho, Eom, Kim, Lee, Kim, Min, Lee and Cho. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.