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Epstein-Barr virus-specific T-cell response in pediatric liver transplant recipients: a crosssectional study by multiparametric flow cytometry

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Background: Epstein-Barr virus (EBV) specific T-cell response measurement can help adjust immunosuppression in transplant patients with persistent infections. We aim to define T-cell responses against EBV in a cohort of pediatric liver-transplant patients.

Methods: Thirty-eight immunosuppressed pediatric liver-transplant patients (IP) and 25 EBV-seropositive healthy-adult controls (HC) were included in our cross-sectional study. Based on their EBV serological (S) and viral load (VL) status, patients were categorized into IP-S^{NEG}, IP-S^{POS}VL^{NEG} and IP-S^{POS}VL^{POS} groups. T-cell response was assessed at two timepoints by stimulating cells with EBV peptides (PepTivator[®]) and performing intracellular-cytokine and activation-induced marker staining. Background subtraction was used to determine EBV-specific T-lymphocyte frequency.

Results: Polyfunctional CD8+ T cells indicated previous EBV contact (IP-S^{NEG} 0.00% vs IP-S^{POS} 0.04% and HC 0.02%; p=0.001 and p=0.01, respectively). Polyfunctional CD8+CD107a+IFNg+IL2-TNF α - profile was increased in serology-positive (IP-S^{NEG} 0.01% vs IP-S_{POS} 0.13% and HC 0.03%; p=0.01 and p=0.50, respectively) and viral-load positive (IP-S^{POS}VL^{POS} 0.43% vs IP-S^{POS}VL^{NEG} 0.07% and HC 0.03%; p=0.03 and p=0.001, respectively) patients. Centralmemory cells were increased among serology-positive adults (IP-S^{NEG} 0.00% vs IP-S^{POS} 0.13% and HC 4.33%; p=0.58 and p=0.002, respectively). At the second timepoint, IP-S^{NEG} patients remained negative (first visit 0.01% vs second visit 0.00%, p=0.44). On the other hand, IP-S^{POS}VL^{POS} patients had cleared viral loads

and, subsequently, decreased polyfunctional CD8+CD107a+IFN γ +IL2-TNF α - cells (first visit 0.43% vs second visit 0.10%, p=0.81).

Conclusion: Polyfunctional CD8+ EBV-specific T-cell response allows detecting EBV previous contact in liver-transplant children. $CD8+CD107a+IFN_Y+IL2-TNF\alpha$ - is increased in patients with positive viral loads. Central memory CD4+ T-cell population more effectively determines prior EBV-exposure in adults.

KEYWORDS

liver transplantation, Epstein-Barr virus infections, cellular immunity, flow cytometry, cytokines, surface antigens

1 Introduction

The progressive improvement of immunosuppressive treatments to prevent graft rejection over the past few decades has contributed to the remarkable improvement in overall graft survival in children receiving liver transplants (1, 2). However, because of this effective immunosuppressive treatment, graft recipients experience a secondary state immunodeficiency, that renders them highly susceptible to infections (3, 4) and malignancies (5).

Most humans are infected with Epstein-Barr virus (EBV), which primarily targets epithelial and B cells, leading to both lytic and latent infections (6). Although EBV has oncogenic potential, it is usually controlled by the immune response. Adults are generally more competent in managing EBV than children, due to a more mature immune system and prior exposure to the virus.

However, in immunosuppressed recipients, EBV reactivation can lead to post-transplant lymphoproliferative disorder (PTLD) (1, 7), a condition characterized by uncontrolled proliferation of EBVinfected cells (8). In liver transplantation, the incidence of PTLD is remarkably higher in pediatric patients (6.3-15.0%) than adults (1.2-2.8%) (9, 10), partially due to their immunologically naïve status for the virus pre-transplantation (seroprevalence in children is approximately 50% vs. 90% in adults) (11) and the incidence of EBV primary infection under immunosuppression (12). In fact, several studies have reported that pre-transplant EBV-seronegative pediatric liver recipients are at higher risk (hazard ratio 12-18) (13) of developing PTLD (14–16).

The Healthcare Working Group of the European Reference Network on Pediatric Transplantation (ERN TransplantChild) has recently published the results of a cross-sectional survey evaluating PTLD strategies for diagnosis and treatment across several pediatric solid organ transplantation programs, from 9 different European countries (17). Over the 2012-2016 period, 1471 pediatric liver transplants were performed and 115 (7.8%) PTLD cases were diagnosed. PTLD preemptive strategies varied across different programs, but all of them included EBV DNA-load measurement by quantitative polymerase chain reaction (qPCR) as the main subrogated biomarker for EBV-specific immunity. Although EBV-load informative capacity appears to be widely integrated into daily clinical practice, its interpretation for PTLD diagnosis and surveillance is still controversial (18). Actually, no specific EBV viremia cutoff value has been defined to initiate preemptive treatment of PTLD (17). Regarding pediatric liver recipients, the association between high viral load and risk of PTLD development seems to be very poor (19), highlighting the necessity of new biomarkers.

Several techniques have been previously validated in different transplantation settings to estimate T-cell EBV response, being the most standardize one the detection of interferon gamma (IFN χ) by either enzyme-linked immunospot (ELISpot) (20–23) or enzyme-linked immunosorbent (ELISA)-based (QuantiFERON[®]) assays (24–29). Other promising techniques involve the identification by flow cytometry of antigen-specific cells using mayor histocompatibility complex class I and class II multimers (23, 30–32), intracellular cytokine staining (ICS) (30, 31, 33–42) or activation-induced marker staining (AIMS) (43, 44).

Research on the T-cell compartment against EBV in pediatric liver recipients is scarce and primarily utilizes ELISpot and tetramer assays. There seems to be a correlation between immunosuppression doses and frequency of EBV-specific cells by ELISpot (45–47). Similar results were reported prospectively, measuring cellular response by tetramers (46). Nevertheless, neither tetramers nor dextramers are apparently effective in discriminating transplanted patients according to EBV viral load (32, 48).

Ning et al. used ICS to measure specific T-cell response in two pediatric liver recipients with detectable viral loads and diagnosis of PTLD. Those patients presented a reduction in T-cell polyfunctionality, with an increment in the expression of CD107a and tumor necrosis factor alpha (TNF α) (49). Another study, examining 20 pediatric-transplanted patients (7 liver-graft recipients), presented findings on T-cell response by ICS. Authors reported a significant increment in EBV-specific T cells in PTLD patients during rituximab treatment, which correlated with a reduction in viral load and subsequent control of EBV by T-cell responses following B-cell recovery (50). To our knowledge, the use of AIMS in this field has never been reported, although OX40 (CD134) has been previously defined as a potential biomarker of T-cell activation status in various types of transplants (51, 52). The aim of this cross-sectional study is to characterize the specific helper and cytotoxic T-cell response to EBV in a cohort of pediatric liver recipients by both ICS and AIMS flow-cytometry methods, and compare it with a cohort of EBV-seropositive healthy adult controls (HC). Furthermore, we aim to identify cellular profiles that allow the discrimination of liver recipients according to both their EBV serological and viral-load status. We hypothesize that pediatric liver recipients controlling EBV will exhibit higher percentages of EBV-specific T cells compared to non-controllers, while also displaying specific cellular profiles.

2 Materials and methods

2.1 Patients and samples

Our cross-sectional study included 38 immunosuppressed pediatric patients (IP) at University Hospital La Paz, who received a liver graft between March 2018 and November 2022, and 25 EBV seropositive HC. All patients gave informed consent, approved by the ethics committee of our institution (reference PI-4000).

Demographic and clinically relevant information from each patient were collected (Table 1). PTLD diagnosis was based on histopathologic criteria. Transplant indication was categorized in five groups (Table 1) (53).

Specific T-cell response against EBV was assessed at two different timepoints, determined by patient availability (median time 3.7 months interquartile range [IQR] 3.2-4.9 between visits). EBV serology and viral load were measured in parallel with each immune response assessment for every patient.

Based on their serological status at the first visit, patients were categorized into EBV seronegative (IP-S^{NEG}) and EBV seropositive (IP-S^{POS}) individuals; the latter group was further classified according to EBV viral load into negative (IP-S^{POS}VL^{NEG}) and positive (IP-S^{POS}VL^{POS}). At the time of the second visit, patients were reclassified based on their updated serological and viral load status at that time. Due to sample exclusions for technical reasons, the immune response was not measured in all samples. ICS was performed to 38/38 (100%) and 28/38 (74%) patients at first and second visit, respectively. AIMS was performed to 27/38 (71%) and 29/38 (76%) patients at first and second visit, respectively, although results for both timepoints were available for only 21 individuals. Among HC participants, ICS and AIMS were successfully performed to 24/25 (96%) and 20/25 (84%) samples, respectively.

Heparinized blood samples from all individuals were collected to isolate peripheral blood mononuclear cells (PBMCs) by density gradient.

2.2 Intracellular cytokine staining

Analysis of specific T-cell response to EBV by ICS was assessed as described by Lovelace and Maecker (54). Details of the method are provided in the Supplementary Material.

T-cell responses were further categorized as monofunctional, when only one response marker was displayed (CD107a, IFN_x,

Interleukin 2 [IL2] or $TNF\alpha$), and polyfunctional, when more than one response marker was expressed. Integrated median fluorescence intensity (iMFI) was calculated for each response marker, multiplying MFI by the frequency of the corresponding specific population.

Unstimulated PBMCs background was subtracted from all test samples to obtain the frequency of EBV-specific T lymphocytes. Gating strategy is displayed in Supplementary Figure S1A.

2.3 Activation-induced cell marker staining

Analysis of AIMS was performed by flow cytometry. Details of the method are provided in the Supplementary Material.

CD4+ T lymphocytes were distributed in naïve (Tn, CD27+ CD45RO-), effector (Teff, CD27-CD45RO-), central memory (Tcm, CD27+CD45RO+) and effector memory (Tem, CD27-CD45RO+) subpopulations. Memory compartment was calculated by the sum of Tcm and Tem subpopulations.

Unstimulated PBMCs background was subtracted from all test samples to obtain the frequency of EBV-specific T lymphocytes. Gating strategy is displayed in Supplementary Figure S1B.

2.4 Immunophenotype analysis

Immunophenotype of T, B, natural killer (NK) and natural killer T (NKT) lymphocytes was performed by multiparametric flow cytometry. Details of the method are provided in the Supplementary Material.

2.5 EBV viral load measurement

EBV viral loads were quantified in whole blood by a specific qPCR assay following manufacturer's instructions (RealStar[®] EBV PCR-Kit 1.0, Altona). Results were informed in International Units per milliliter (IU/mL). The negative group for viral load comprised exclusively patients with zero IU/mL.

2.6 EBV serology

EBV serological status was determined by a chemiluminescent microparticle immunoassay. Presence of IgM antibodies against viral capsid antigen (VCA) and/or IgG antibodies against VCA and nuclear antigen (EBNA) were measured following manufacturer's instructions (Abbott, Germany). EBV seropositive status was defined by the positivity of at least one of the analyzed antibodies.

2.7 Statistics

Descriptive data are presented as median with IQR. Categorical data are presented as absolute number and proportion (%). The software package Prism 8 (GraphPad, USA) was used for statistical analysis. Significance of differences comparing frequencies was determined by Pearson χ 2-test and by *t* test or analysis of variance (Mann Whitney or Kruskal-Wallis tests) when comparing median values. Median frequencies between timepoints were compared by

Characteristics	HC (n=25)	IP-S ^{POS} (n=32)	IP-S ^{NEG} (n=6)	P-value	HC (n=25)	IP-S ^{POS} VL ^{POS} (n=8)	IP-S ^{POS} VL ^{NEG} (n=24)	P-value
Sex, n (%)				0.10				0.07
Male	17 (68)	13 (41)	4 (67)		17 (68)	2 (25)	11 (46)	
Female	8 (32)	19 (59)	2 (33)		8 (32)	6 (75)	13 (54)	
Age, years (IQR)	54 (41-62)	5 (3-9)	5 (3-6)	<0.001	54 (41-62)	3 (2-4)	5 (4-9)	<0.001
Donor sex, n (%)				0.62				0.34
Male		8 (25)	3 (50)			1 (13)	7 (29)	
Female		13 (41)	2 (33)			5 (63)	8 (33)	
Donor age, years (IQR)		28 (19-34)	27 (15-38)			30 (26-31)	27 (16-34)	
Type of donor, n (%)				0.25				0.52
Living donor		10 (31)	4 (67)			3 (38)	7 (29)	
Deceased donor - whole graft		9 (28)	1 (17)			1 (13)	8 (33)	
Deceased donor - split graft		13 (41)	1 (17)			4 (50)	9 (38)	
ABO compatibility, n (%)				>0.99				>0.99
Compatible		25 (78)	6 (100)			7 (88)	18 (75)	
Incompatible		3 (9)	0 (0)			1 (13)	2 (25)	
Indication for transplantation, n (%)				0.23				0.31
Cholestasis/biliary atresia		19 (59)	3 (50)			6 (75)	13 (54)	
Metabolic diseases		7 (22)	2 (33)			0 (0)	7 (29)	
Cirrhosis (other)		2 (6)	0 (0)			1 (13)	1 (4)	
Severe acute liver failure		2 (6)	0 (0)			1 (13)	1 (4)	
Liver tumours		2 (6)	0 (0)			0 (0)	2 (8)	
Metabolic diseases and liver tumours		0 (0)	1 (17)			0 (0)	0 (0)	
Time since transplantation, months (IQR)		40 (31-47)	48 (26-53)	0.47		32 (12-36)	44 (35-47)	0.002
Immunosuppressive treatment, n (%)				0.47				0.70
CE		5 (16)	0 (0)			2 (25)	3 (13)	
TAC		2 (6)	0 (0)			0 (0)	2 (8)	

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Characteristics	HC (n=25)	IP-S ^{POS} (n=32)	IP-S ^{NEG} (n=6)	P-value	HC (n=25)	IP-S ^{POS} VL ^{POS} (n=8)	IP-S ^{POS} VL ^{NEG} (n=24)	P-value
CE+TAC		22 (69)	6 (100)			5 (63)	17 (71)	
CE+TAC+MMF		3 (9)	0 (0)			1 (13)	2 (88)	
TAC blood levels, ng/ml (IQR)		3.4 (2.7-4.5)	4.4 (3.8-6.7)	0.07		4.1 (3.3-6.5)	3.3 (2.4-4.4)	0.18
EVB-serology pre-transplantation, n (%)				0.03				0.65
Positive		14 (56)	0 (0)			4 (50)	10 (42)	
Negative		13 (52)	6 (100)			2 (25)	11 (46)	
CMV-serology pre-transplantation, n (%)				0.66				>0.99
Positive		15 (47)	2 (33)			4 (50)	11 (46)	
Negative		14 (44)	4 (67)			3 (38)	11 (46)	
PTLD diagnosis, n (%)		5 (16)	0 (0)	0.57		2(25)	3(13)	0.58
Time since diagnosis of PTLD, days (IQR)		35 (12-133)	NA	NA		8 (6-10)	133 (84-311)	0.20
Lymphocyte number x10 ³ , cells/µL (IQR)		3.0 (2.4-3.9)	4.0 (2.9-4.3)	0.24		2.7 (2.2-3.3)	3.0 (2.4-4.7)	0.48
Immune phenotype, % (IQR)								
CD3+ T lymphocytes		78 (72-85)	72 (65-76)	0.20		74 (69-76)	83 (75-87)	0.11
CD4+ T lymphocytes		45 (38-55)	42 (37-49)	0.55		46 (34-55)	43 (39-56)	>0.99
CD8+ T lymphocytes		22 (17-30)	21 (12-29)	0.41		18 (14-28)	24 (20-36)	0.20
B lymphocytes		8 (6-14)	11 (8-14)	0.55		15 (7-20)	7 (5-9)	0.11
NK lymphocytes		10 (6-16)	19 (10-23)	0.13		13 (8-17)	7 (5-16)	0.33
NKT lymphocytes		0.7 (0.6-1.8)	0.6 (0.5-0.9)	0.20		0.8 (0.5-2.5)	0.7 (0.6-1.8)	0.80

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CE, corticosteroids; CMV, Cytomegalovirus; EVB, Epstein-Barr virus; HC, healthy controls; IQR, interquartile range; MMF, mycophenolate mofetil; NA, not applicable; NK, natural killer; NKT, natural killer T; PTLD, post-transplant lymphoproliferative disorder; TAC, tacrolimus.

Statistically significant p-values are indicated in bold.

05

Wilcoxon signed-rank test. Correlation between ICS and AIMS results was assessed by linear regression. P-values under 0.05 were considered significant.

3 Results

3.1 Demographic and clinical characteristics

A first classification of our cohort (n=38) was performed according to EBV serological status: IP-S^{NEG} (6/38, 16%) and IP-S^{POS} (32/38, 84%). A second distribution of EBV-seropositive patients (n=32) was made according to EBV viral loads: IP-S^{POS}VL^{POS} (8/32, 25%) and IP-S^{POS}VL^{NEG} (24/32, 75%).

First analysis of our cohort (Table 1) showed statistically significant differences when comparing age (HC adults vs. IP) and EBV-serology pre-transplantation (seropositive vs. seronegative). Interestingly, 52% of post-transplant EBV seropositive patients were negative pre-transplantation. When time since transplantation was analyzed, we observed that IP with positive EBV viral loads had been more recently transplanted (IP-S^{POS}VL^{POS} 32 months IQR 12-36 vs. IP-S^{POS}VL^{NEG} 44 months IQR 35-47, p=0.002). Percentages of T, B and NK subpopulations were similar among groups (Table 1).

3.2 EBV-specific T-cell response by intracellular cytokine staining

At first visit, EBV-specific %CD3+ T cells by ICS was higher in HC than IP groups (IP-S^{NEG} 0.03% vs. IP-S^{POS} 0.04% and HC 0.06%; p=0.41 and p=0.24, respectively), but differences did not reach statistical signification. Nonetheless, all three groups showed similar positive-control stimulation (IP-S^{NEG} 0.75%, IP-S^{POS} 0.79% and HC 0.45%; p=0.08), indicating that cellular response *in vitro* was not impaired by immunosuppression. Positive-control responses remained comparable when dividing mono/polyfunctional CD4+ and CD8+ T-cell subpopulations (data not shown).

However, when splitting EBV-specific response between T-cell subpopulations (Figure 1A), statistically significant differences were observed. Median frequency of monofunctional CD4+ and CD8+ T cells was higher than polyfunctional cells in all groups (Figure 1A). Interestingly, although monofunctional responses were detected in higher frequencies, only polyfunctional CD8+ T cells significantly discriminated EBV seronegative patients from seropositive HC and IP groups (IP-S^{NEG} 0.00% vs. IP-S^{POS} 0.04% and HC 0.02%; p=0.01 and p<0.001, respectively; Figure 1A).

For a more detailed analysis, the different EBV-specific CD8+ Tcell profiles were analyzed (Figures 1B, C). Three different CD8+ polyfunctional profiles differentiated seronegative recipients from both HC and IP seropositive individuals: CD8+CD107a+IFNγ+ IL2+TNFa+, CD8+CD107a+IFNγ+IL2-TNFa+ and CD8+CD107a-IFNγ+IL2-TNFa+ (Figure 1B). Furthermore, seropositive IP had higher frequencies of EBV-specific CD8+ polyfunctional cells than seronegative IP in two other subsets: CD8+CD107a+IFNγ+IL2-TNFa- and CD8+CD107a+IFNγ-IL2-TNFa+. The profile CD8 +CD107a+IFNγ+IL2-TNFa- was the most frequent one (0.13%).

Precisely, this CD8+CD107a+IFNy+IL2-TNFa- subset significantly discriminated patients with positive viral loads from the rest of individuals (IP-SPOSVLPOS 0.43% vs. IP-SPOSVLNEG 0.07% and HC 0.03%; p=0.03 and p=0.001, respectively; Figure 1C). IP-S^{POS}VL^{POS} patients also showed higher %CD8+CD107a+IFNy-IL2+ TNFa- compared to HC, but not to IP-S^{POS}VL^{NEG} group (Figure 1C). Remarkably, polyfunctional response was more intense than monofunctional response in seropositive individuals (Supplementary Table S1). All three cytokine markers IFNy, IL2 and TNFa had significantly higher iMFI values in polyfunctional response, both in CD4+ and CD8+ T cells, whereas CD107a only showed higher intensity in polyfunctional CD8+ T cells (Supplementary Table S1). In line with our previous results, CD8+ polyfunctional subpopulation allowed discriminating seropositive from seronegative status, according to iMFI from all four response markers (Table 2). However, regarding EBV viral load, only total (mono and polyfunctional) CD107a iMFI on CD8+ T cells significantly differentiated IP with positive viral loads from the other groups (IP-S^{POS}VL^{POS} 123,398 vs. IP-S^{POS}VL^{NEG} 20,708 and HC 21,207; p=0.03 and p=0.01, respectively).

To simplify cytometry panels, we studied whether CD8+ T cells expressing only CD107a and IFN γ markers could be distinctive, regardless of other cytokines. As expected, %CD8+CD107a+IFN γ + T cells were significantly higher among seropositive individuals compared to seronegative IP (IP-S^{NEG} 0.02% vs. IP-S^{POS} 0.37% and HC 0.12%; p=0.002 and p=0.04, respectively). However, considering EBV viral load, although the frequency of specific cells was also increased within positive IP group, differences were significant compared to controls, but not to negative patients (IP-S^{POS}VL^{POS} 0.70% vs. IP-S^{POS}VL^{NEG} 0.19% and HC 0.12%; p=0.10 and p=0.03, respectively).

Finally, we explored whether CD3+IFN χ + T cells, the main population targeted by other methods, allowed discrimination of serology and/or viral-load status. While frequency of CD3+IFN χ + T cells was insufficient for serology discrimination (IP-S^{NEG} 0.10% vs. IP-S^{POS} 0.31% and HC 0.12%; p=0.31 and p>0.99, respectively), iMFI was significantly higher in seropositive than seronegative individuals (IP-S^{NEG} 182 vs. IP-S^{POS} 3,628 and HC 1,380; p=0.003 and p=0.01, respectively). When comparing viral-load status no statistically differences were reached either with CD3+IFN χ + frequency (IP-S^{POS}VL^{POS} 0.37% vs. IP-S^{POS}VL^{NEG} 0.24% and HC 0.12%; p>0.99 and p=0.40, respectively) or iMFI (IP-S^{POS}VL^{POS} 3,834 vs. IP-S^{POS}VL^{NEG} 2,809 and HC 1,380; p=0.62 and p=0.82, respectively).

3.3 EBV-specific T cell response by activation-induced cell marker staining

At first visit, differences were found in EBV-specific T-cell frequencies by AIMS between HC and IP groups (Figures 2A, B). As expected, CD4+ Tn EBV-specific subset was the lowest within each group, compared to Teff and memory compartment (Figures 2A, B). On the other hand, seropositive groups (Figure 2A) had higher frequency of EBV-specific memory compartment than Teff cells (IP- S^{POS} memory compartment 0.99% vs. Teff 0.04%, p=0.12; HC memory compartment 8.24% vs. Teff 0.37%, p<0.001), although differences were statistically significant only within HC group. Interestingly,



FIGURE 1

Specific T-cell response to Epstein-Barr virus (EBV) performed by intracellular cytokine staining of CD107a, IFN_X, IL2 and TNFa markers in EBV seropositive healthy adult controls (HC) and immunosuppressed pediatric liver recipients (IP). (A) Median frequency of EBV-specific monofunctional (one response marker) or polyfunctional (more than one response markers) CD4+ and CD8+ T cells in HC and IP groups, the last one categorized according to their positive/negative serological status (IP-S^{POS} and IP-S^{NEG}, respectively). (B) Median frequency of EBV-specific polyfunctional CD8+ T-cell subpopulations according to the different response markers in HC, IP-S^{POS} and IP-S^{NEG} groups. (C) Median frequency of EBV-specific polyfunctional CD8+ T-cell subpopulations in HC and IP-S^{POS} groups, the last one segregated according to their positive/negative EBV viral-load status (IP-S^{POS}VL^{POS} and IP-S^{POS}VL^{NEG}, respectively). The same HC group was used as the control group in panels (A–C). Significance levels are denoted as *p-value<0.05, **p-value<0.001.

TABLE 2 Median of the integrated median fluorescence intensity (iMFI) for each marker (CD107a, IFN $_{Y}$, IL2 or TNF α) in both polyfunctional CD4+ and CD8+ T cells in EBV-seropositive adult healthy controls (HC) and immunosuppressed pediatric liver-transplanted patients, categorized as positive/ negative serology status (IP-S^{POS} vL^{POS} and IP-S^{NEG}, respectively) or positive/negative viral load status (IP-S^{POS} vL^{POS} and IP-S^{NEG}, respectively).

iMFi		HC	IP-S ^{POS}	IP-S ^{NEG}	Dualua	НС	IP-S ^{POS} VL ^{POS}	IP-S ^{POS} VL ^{NEG}	Duchuc
T-cell subset	Parameter	(n=24)	(n=32)	(n=6)	P-value	(n=24)	(n=8)	(n=24)	r-value
Polyfunctional CD4+ T cells	CD107a+	1,060	1,813	2,588	0.43	1,060	2,651	1,813	0.35
	IFNy+	475	179	49	0.07	475	181	179	0.13
	IL2+	122	93	237	0.69	122	95	93	0.93
	TNFα+	1,248	1,422	1,838	0.84	1,248	1,705	1422	0.85
Polyfunctional CD8+ T cells	CD107a+	17,538	42,771	373	0.002	17,538	105,001	18,628	0.03
	IFNy+	3,366	7,714	36	0.002	3,366	10,042	5,228	0.34
	IL2+	377	424	0	0.001	377	329	488	1.00
	TNFα+	7,370	8,307	0	0.001	7,370	11,509	7,391	0.86

Statistically significant p-values are indicated in bold.

seronegative IP showed higher frequencies of Teff than memory EBVspecific cells (memory compartment 0.00% vs. Teff 0.25%, p=0.14), although differences did not reach statistical signification.

Compared to IP seronegative group, percentages of EBV-specific cells were higher in HC group for Tn (0.14% vs. IP-S^{NEG} 0.00%, p=0.03), Tem (4.31% vs. IP-S^{NEG} 0.00%, p=0.01) and Tcm

(4.33% vs. IP-S^{NEG} 0.00%, p=0.002) subpopulations (Figure 2A). However, comparing frequencies between seropositive and seronegative IP did not yield any statistically significant differences. Again, positive-control stimulation in CD4+ T cells was comparable by AIMS (IP-S^{NEG} 24.52%, IP-S^{POS} 32.67% and HC 26.42%; p=0.62).



FIGURE 2

Specific T-cell response to Epstein-Barr virus (EBV) performed by activation-induced marker staining (CD4+CD25+CD134+) in EBV seropositive healthy adult controls (HC) and immunosuppressed pediatric liver recipients (IP). **(A)** Median frequency of EBV-specific CD4+CD25+CD134+ T cells in HC and IP groups, the last one categorized according to their positive/negative serological status (IP-S^{POS} and IP-S^{NEG}, respectively). **(B)** Median frequency of EBV-specific CD4+CD25+CD134+ cells in HC and IP-S^{POS} groups, the last one segregated according to positive/negative EBV viral-load status (IP-S^{POS} VL^{POS} and IP-S^{POS} VL^{NEG}, respectively). The same HC group was used as the control group in both panels **(A, B)**. Significance levels are denoted as *p-value<0.05, **p-value<0.001.

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We next studied seropositive IP grouped by viral-load status (Figure 2B) and observed that %CD4+ Tcm cells was significantly higher in HC than seropositive IP with detectable EBV (IP-S^{POS}VL^{POS} Tcm 0.13% vs. HC Tcm 4.32%, p=0.02). Interestingly, median %CD4+ Tem cells in seropositive IP-S^{POS}VL^{POS} was higher than in IP-S^{POS}VL^{NEG} (0.90% vs. 0.34%, respectively; p>0.99), although no significant differences were found.

Finally, we further investigated the potential correlation between the parameters defined by ICS (%CD8+CD107a+IFNy+IL2-TNFa-) and AIMS (%CD4+CD134+CD25+ Tcm), which effectively distinguished positive from negative individuals. We noted that those parameters exhibited no correlation (data not shown), likely due to the comparison involving distinct T-cell subpopulations. Thus, a potential correlation between EBV-specific CD4+ T cells by ICS (% and iMFI CD4+INFy) and AIMS (%CD4+CD134+CD25+ Tcm) techniques was next sought, although we did not observe any correlation, by either frequency or iMFI (data not shown). However, within HC group, after excluding the data from one individual lacking EBV-specific CD4+ CD134+CD25+ memory T cells, a significant correlation between iMFI CD4+INFy by ICS and %CD4+CD134+CD25+ by AIMS ($r^2 = 0.24$ and p=0.04, Figure 3) was found.

3.4 EBV-specific T cell response at two different timepoints

EBV-specific response was measured on a second visit by both techniques. Serological and viral-load status was re-evaluated and patients were reclassified accordingly. Tacrolimus blood levels remained similar at first and second timepoints in all three groups (data not shown).

At the second timepoint, identical results to those reported at the first visit were found when comparing frequencies of EBV-specific cells detected by ICS and AIMS. We confirmed that only polyfunctional CD8+ specific T cells significantly discriminated EBV seronegative patients from seropositive individuals (IP-S^{NEG} 0.00% vs. IP-S^{POS} 0.03% and HC 0.02%; p=0.01 and p=0.02, respectively). Likewise, %EBV-specific cells from HC were higher than those detected in seronegative patients for Tn (0.14% vs. 0.01%, p=0.05), Tem (4.31% vs. 0.00%, p=0.01) and Tcm (4.33% vs. 0.00%, p=0.003) subpopulations.

Regarding IP-SPOSVLPOS group, all recipients with detectable EBV at first timepoint cleared viral loads at the second visit (median time 3.5 months IQR 2.2-5.3 between timepoints). Consequently, median % CD8+CD107a+IFNy+IL2-TNFa- (first visit 0.26% vs. second visit 0.10%, p=0.81; Figure 4A) and %CD4+CD25+CD134+ Tcm cells (first visit 0.39% vs. second visit 0.14%, p=0.88; Figure 4B) decreased at the second timepoint, although no significant differences were found.

On the other hand, only two IP-SPOSVLNEG patients at first timepoint had detectable EBV at the second visit (black arrows in Figures 4C, D) (median time 3.7 months IQR 3.3-4.8 between timepoints). Interestingly, significant differences were found by ICS (first visit 0.08% vs. second visit 0.06%, p=0.04; Figure 4C), but not by AIMS (first visit 0.00% vs. second visit 0.76%, p=0.31; Figure 4D). Frequency of CD8+CD107a+IFNy+IL2-TNFa- cells decreased or remained similar for all patients, except for the individual who tested positive for EBV at the second visit, whose frequency increased from 0.06% to 0.09% (black arrow in Figure 4C). This patient's frequency of specific CD4+CD25+CD134+ Tcm cells also increased from 0.00% to 1.06% (black arrow in Figure 4D). Nevertheless, that variation in the specific response by AIMS was not observed in the other patient who tested positive for viral load at the second timepoint (black arrow in Figure 4D). One IP-S^{POS}VL^{NEG} patient had his immunosuppression regimen changed between visits, incorporating mycophenolate to his treatment with corticosteroids and tacrolimus. Interestingly, frequency



by intracellular cytokine staining and EBV-specific CD4+CD25+CD134+ central memory T (Tcm) cells measured by activation-induced marker staining

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FIGURE 4

Frequency of Epstein-Barr virus (EBV)-specific CD8+CD107a+IFNy+IL2-TNFa- T cells (**A**, **C**, **E**) and CD4+CD25+CD134+ central memory T (Tcm) cells (**B**, **D**, **F**) measured at two different timepoints (1^{st} and 2^{nd} visit) in immunosuppressed pediatric liver recipients (IP). Patients were classified at their first visit according to their positive/negative EBV serological and viral-load status: IP-S^{POS}VL^{POS} (**A**, **B**), IP-S^{POS}VL^{NEG} (**C**, **D**) and IP-S^{NEG} (**E**, **F**) groups. At the second visit, their updated serological and viral-load status is represented by rhomboid, circular or square markers, respectively. Changes in their classification at second visit are marked with a black arrow. Patients that suffered changes in his immunosuppression regimen at second visit are marked with a white arrow. Significance levels are denoted as *p-value<0.05, **p-value<0.01 and ***p-value<0.001.

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of EBV-specific T cells by ICS decreased from 0.44% to 0.25%, while frequency of CD4+CD25+CD134+ Tcm cells remained at 0.00% (white arrows in Figures 4C, D).

Finally, all seronegative patients (Figures 4E, F) kept their negative serological status at the second visit (median time of 3.9 months IQR 3.6-4.0 between timepoints). Consequently, no differences in EBV-specific response were found, either by ICS (0.01% vs. 0.00%, p=0.44; Figure 4E) or AIMS (0.00% vs. 0.00%, p>0.99; Figure 4F).

4 Discussion

In our study, we have first explored EBV-specific T-cell response combining ICS and AIMS techniques in a cohort of pediatric liver transplanted recipients. We found significant differences in polyfunctional CD8+ T-cell response between EBV-seronegative and seropositive individuals, and among patients with positive and negative viral loads.

Firstly, we found higher percentages of monofunctional than polyfunctional EBV-specific T cells. This could be attributed to crossreactivity resulting from heterologous immunity (55), although recent studies state that it is less generalized than previously reported (56). Since we have also confirmed that monofunctional responses showed lower iMFI (49), we can attribute this result to a potential unspecific bystander activation *in vitro*. The presence of monofunctional EBV-specific cells among seronegative patients provides additional support for that hypothesis.

The predominance of CD8+ over CD4+ T-cell responses in controlling EBV infection is well documented (49, 57). Although both CD4+ and CD8+ T cells show polyfunctional responses after primary infection, only the CD8+ polyfunctional subset increases over time (58). Accordingly, we observe that polyfunctional CD8+ T cells are significantly increased in seropositive individuals (Figure 1B). Remarkably, we expected a reduction in the frequency of this population in patients with detectable viral loads, in concordance with Ning et al., who demonstrated this in two pediatric liver recipients with PTLD (49). Conversely, IP-S^{POS}VL^{POS} in our cohort showed high %CD8+ EBV-specific cells (Figure 1C), including 2 patients who were studied at the time of PTLD diagnosis.

The increment of CD8+CD107a+IFNy+IL2-TNFa- in recipients with detectable EBV viral loads reflects the critical role of cytotoxicity, as indicated by CD107a, and the antiviral function of IFNy in the response against EBV. Frequent EBV reactivation might inflate this cell compartment, exhausting and rendering cells dysfunctional, thus requiring larger numbers to control the virus (59). Previously suggested explanation for pediatric graft recipients carrying chronically high EBV loads involves an exhausted phenotype (23). We did not include exhaustion markers in our study, but we consider that they would be useful to better describe viral responses in future investigations. Other protocols, such as expanding T cells in the presence of EBV peptides for 7-10 days before analysis (60), could also be applied. However, we chose shorter incubation times (54), as this approach was more compatible with the workflow of our routine laboratory.

We further evaluated CD107a combined with IFNy, as exclusive response markers to distinguish patients with positive viral loads.

While %CD8+CD107a+IFN χ + helped in serostatus discrimination, it did not show significant differences in EBV viral loads, likely due to our limited sample size. Similar results were reported by Wilsdorf et al., who measured intracellular IFN χ after EBV-peptide stimulus in pediatric transplanted patients with PTLD (4/16 liver-graft recipients) or positive viral-loads (3/4 liver-graft recipients) and 18 HC. Median %CD4+IFN χ + and %CD8+IFN χ + was higher in recipients with EBV reactivation, yet they did not find significant differences either (50).

In our cohort, %CD3+IFN χ + cells do not differentiate seropositive from seronegative patients, contrary to prior studies using ELISpot (47). Instead, we found that the intensity of IFN χ response in CD3+ T cells effectively differentiated these two groups. While no equivalent parameter to iMFI has been described in ELISpot assays, mean spot size could be comparable (61). Our findings may be influenced by the constraints of our sample size, but conducting additional studies to explore iMFI further would be valuable.

On the other hand, the use of AIMS to measure specific viral response is not extended, although it has been validated in HC for Varicella-Zoster Virus, Cytomegalovirus, EBV (44) and Hepatitis C (43). Regarding EBV, it seems feasible to distinguish seronegative from seropositive individuals measuring %CD4+CD134+CD25+ specific-cells (44). We confirmed these findings by examining Tcm and Tem CD4+ subsets, consistent with the predominant memory CD4+ response to EBV (62). Interestingly, we detected specific CD4+ Teff cells in seronegative individuals, probably reflecting antigen exposure in some patients (63), although we cannot exclude unspecific activation, since percentages were similar among groups. The increase of EBV-specific CD4+ Tn in HC (Figure 1A) has been previously reported as a genuine memory population transitioning to express naïve surface markers (62). This stem memory T-cell population (CCR7+CD27+CD45RO-) shares some features with Tn and requires staining with specific markers (CD95) for proper selection (64). These cells emerge rapidly post-antigen exposure, transitioning into effector cells, while retaining self-renewal and multipotent abilities, making them ideal for adoptive T-cell therapies, including EBV infection in transplant recipients (65, 66).

Due to the age gap, memory response in HC is the highest, reflecting repeated exposures to EBV antigens over their lifetimes, which expands the clonal repertoire against the virus (23). Interestingly, Tem specific subset is increased in patients with active viral replication, consistent with findings by Amyes et al. (67). They observed a primary burst of CD4+ Teff cells in response to EBV, persisting throughout the chronic phase of infection. However, our stimulation with a cocktail of lytic and latent EBV peptides does not differentiate between viral phases (58, 67).

Regarding correlation between ICS and AIMS, Sadler et al. demonstrated that EBV-specific production of IFNy significantly correlated with %CD4+CD134+CD25+ cells in HC (44). While we did not replicate this result, we found %CD4+CD134+CD25+ cells from memory compartment correlated with CD4+ response measured by IFNy iMFI, confirming AIMS reliability to infer specific CD4+ Tcell response. Further studies on CD8+ T-cell activation markers, such as CD38 and HLA-DR (44), are recommended.

At the second visit, we confirmed that polyfunctional CD8+ EBV-specific T-cell response detects serology and viral-load positive individuals. Furthermore, we replicated our findings on CD4+ Tcm population in adults. Compared to first visit, we observed changes in EBV-specific cellular response among patients with viral-load status shifts, significant in the largest sample group (Figure 4C). Interestingly, tacrolimus blood levels remained similar between visits, questioning the effect of immunosuppression treatment. Positive control stimulus elicited a similar reaction in HC and immunosuppressed patients, suggesting that anti-CD3/28 beads override immunosuppression. Similarly, Arasaratnam et al. found comparable IFNy production by Staphylococcal enterotoxin B in pediatric liver recipients post-immunosuppression (45).

On the other hand, other authors observed that immunosuppression treatment modifications for PTLD management lead to changes in frequency of EBV-specific cells detected by ELISpot (45–47). Moreover, OX40 has been postulated as an indicator of the immunosuppressive status of patients after stem cell transplantation (68), although Lamb et al. reported a recipient of stem cell transplantation whose changes in the immunosuppressive treatment did not reflect differences in %CD4+CD134+ cells (69). To elucidate the potential use of ICS or AIMS in evaluating immunosuppressive status of transplanted patients further prospective studies are needed.

In conclusion, our study reveals significant insights into EBVspecific T-cell responses in pediatric liver transplant recipients. We demonstrated that polyfunctional CD8+ T cells were markedly increased in seropositive individuals, underscoring their role in controlling EBV infection. Despite a limited sample size, our findings support the utility of CD107a/IFNy response markers for distinguishing EBV serostatus and viral load. Additionally, OX40 proved reliable in assessing CD4+ memory responses, suggesting its potential for broader application in evaluating antiviral immunity. Further prospective research is recommended to refine our understanding of EBV-specific T-cell dynamics in transplant recipients.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material. Further inquiries can be directed to the corresponding author.

Ethics statement

The studies involving humans were approved by University Hospital La Paz ethics committee. The studies were conducted in accordance with the local legislation and institutional requirements. Written informed consent for participation in this study was provided by the participants' legal guardians/next of kin.

Author contributions

RC: Conceptualization, Data curation, Investigation, Methodology, Writing – original draft, Writing – review & editing. AT: Investigation, Methodology, Writing – review & editing. LM: Investigation, Methodology, Writing – review & editing. KR: Investigation, Methodology, Writing – review & editing. EF: Writing – review & editing, Conceptualization, Investigation. LH: Writing – review & editing, Conceptualization, Investigation. GM: Writing – review & editing, Investigation. ML: Writing – review & editing, Investigation. AG: Writing – review & editing, Methodology. AM: Writing – review & editing, Investigation. EL: Supervision, Writing – original draft, Writing – review & editing, Conceptualization, Investigation. ES: Supervision, Writing – original draft, Writing – review & editing, Conceptualization.

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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.

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Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fimmu.2024.1479472/full#supplementary-material

SUPPLEMENTARY FIGURE 1

Flow-cytometry gating strategy for (A) intracellular cytokine staining or (B) activationinduced marker staining in a representative healthy control. Cells were left unstimulated (NC) or stimulated using a positive control (PC) or Epstein-Barr virus peptides (EBV).

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