

## Laboratory diagnostic testing for cytomegalovirus infection in solid organ transplant patients

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Human cytomegalovirus (CMV) infection, which is one of the most common complications in transplant recipients, increases the risk of graft loss and rejection. Laboratory strategies for diagnosing CMV infection rely on the measurement of viral DNAemia and CMV-specific cell-mediated immunity (CMV-CMI). The CMV quantitative nucleic acid amplification test (QNAT) enabled the spread of preemptive therapy and prompted recommendations for surveillance, diagnosis, and monitoring. Despite the implementation of the World Health Organization international standard for calibration, variability of QNAT persists due to technical issues. CMV immunoglobulin G serology is the standard method for CMV immune screening of transplant candidates and donors. Assays for CMV-CMI play an important role in helping to predict the risk and to develop an individualized CMV management plan. Genotypic testing for resistance is needed when drug-resistant CMV infection is suspected. Here, we review the state of the art of laboratory tests for CMV infection in solid organ transplantation.

**Keywords:** Transplantation; Cytomegalovirus; Cell-mediated immunity; Serology; Viral load

### INTRODUCTION

Cytomegalovirus (CMV) infection is generally asymptomatic and relatively benign in the immunocompetent population [1]. However, CMV infection in solid organ transplantation (SOT) is significantly associated with increased morbidity and mortality [2,3]. In SOT, CMV is transmitted via a transplanted organ, and the risk of CMV infection is increased if the recipient does not have pre-existing CMV-specific cell-mediated immunity (CMV-CMI) and the graft has a certain amount of lymphoid tissue [4,5]. Long-

term immunosuppressive treatment to prevent rejection increases the risk of viral reactivation in seropositive recipients who had previously acquired a latent infection [6].

The worldwide CMV seroprevalence has been estimated to be approximately 83% in the general population [7]. A high seroprevalence of CMV has been reported in Korea, and a recent study estimated the overall CMV immunoglobulin G (IgG) seropositivity rate to be 94% [8]. Therefore, Koreans were generally considered seropositive donors and recipients (D+/R+), who are at intermediate risk for CMV infection and/or latent virus reactivation after trans-

## HIGHLIGHTS

- Cytomegalovirus (CMV) quantitative nucleic acid amplification test is a rapid and sensitive method of diagnosing CMV infection and is preferred for CMV surveillance to guide preemptive treatment.
- CMV-specific cell-mediated immunity (CMV-CMI) in transplant recipients can be assessed before transplantation to determine their baseline immunity.
- Immune monitoring by measuring CMV-CMI might be useful for stratifying the risk of CMV infection and managing the posttransplant strategy.
- Genotype resistance testing should be performed when drug-resistant CMV infection is suspected.

plantation [9].

CMV may indirectly affect transplant recipients, in part due to its ability to regulate the immune system [2]. CMV is associated with increased rates of bacteremia, invasive fungal infection, and Epstein-Barr virus-mediated posttransplant lymphoproliferative disorders [10-13]. CMV infection has been reported to be associated with acute rejection and chronic allograft nephropathy in renal recipients [14-17], bronchiolitis obliterans in lung recipients [18], and cardiac allograft vasculopathy [19,20]. Many studies have presented an association between CMV infection and decreased patient survival [16,21].

The definition of CMV infection is the presence of CMV replication in tissues, blood, or other body fluids, irrespective of symptoms. CMV replication can be detected by an antigen test, a nucleic acid amplification test, and virus culture. Therefore, CMV replication can be termed CMV antigenemia (antigen detection), CMV DNAemia or RNAemia (nucleic acid detection), and CMV viremia (culture) depending on the laboratory method used [22]. CMV disease refers to viral infection with clinical symptoms and signs. CMV disease includes CMV syndrome (fever, malaise, atypical lymphocytosis, cytopenia, and elevated liver enzymes) and end-organ CMV disease (gastrointestinal disease, encephalitis, retinitis, hepatitis, nephritis, pneumonitis, myocarditis, and pancreatitis). Asymptomatic CMV infection refers to CMV replication without clinical symptoms or signs [2].

The incidence of CMV infection or disease in SOT has been reported to be 12%–20% and 6%–30%, respectively [23]. The adjusted incidence rate of CMV disease in Korea,

excluding CMV syndrome, was recently reported to be 33.1 (95% confidence interval [CI], 28.8–38.0) per 1,000 person-years in SOT and 5.1 (95% CI, 4.6–6.1) per 1,000 person-years in hematopoietic stem cell transplant (HSCT) recipients. In the SOT cohort, heart transplant recipients had the highest rate of CMV disease, at 104.2 (95% CI, 66.4–163.7) per 1,000 person-years, liver transplant recipients showed the lowest rate, at 11.1 (95% CI, 7.7–16.3) per 1,000 person-years, and renal transplant recipients had an intermediate rate, at 44.3 (95% CI, 37.7–52.1) per 1,000 person-years [24].

The development and implementation of accurate and reliable diagnostic assays for CMV infection are essential for the improvement of early posttransplant CMV prevention and disease management [25]. In this review, we describe the laboratory tests for the diagnosis of CMV infection that are currently widely used in clinical laboratories. Tests for direct virus detection and measurement of host immune response are used in SOT recipients (Table 1). Laboratory tests that directly detect CMV are recommended for diagnosis, surveillance, and monitoring, whereas immune status analysis is utilized for CMV risk assessment and risk factor stratification [26].

## ASSAYS OF VIRUS DETECTION

Laboratory methods for virus detection include molecular assays, antigenemia, culture, and histopathology.

### Molecular Assay

The commercial quantitative nucleic acid amplification test (QNAT) is the preferred assay for diagnosing CMV infection, guiding preemptive treatment, and monitoring the response to therapy [27]. QNAT is very sensitive, provides a rapid turnaround, and is preferred over antigenemia [26]. Quantitative detection is preferred over qualitative analysis, as CMV QNAT can distinguish CMV replication with a high viral load from latent viral infections with low levels of CMV DNAemia [28,29]. Various commercial assays have been developed to detect and quantify CMV DNA [26]. Commercially available Food and Drug Administration (FDA)-cleared or approved platforms in the USA include the Cobas AmpliPrep/Cobas TaqMan CMV test (Roche, Basel, Switzerland), the Artus CMV RGQ MDX kit (Qiagen, Hilden, Germany), and the Abbott RealTime CMV assay (Abbott, Chicago, IL, USA) [27,30].

There are no commercially available assays for the detection of CMV RNA, although CMV RNA is a specific indicator of CMV replication. CMV DNA may not accurately reflect CMV replication, because highly sensitive QNATs can simply amplify latent viral DNA [2]. Whole blood and plasma are common specimens, but bronchoalveolar lavage (BAL) fluid and cerebrospinal fluid (CSF) are also available [31,32]. CMV DNA in CSF suggests the possibility of central nervous system disease [22]. Another challenging area for the diagnosis of CMV in HSCT and SOT recipients is CMV pneumonia. CMV QNAT in BAL fluid can be used as a less invasive tool for diagnosing CMV pneumonia, especially if performing a transbronchial biopsy would be risky. Although the reported viral load values in BAL fluid vary widely across studies, higher viral loads in BAL fluid compared to asymptomatic shedding correlated with biopsy-proven CMV pneumonia [22,31]. Further studies are warranted to develop a standardized method for collecting BAL fluid, analyzing and reporting the results, and identifying an optimal cut-off applicable in various clinical contexts. If successful, this approach could replace invasive biopsy and viral cultures for the diagnosis of CMV pneumonia in clinical settings [32]. The use of QNAT for intestinal biopsy is an evolving field. CMV QNAT on tissue biopsy specimens alone is insufficient for diagnosing CMV gastrointestinal disease. However, the guideline developed by the CMV Drug Development Forum for clinical trial standardization considers that QNAT of a biopsy specimen in a compatible clinical setting is indicative of possible gastrointestinal disease [22].

Viral load trends over time are directly related to the possibility of severe CMV disease. Higher or rapidly increasing viral loads are associated with the high risk of serious CMV disease [26]. Conversely, reduction of the viral load during antiviral therapy correlates with the clinical resolution of disease [28]. A sustained increase or minimal decrease in the viral load suggests refractory or drug-resistant CMV [33]. The guideline recommends treating patients until a negative threshold is reached, as persistent viremia at the end of antiviral therapy is a risk factor for relapse [2].

A limitation of the CMV QNAT is the absence of applicable viral load thresholds for various clinical indications. The World Health Organization (WHO) international standard for CMV nucleic acid amplification techniques was developed in October 2010 [34]. Implementation of the WHO international standard for calibration has significantly improved the degree of agreement of viral load values

between different assays and allows results to be reported in units of IU/mL [27,35]. A multicenter study for intra-assay harmonization of the CMV QNAT on plasma samples has been conducted, but clinically relevant differences in viral load values have been reported between the various assays [36]. Viral load variability among assays calibrated by the WHO international standard occurred due to differences in the assay platform, gene targets, and amplicon size [36,37]. Therefore, the same assay platform should be used for serial CMV surveillance and monitoring using the same type of sample [27,38,39]. It is recommended that transplant centers work with their laboratories to define and validate center-specific viral load thresholds for each clinical application [2,40].

As another method, droplet digital polymerase chain reaction shows potential to reduce the variability of CMV DNA load measurements, but it is not yet widely used [41-43]. Novel strategies are being tried to improve adherence by allowing patients to collect and submit their blood samples for monitoring without visiting hospitals or standard phlebotomy [44]. Dried blood spots, a previously studied method for the diagnosis of congenital CMV, can be used to assess the CMV viral load using finger-stick blood samples [44]. Dried blood spot quantification was validated in 35 SOT recipients [45] and is currently being evaluated in a multicenter randomized controlled trial using mobile device-assisted CMV monitoring in HSCT recipients at high risk of late CMV disease (ClinicalTrials.gov identifier NCT03910478) [44].

### **CMV Antigenemia**

CMV pp65 antigenemia is an indirect immunofluorescence-based assay that identifies the pp65 antigen of CMV in peripheral blood leukocytes [46]. This assay uses monoclonal antibodies to detect the pp65 antigen, an immediate early antigen of CMV [26]. This assay has limitations, including a lack of standardization with significant inter-laboratory variability, lack of automation, the need for sufficient leukocytes in the sample (limited to leukopenia), a labor-intensive nature, and subjective interpretation [26]. The use of antigenic assays has declined significantly, and antigenic assays are now being replaced by molecular assays in most transplant laboratories [28]. CMV pp65 antigenemia was comparable to QNAT in previous studies [2,46].

**Table 1.** Characteristics of cytomegalovirus assays in solid organ transplantation

Assay	Technique	Advantage	Limitation
For virus detection			
CMV QNAT	<ul style="list-style-type: none"> <li>• Detects and quantifies CMV DNA</li> <li>• Reporting unit: IU/mL</li> </ul>	<ul style="list-style-type: none"> <li>• Various commercial assays are available</li> <li>• Rapid and sensitive tool for diagnosis of CMV infection</li> <li>• Surveillance for preemptive treatment</li> <li>• Monitoring the response of antiviral therapy</li> <li>• Implementation of the WHO internal standard for calibration</li> </ul>	<ul style="list-style-type: none"> <li>• Lack of universal viral load threshold</li> <li>• Lack of standardization despite WHO internal standard material due to various aspects of assay (limit of detection, quantification range, sample type, amplicon size, gene target)</li> </ul>
Antigenemia	<ul style="list-style-type: none"> <li>• Immunofluorescence-based assay</li> <li>• Detect CMV pp65 antigen expressed in leukocytes using monoclonal antibody</li> <li>• Reporting unit: number of pp65 positive cells per number of leukocytes</li> </ul>	<ul style="list-style-type: none"> <li>• Monitoring CMV infection</li> <li>• Monitoring the response of antiviral therapy</li> </ul>	<ul style="list-style-type: none"> <li>• Lack of assay standardization</li> <li>• Need for enough leukocytes in sample (limited in neutropenia)</li> <li>• Lack of automation</li> <li>• Interpretation is subjective</li> <li>• Labor-intensive</li> </ul>
For CMV-specific cell-mediated immunity			
Serology	<ul style="list-style-type: none"> <li>• Usually detects CMV IgG antibodies</li> </ul>	<ul style="list-style-type: none"> <li>• The risk of CMV infection is determined depending on the CMV serology in donor and transplant candidate</li> </ul>	<ul style="list-style-type: none"> <li>• IgM is not recommended due to false positivity</li> <li>• The use of serology is limited for diagnosis of CMV infection after transplantation</li> </ul>
QuantiFERON-CMV	<ul style="list-style-type: none"> <li>• ELISA-based</li> <li>• Measures IFN-<math>\gamma</math></li> <li>• Collecting whole blood into tubes containing the CMV peptide pool</li> </ul>	<ul style="list-style-type: none"> <li>• Commercial assay</li> <li>• Standardized high-throughput assay</li> <li>• Can be performed routinely in laboratories</li> </ul>	<ul style="list-style-type: none"> <li>• HLA class I restricted</li> <li>• Only measures CD8+ T cells (not CD4+ T cells)</li> <li>• Indeterminate results in immunosuppressed patients</li> </ul>
ELISpot	<ul style="list-style-type: none"> <li>• Measures IFN-<math>\gamma</math></li> <li>• Stimulates PBMCs with CMV-overlapping peptides</li> <li>• Reporting unit: spot forming units/PBMCs</li> </ul>	<ul style="list-style-type: none"> <li>• Commercial assays are available: T-SPOT.CMV, T-Track CMV</li> <li>• Highly sensitive</li> <li>• Not limited by HLA</li> <li>• Measures both CD4+ T cells and CD8+ T cells</li> </ul>	<ul style="list-style-type: none"> <li>• Requires PBMC isolation procedure</li> <li>• Lack of proper cut-offs for positivity</li> <li>• Requires ELISpot reader</li> <li>• Unable to differentiate between CD4+ T cells and CD8+ T cell response</li> <li>• Lack of standardization as many laboratories use in-house methods</li> </ul>
Intracellular staining and flow cytometry	<ul style="list-style-type: none"> <li>• Detects intra-cytoplasmic cytokines produced by stimulation of whole blood or PBMCs with CMV peptides using a fluorochrome antibody</li> </ul>	<ul style="list-style-type: none"> <li>• Simultaneous detection of multiple cytokines and cell surface markers</li> <li>• Can differentiate T-cell phenotypes</li> <li>• Can differentiate between CD4+ T cells and CD8+ T cell response</li> </ul>	<ul style="list-style-type: none"> <li>• Requires flow cytometer</li> <li>• Lack of standardization</li> <li>• Expensive</li> <li>• Labor-intensive</li> <li>• Limited to research use only</li> </ul>

CMV, cytomegalovirus; QNAT, quantitative nucleic acid amplification test; WHO, World Health Organization; IgG, immunoglobulin G; IgM, immunoglobulin M; ELISA, enzyme-linked immunosorbent assay; IFN- $\gamma$ , interferon-gamma; HLA, human leukocyte antigen; ELISpot, enzyme-linked immunosorbent spot; PBMC, peripheral blood mononuclear cell.

## ASSAYS FOR IMMUNE RESPONSES AFTER CMV INFECTION

Assays for detecting the immune response to CMV in SOT include serology and CMV-specific cell-mediated immune assays [26].

### CMV Serology

Pretransplant CMV IgG measurement in the donor and recipient is the most commonly recommended test for predicting and stratifying the risk of CMV infection [47]. Based on CMV IgG results, recipients are classified as high-risk (seropositive donors and seronegative recipients), intermediate-risk (seropositive recipients), or low-risk (seronegative donors and recipients) [26]. Although CMV-seropositive recipients are considered to have a lower risk of CMV infection than CMV-seronegative recipients, the risk of CMV infection after transplantation remains. A recent study found that in seropositive recipients, patients with low pretransplant CMV IgG titers (<20 AU/mL) were more likely to have a CMV infection (hazard ratio, 2.98) after kidney transplantation (KT) than patients with CMV IgG titers greater than 20 AU/mL [48].

### Assay for CMV-CMI

The CMV-CMI test was developed to complement conventional CMV viral load assays and provide an opportunity to identify immunocompetent patients capable of controlling viral replication via host immune mechanisms without antiviral therapy [44]. The posttransplant risk of CMV infection can be predicted by pretransplant CMV-CMI assessment. CMV-CMI can be measured during or at the end of CMV prophylaxis to predict the risk of CMV infection, or at the completion of treatment to predict the risk of CMV recurrence or to determine whether secondary prophylaxis is needed [47].

CMV-CMI studies, mostly performed in KT recipients, showed that measured immunity levels correlated with virologic outcomes. Several studies have reported that increased CMV-CMI levels measured by CMV-specific interferon-gamma (IFN- $\gamma$ ) release assays are associated with reduced incidence of CMV infection or disease, lower initial and peak viral load, reduced CMV recurrence, and freedom from CMV events [49-64]. It has been shown that a lack of adequate CMV-specific CD4+ and/or CD8+ T cell immunity is correlated with a higher likelihood of CMV disease, recurrence, and treatment failure [2]. However, to date, well-designed large-scale intervention studies

demonstrating the clinical utility of CMV-CMI are still lacking.

Commercially available CMV-CMI assays have the advantages of relative ease of use, standardized approaches, and suitability for comparing results between studies of different populations [27]. Commercially available assays include the QuantiFERON-CMV (CMV-QF) enzyme-linked immunosorbent assay (ELISA; Qiagen), T-SPOT.CMV (Oxford Immunotec, Abingdon, UK) and T-Track CMV (Lophius Biosciences, Regensburg, Germany), which are enzyme-linked immunosorbent spot (ELISpot) assays [44]. CMV-QF, T-SPOT.CMV and T-Track CMV are not FDA-approved, but are CE (Conformité Européenne) marked in Europe [65]. Recently, the novel CMV T Cell Immunity Panel (Viracor Eurofins Inc. Laboratories, Lee's Summit, MO, USA), which measures CMV-specific cellular immunity by intracellular cytokine staining (ICS) and flow cytometry, has become available in the USA [66,67] (Table 2).

### QuantiFERON-CMV

CMV-QF measures the IFN- $\gamma$  response of CD8+ T cells to various human leukocyte antigen (HLA) class I restricted synthetic CMV T cell epitopes. The peptides are designed to target A1, A2, A3, A11, A23, A24, A26, B7, B8, B27, B35, B40, B41, B44, B51, B52, B57, B58, B60, and Cw6 (A30, B13) HLA class I haplotypes covering >98% of the human population (<https://www.quantiferon.com/wp-content/uploads/2018/10/L1075110-R05-QF-CMV-ELISA-IFU-CE.pdf>). These peptide epitopes include pp65, pp50, IE-1, IE-2, and the glycoprotein B antigens. Whole blood is incubated with CMV peptides for 18–24 hours, and then the supernatant is harvested and the level of IFN- $\gamma$  is measured based on ELISA. The cut-off recommended by the manufacturer is 0.2 IU/mL [68]. One major limitation of CMV-QF is that the use of HLA class I-restricted CMV epitopes may not reflect the ability of some individuals to recognize and respond to the epitope [69]. About 60% of pretransplant CMV-seropositive recipients were negative for CMV-QF [70]. A recent study performed CMV-CMI in healthy individuals and found 18.3% (13/71) revealed humoral/cellular discordance, showing CMV-seropositivity with CMV-QF negativity [71]. Individuals with inconsistent results had lower levels of CD4+ and CD8+ T cell proliferation in response to CMV lysate stimulation and expressed a lower level of anti-CMV IgG. This suggests that immune response to CMV was highly heterogeneous in healthy subjects [71]. In addition, CMV-QF only measures the CMV-specific CD8+ T cell response. However, CMV-specific CD4+ T cells are

**Table 2.** Characteristics of commercially available CMV-specific cell-mediated immunity assays

	QuantiFERON-CMV	T-SPOT.CMV	T-Track CMV	CMV T cell immunity panel
Assay principle	ELISA	ELISpot	ELISpot	Intracellular staining and flow cytometry
Product company	Qiagen	Oxford Immunotec	Lophius Bioscience	Viracor Eurofins
Sample	3 mL of whole blood	Purified PBMCs from 12 mL of whole blood	Purified PBMCs from 15 mL of whole blood	10 mL of whole blood
PBMCs required	No	Yes	Yes	Yes
CMV antigen	Various HLA class I restricted 21 CMV peptides	pp65, IE-1	T-activated CMV-specific pp65 and IE-1	Whole viral lysate, pp65, IE-1
Measurement	IFN- $\gamma$	IFN- $\gamma$ -specific spot-forming cells	IFN- $\gamma$ -specific spot-forming cells	% CMV-specific CD4+ and CD8+ T-cells
Cut off for positivity	CMV antigen minus Nil control $\geq 0.2$	NA	Either pp65 or IE-1 $\geq 10$ spots	CMV-specific CD4+ or CD8+ responses $> 0.2\%$
Measuring range (linearity)	Up to 10 IU/mL	NA	10–1,000 spots	NA
Clinical sensitivity	80.5% (insert)	93.3% (insert)	89.6% (insert)	79%–82% [66]
Time to test	16–24 hours	30–40 hours	30–40 hours	3–4 business days from receipt of specimen in USA
Quality control	Nil control $\leq 8.0$ Mitogen control	Nil control $< 10$ spots Positive control (mitogen solution containing PHA) $> 20$ spots	Negative control $< 10$ spots Positive control (SEB) $> 400$ spots	Negative control Positive control (SEB)
Comment	CE-marked Not FDA-approved	CE-marked Not FDA-approved	CE-marked Not FDA-approved	Not FDA-approved

CMV, cytomegalovirus; ELISA, enzyme-linked immunosorbent assay; ELISpot, enzyme-linked immunosorbent spot; PBMC, peripheral blood mononuclear cell; HLA, human leukocyte antigen; IFN- $\gamma$ , interferon-gamma; NA, not available; PHA, phytohemagglutinin; SEB, *Staphylococcus aureus* enterotoxin type B; CE, Conformité Européenne; FDA, Food and Drug Administration.

important for generating a pool of memory CD8+ T cells capable of controlling CMV infection [66], and evidence for antiviral properties of CD4+ T cells against CMV has also been reported [72]. Caution is needed in interpreting indeterminate results of CMV-QF, which may occur in up to 38% of HSCT recipients [73]. Indeterminate results may result from improper processing of the samples, but many of them are caused by inadequate T cell responses to the mitogen, which reflects low or dysfunctional T cells [73,74].

### ELISpot Assays

The ELISpot assay measures IFN- $\gamma$  production by both CD4+ and CD8+ T cells in response to stimulation by CMV antigens. Production levels are quantified by counting the number of spot forming units per given number of target cells, such as peripheral blood mononuclear cells (PBMCs). Currently commercially available assays are the T-Track CMV and T-SPOT.CMV. For analysis, PBMCs are isolated

from whole blood, placed in wells, and stimulated with CMV pp65 and IE-1 antigens (urea-formulated proteins in T-Track CMV, peptides in T-SPOT.CMV). The T-Track CMV assay uses recombinant urea-formulated pp65 and IE-1 antigens that activate a wide range of effector cells such as CD4+, CD8+ T cells, natural killer (NK), and NK T cells [75]. In both assays, after 17–21 hours of incubation, secreted IFN- $\gamma$  binds to IFN- $\gamma$  specific capture antibodies in the well, and the binding of enzyme-linked secondary antibody generates insoluble spots for detecting antibody-bound IFN- $\gamma$  [47]. CMV-QF is easier to perform than ELISpot because it does not require additional procedures and laboratory instruments for PBMC purification [65]. However, ELISpot assays are more sensitive than ELISA-based assays and produce quantitative spot results [76]. A limitation of the ELISpot assay is the inability to differentiate between CD4+ and CD8+ T cells [27]. Moreover, since many laboratories use in-house methods, there are no adequate cut-

offs and a lack of standardization [77].

### ICS and Flow Cytometry

ICS using flow cytometry may be a superior method due to its ability to simultaneously measure multiple cell surface molecules and cytokines in real time and provide quantitative characteristics of CMV-specific CD4+ and CD8+ T cells [27]. Whole blood or PBMCs are stimulated with CMV peptides, and then the cells produce cytokines such as IFN- $\gamma$ , tumor necrosis factor-alpha, and interleukin (IL)-2. Intracellular cytokines of interest are stained with fluorescein-coated antibodies [77]. ICS is the only assay capable of analyzing CD4+ and CD8+ T cell responses separately [66]. However, the ICS assay has several disadvantages, including a lack of standardization, resource intensiveness, a high cost to perform, and the need for expert interpretation [77]. Additionally, there are only a few studies supporting ICS assays, and to date, they have not been widely used in clinical laboratories [78-81]. In a recent multicenter cohort study of 124 CMV-seropositive KT recipients not receiving anti-thymocyte globulin (ATG), ICS was measured before and 15 days after transplantation [81]. Low levels of CMV-CMI (defined as <2.0 CD8+ T cells/mL or <1.0 CD4+ T cells/mL) measured on day 15 were associated with a higher likelihood of CMV events (asymptomatic infection or disease).

The CMV inSIGHT T cell immunity panel from Viracor Eurofins Laboratories was recently commercialized in the USA (<https://www.eurofins-viracor.com/clinical/test-menu/30360-cmv-insight-t-cell-immunity/>). This assay measures CMV-specific CD4+ and CD8+ T cell responses based on the percentage of cells expressing surface CD69 and IFN- $\gamma$  after stimulation with CMV epitopes (including whole viral lysate, pp65, and IE-1) [82]. Although further studies are needed, this assay has shown good diagnostic performance in predicting CMV events [66,67].

### Factors Influencing CMV-CMI

It is necessary to review the factors that may affect CMV-CMI in SOT. First, it has been demonstrated that the kinetics of CMV-specific CD4+ T cell responses are different from those of CD8+ T cells [83]. The level of CMV-specific CD4+ T cells significantly decreased at 2 months post-KT compared to the pretransplantation level and gradually increased to the pretransplant level by 12 months. In contrast, it was found that the CMV-specific CD8+ T cell response decreased rapidly within the first 2 weeks post-KT, but returned to the pretransplant levels by 2 months post-

KT [83]. Second, posttransplant CMV-specific T cell levels may be affected by induction immunosuppression. Abate et al. [84] reported that ATG treatment had no impact on CMV-specific T cell responses. However, recent prospective studies revealed that the ATG group showed significantly lower cellular immunity than the group of recipients treated by anti-IL 2 receptor antibodies (basiliximab) [49,50]. Calcineurin inhibitors have also been reported to have direct inhibitory effects on CMV-specific T cell reactivity [80,85]. Third, the influence of antiviral prophylaxis on CMV-CMI levels has yet to be conclusively determined [47]. In addition to promising data showing the clinical utility of CMV-CMI assessment for risk stratification, a proportion of SOT recipients (4%–10%) developed late-onset CMV disease [86,87]. This suggests that a higher cut-off for positive CMV-CMI, which can reliably predict protection against CMV disease, may be necessary, or that other aspects of the immune response, such as neutralizing antibodies, should also be considered [88]. A previous study showed that quantifying epithelial cell neutralizing antibodies was useful for identifying liver transplant recipients with lower risk for CMV disease [86].

Although many studies over the past 20 years have consistently evaluated the clinical application of CMV-CMI assays, they have not been routinely incorporated into clinical practice. The reasons are the absence of well-defined thresholds for positivity and negativity, the variability in CMV antigens in a protocol (whole cell lysate vs. peptide pools, pp65 vs. IE-1), and the heterogeneity of study populations (transplanted organ, serostatus of the donor/recipient, use of antiviral prophylaxis vs. preemptive therapy). In addition, most of the data were from observational studies, and there were only a few intervention studies in which treatments were made based on the CMV-CMI results [47].

### Recent Observational Studies of CMV-CMI Assays

Observational studies have confirmed that CMV-CMI assays can predict the risk of subsequent CMV replication and disease development. Strong evidence indicates that assessing CMI using standardized assays such as CMV-QF and ELISpot (T-Track CMV and T-SPOT.CMV) can stratify recipients according to their risk of CMV events [77].

A large, prospective observational study of KT recipients showed that CMV-specific immune assessment using the ELISpot assay can predict protection from CMV infection [49]. In 583 recipients consisting of 260 seronegative recipients with a seropositive donor (D+/R-) and 277 seropositive recipients (R+), the CMV ELISpot assay was per-

formed at prophylaxis discontinuation and once per month for 4 months and at 6 months. CMV events were significantly lower in ELISpot-positive patients (>40 spot-forming units per  $2.5 \times 10^5$  cells for either pp65 or IE-1) vs. negative patients (3.0% vs. 19.5%,  $P < 0.0001$ ). However, the positive predictive value was low (less than 20%), showing poor sensitivity for distinguishing patients at high risk of CMV replication. This study showed the possibility of performing real-time CMV-CMI in a central laboratory, and also confirmed that this assay is applicable in routine clinical practice.

Two recent studies evaluated the role of CMV-CMI in lung transplant patients, who constitute a high-risk group for CMV-related complications. The first Spanish study tested the T-SPOT.CMV assay in 60 CMV seropositive recipients at posttransplant 6 months when prophylaxis was withdrawn [89]. Recipients who developed late-onset CMV DNAemia showed significantly lower ELISpot responses, particularly to the IE-1 antigen, than patients who did not. In the second study, 39 lung transplant candidates underwent CMV-QF prior to transplantation [90]. CMV-QF negativity was associated with a higher likelihood of CMV reactivation after transplantation. This is consistent with previous studies, in which CMV-CMI at the time of transplantation had better predictive value than CMV serostatus for the development of CMV replication.

#### Recent Interventional Clinical Trials of CMV-CMI Assays

A few interventional clinical studies on CMV-CMI assay have been published [50,63,64,91]. The first interventional trial using CMV-CMI was a pilot nonrandomized uncontrolled trial in 27 SOT recipients with CMV replication and/or disease, in whom a CMV-QF assay was performed after 2–3 weeks of administration of antiviral therapy [64]. Fourteen patients (51.9%) had positive CMV-QF responses at the end of treatment, and the antiviral therapies were discontinued. The remaining 13 patients (48.1%) had negative results and received secondary antiviral prophylaxis for 2 months. Patients with detectable CMV-CMI had a lower rate of CMV relapse than recipients who were CMV-CMI negative and received longer-term antiviral therapy (7.1% vs. 69.2%,  $P = 0.001$ ). This was the first interventional study to demonstrate the validity and safety of real-time CMV-CMI assessments to guide changes in CMV management.

A second randomized controlled trial was conducted to determine the duration of antiviral prophylaxis according to the results of CMV-QF [63]. In this study, lung transplant recipients ( $n = 118$ ) were randomized to receive a fixed du-

ration of prophylaxis (5 months) or a duration determined by the CMV-QF assay, performed at 5, 8 and 11 months posttransplant. The incidence of CMV infection (>600 copies/mL in BAL fluid) was significantly lower in the CMV-QF guided group than the standard of care group (36.6% vs. 58.3%,  $P = 0.03$ ). Of the 80 patients who ceased prophylaxis (36 in the standard-of-care group and 44 in the CMV-QF-directed group), the incidence of severe viremia (>10,000 copies/mL) was lower in recipients with positive CMV-QF than in those with negative CMV-QF (3% vs. 50%,  $P < 0.001$ ). CMV-CMI monitoring allows an individualized approach to CMV prophylaxis and reduces late CMV infection within the lung allograft.

A third trial used pretransplant cell-mediated immune status by means of an ELISpot-CMV to determine the risk for developing posttransplant CMV replication [50]. Using pretransplant T-SPOT.CMV (cut off, 20 spots/ $3 \times 10^5$  PBMCs), patients were divided into two groups: group A (low-risk T-SPOT.CMV,  $\geq 21$  spots/ $3 \times 10^5$  PBMCs) and group B (high-risk T-SPOT.CMV,  $\leq 20$  spots/ $3 \times 10^5$  PBMCs). Each group was randomized at a 1:1 ratio and divided into a 3-month antiviral drug prophylaxis group (subgroups A1 and B1) or a preemptive treatment group (subgroups A2 and B2). Patients at high risk according to the pretransplant T-SPOT.CMV results showed a higher risk of CMV infections than those at low risk for both the prophylactic (33.3% vs. 4.1%) and preemptive approach (73.3% vs. 44.4%). However, the predicted capacity of CMV-CMI to identify recipients at high risk of CMV infection was only found in patients treated with basiliximab (not receiving T cell-depleting antibodies) for both prophylactic and preemptive therapy. Furthermore, 15-day posttransplant T-SPOT.CMV was a better predictor of CMV infection than CMV-CMI measured before transplantation in basiliximab-treated patients. This study showed that CMV-CMI monitoring may guide decisions regarding the type of CMV preventive strategy in KT.

A randomized clinical trial by Singh et al. [91] compared preemptive therapy versus prophylaxis in 205 CMV seronegative liver transplant recipients from seropositive donors. Patients received either valganciclovir (900 mg) prophylaxis daily for 100 days, or valganciclovir (900 mg) twice a day if viremia was detected during weekly monitoring with CMV QNAT for 100 days. The incidence of symptomatic CMV disease was significantly lower in the preemptive group than in the prophylaxis group (9.0% vs. 19.0%,  $P = 0.04$ ). The incidence of opportunistic infections, rejection, graft loss, and mortality showed no differences

between the two groups. In this study, CMV pp65 ELISpot responses were stronger after preemptive therapy than prophylaxis for both CD4+ and CD8+ T cells. Among the preemptive recipients, CD8+ T cell responses were significantly higher in patients with preceding viremia than in those without viremia.

Two additional randomized interventional trials have been completed, and we are looking forward to the upcoming results [65]. In one clinical trial (Clinicaltrials.gov identifier NCT03123627), CMV seropositive KT recipients who underwent ATG induction therapy were randomized to receive a fixed period (3 months) of valganciclovir prophylaxis or a period determined by CMV-QF results. Therapy was stopped in case of a positive CMV-QF. A second trial (Clinicaltrials.gov identifier NCT02538172) used an ELISpot (T-Track CMV) and included D+/R- and R+ kidney and liver transplant recipients receiving ATG. The control group received valganciclovir prophylaxis up to posttransplant 90 days, and the prophylaxis period of the intervention group was determined by the T-Track CMV assay. We expect these studies to support a promising role for the CMV-CMI assay in modulating the duration of antiviral prophylaxis.

## ASSAYS FOR CMV DRUG MUTATION

Genotyping should be performed to detect specific drug mutations when drug-resistant CMV infection is suspected [2]. A 1-log reduction in the CMV viral load is expected when treated with an appropriate dose after at least 2 weeks. Refractory CMV infection is defined as an increase of  $>1 \log_{10}$  in CMV DNA levels between baseline and after at least 2 weeks of an appropriate dose of antiviral treatment [33]. Probable refractory infection is defined as CMV DNA levels that persist (at the same level or increasing by less than  $1 \log_{10}$ ) after at least 2 weeks of antiviral therapy. However, persistent CMV DNA titers less than 1,000 IU/mL, particularly detected but non-quantifiable levels ( $<137$  IU/mL), should not be considered as refractory [33]. Genotyping is possible for viral sequences that are amplified from blood (plasma, whole blood, or leukocytes), body fluids (CSF, BAL fluid, urine, or vitreous humor), or tissue [2]. The sample should have sufficient CMV DNA levels. The results are more reliable when the CMV copy number is greater than 1,000 IU/mL [27]. The accuracy of variant sub-population detection was lower at 1,000 copies/mL than at 10,000 copies/mL [92]. CMV genes associated with com-

mercially available or novel antiviral agents are *UL97* and *UL54* for ganciclovir; *UL97* and *UL27* for maribavir; *UL54* for cidofovir and foscarnet; and *UL51*, *UL56*, and *UL89* for letermovir [33].

## CONCLUSION

Advances in diagnostics have been essential for improving our understanding of CMV immunity and its role in disease. The use of sensitive CMV QNAT has become the standard of care as a means to monitor and treat CMV disease in transplant patients. CMV immune monitoring to better identify individuals at high risk for CMV-related complications is an area of continuing high clinical need and interest. With the development of standardized CMV-CMI assays such as CMV-QF and ELISpot, CMV immune monitoring is being integrated into routine clinical care, moving one step closer to personalized medicine.

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### Conflict of Interest

Eun-Jee Oh is an editorial board member of the journal but was not involved in the peer reviewer selection, evaluation, or decision process of this article. No other potential conflict of interest relevant to this article was reported.

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