



## Review

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# Cytomegalovirus and immunotherapy: opportunistic pathogen, novel target for cancer and a promising vaccine vector

Cytomegalovirus (CMV) is a  $\beta$ -herpesvirus that infects most people in the world and is almost always asymptomatic in the healthy host. However, CMV persists for life, requiring continuous immune surveillance to prevent disease and thus, CMV is a frequent complication in immune compromised patients. Many groups have been exploring the potential for adoptive T-cell therapies to control CMV reactivation as well as the progression of solid tumors harboring CMV. In addition, CMV itself is being explored as a vaccine vector for eliciting potent T-cell responses. This review will discuss key features of the basic biology of CMV-specific T cells as well as highlighting unanswered questions and ongoing work in the development of T-cell-based immunotherapies to target CMV.

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Cytomegalovirus (CMV) is a member of the  $\beta$ -subfamily of herpesviruses, a family of viruses that are thought to have been co-evolving with their hosts for ~180 million years [1]. CMV establishes an asymptomatic, latent/persistent infection in most people in the world and persists for the life of the host [2]. Importantly, reactivation from latency is thought to occur regularly throughout the body, requiring constant immune surveillance to keep the host disease free. Indeed, patients who are immunosuppressed are at risk for CMV reactivation, which can lead to increased morbidity and mortality (comprehensively reviewed in [3]).

Even in immune-competent individuals, constant CMV reactivations serve as potent stimulators for CMV-specific T cells. Intriguingly, recombinant CMV viruses are now being explored as potential vaccine vectors to generate large numbers of T cells against both infectious diseases and cancer. Importantly, recent work from the Picker lab

has demonstrated that CMV-based vaccines were profoundly protective in a nonhuman primate model of HIV infection [4–7]. Thus, a deeper understanding of how CMV-specific immunity is stimulated and maintained will help to improve CMV immune restoration following immune suppression and help to develop new vaccines using attenuated CMV viral vectors.

Our current understanding of the immunology and virology of CMV infections is derived from both human studies and animal models. Unfortunately, *in vivo* studies using human CMV (HCMV) is difficult because HCMV is often asymptomatic during the acute phase of infection and produces undetectable levels of transcripts during the latent phase. Additionally, CMV species selectivity is so strict that HCMV will not replicate in nonhuman cells [8]. Thus, studies using HCMV are usually performed *in vitro* and animal studies require species-specific CMV orthologs.

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Remarkably, the overall host–pathogen balance has been highly conserved despite divergence between these species-specific CMV viruses. In particular, the natural mouse pathogen murine CMV (MCMV) has been a well-described tool for investigating CMV-specific pathogenesis and immunity [9,10]. It is important to note that HCMV and MCMV differ in multiple ways, including the expression of many unique genes and aspects of viral pathogenesis that are dependent on the host species (e.g., *in utero* transmission [11]). However, the overall viral life cycles are overlapping and there are several examples of unique viral genes in each virus that have overlapping functions. Importantly for studies of immune control, both HCMV and MCMV use similar mechanisms to evade or limit immune control, both establish latency in the same cell types and both viruses require constant immune surveillance to prevent viral replication and disease [9,12–16]. Although additional studies are needed to further understand and appreciate the similarities and differences between MCMV and HCMV, the MCMV model has provided directly translatable insight into HCMV, particularly in the arena of immune control.

Investigations over the last 20 years with HCMV and animal models of CMV infection have revealed that immune control of CMV is a layered process. Type-I IFN, NK cells,  $\gamma\delta$ -T cells, B cells, CD4<sup>+</sup> T cells and CD8<sup>+</sup> T cells all play a recognized (if not yet fully defined) role in suppressing viral activity [3]. In terms of CMV-specific T cells, it is clear that CMV-specific CD8<sup>+</sup> T cells can, in isolation, restrict CMV replication as first shown in mice by Reddehase *et al.* [17] and subsequently in humans by Riddell *et al.* [18]. Moreover, long-term control of CMV in humans requires both CD4<sup>+</sup> and CD8<sup>+</sup> T cells [19,20]. Precisely how CMV-specific T cells remain functional and control this infection for life remains a mystery. Importantly, the biology of CMV-specific CD8<sup>+</sup> T cells is parallel between mouse and man. In both species, CMV-specific CD8<sup>+</sup> T cells persist in large numbers and slowly accumulate over time, a process that has been called ‘memory inflation’ [21,22]. Moreover, the phenotype, genetic signature and half-life of MCMV- and HCMV-specific T cells overlap, indicating the relevance of the MCMV model to study HCMV-specific CD8<sup>+</sup> T cells [23–27]. Therefore, the MCMV model is an outstanding tool to explore HCMV-specific T-cell populations.

CMV-specific T cells are at the core of multiple therapeutic approaches designed to control CMV, other pathogens and even cancer. The purpose of this review is to discuss the current state of knowledge about CMV-specific T cells, as well as their use in the settings of adoptive immunotherapy and vaccination.

## Part I: CMV reactivation & control

The conditioning regimens necessary for successful hematopoietic stem cell transplantation (HSCT) cause severe immunosuppression in the recipient and, typically, a loss of CMV-specific T cells. Rapid recovery of antiviral CD4<sup>+</sup> and CD8<sup>+</sup> T cells is critical for viral control and reducing morbidity and mortality [15,19,28]. Studies have shown that it is beneficial for CMV-seropositive patients to receive stem cells from a CMV-seropositive donor, likely because of the presence of donor CMV-specific immune cells that are transplanted with the graft [29,30]. Additional risk factors for CMV disease include: the use of an MHC mis-matched graft, T-cell depletion from the graft and poor T-cell reconstitution after transplant, treatment with high-dose corticosteroids or other immune suppressive drugs (comprehensively reviewed in [31]). Unfortunately, current clinical therapies for CMV replication leave much to be desired [31] and thus, restoring CMV-specific immunity remains the primary goal. A better understanding of CMV-specific immunity, particularly CMV-specific T-cell populations, could enable the development of therapies that promote the recovery of CMV-specific T-cell populations in immune suppressed patients, or improve the development of adoptive T-cell therapies designed to control CMV in these people.

## CMV latency & reactivation

CMV is primarily spread via mucosal contact with infected secretions and acutely infects various cell types [2]. After this acute infection and systemic spread, CMV establishes latency throughout the body. Studies using MCMV have shown that the viral genome copy number in various organs is vast compared with the number of detectable transcripts [32], suggesting that the majority of MCMV genomes are transcriptionally silent. Likewise, cells harboring viral DNA can be recovered from humans without evidence of productive viral replication [33].

In order to reactivate from the latent state and produce infectious particles, like all herpesviruses, CMV gene transcription progresses through an ordered cascade beginning with expression of the intermediate-early (IE) genes, which serve as transactivators for further viral gene expression [34]. Following the expression of the IE genes, reactivation progresses to the transcription of the early (E) and late (L) genes that are important for host manipulation, DNA replication and viral packaging (reviewed in [2]). Studies of CMV latency and reactivation in the MCMV model have revealed that IE gene expression occurs continuously throughout latency in a focal and stochastic fashion [35–37]. Thus, it appears there is always a basal level of CMV reactivation. Expression of IE genes is controlled by the

major immediate early promoter, which contains consensus binding sites for pro-inflammatory transcription factors (e.g., NF- $\kappa$ B and AP-1) [38–40]. Not surprisingly, inflammatory signals can promote reactivation of HCMV and MCMV [38,41–43]. In this way, CMV is sensitive to its environment and viral activity is directly modulated by the local inflammatory milieu. For this reason, inflammation from a transplant routinely promotes viral reactivation.

### T-cell control of CMV latency & reactivation

A substantial amount of data indicates that T cells play a critical role in the control of latent and reactivating CMV. Elegant studies from the Reddehase lab have shown that MCMV-specific CD8<sup>+</sup> T cells actively suppress viral reactivation in the lungs of latently infected mice by limiting the cascade of viral gene expression [44]. An additional large body of work from this group has shown that adoptively transferred CD8<sup>+</sup> T cells specific for various MCMV proteins could control viral replication (reviewed in [45]). Likewise, infusions of HCMV-specific CD8<sup>+</sup> T cells can restore CMV-specific immunity in people, in both prophylactic and therapeutic settings [20,46].

The Reddehase lab has further shown that adoptive transfer of CD4<sup>+</sup> T cells was ineffective at limiting viral replication in immune compromised mice [17,47]. Interestingly however, mice lacking CD8<sup>+</sup> T cells still controlled viral replication in most organs [48]. Work in humans has suggested that antiviral CD4<sup>+</sup> T cells may be important for sustaining HCMV-specific CD8<sup>+</sup> T-cell populations in some settings [20] or for directly controlling viral replication through the production of IFN- $\gamma$  and/or expression of cytolytic machinery [49–51]. Importantly, mice lacking CD4<sup>+</sup> T cells have persistent viral replication selectively in the salivary gland [52] where CD4<sup>+</sup> T-cell-derived IFN- $\gamma$  is required to control viral replication [53,54]. The salivary gland is a major site of viral shedding for both MCMV and HCMV. Data from the Oxenius lab suggest that CD8<sup>+</sup> T cells cannot recognize virus-infected epithelial cells in the salivary gland as a result of viral immune evasion of class I MHC [53]. Interestingly, these same immune evasion genes have a much more subtle impact in other organs of the body [55], enabling better control of CMV by CD8<sup>+</sup> T cells. These data show that CD4<sup>+</sup> and CD8<sup>+</sup> T cells collaborate to control CMV systemically.

The ongoing immune surveillance of CMV sustains the largest antiviral T-cell populations in the circulation of healthy adults [56]. However, viral transcripts are nearly undetectable. The relationship between the undetectable virus and the overwhelming T-cell response to viral persistence remains one of the major mysteries about CMV immune surveillance. The data

suggest that both peak viral titers achieved during acute infection, as well as reactivation and/or reinfection contribute to the overall size of the CMV-specific T-cell pool [57–59]. Moreover, viral spread from the initially infected cells is not required to sustain these large T-cell pools [60]. Together, these data imply that the number of latently infected cells directly influences the size of the CMV-specific T-cell compartment. It is not at all clear, however, that the number of T cells sustained by CMV persistence is necessary to prevent CMV disease. Nevertheless, a minimum threshold must be reached after HSCT to prevent disease [61]. It might be argued that the systemic nature of the infection and the sheer number of potentially infected cells demand enormous numbers of T cells to guarantee that some are near to every reactivation event.

### Part II: adoptive immunotherapy to control CMV

In humans, there is a clear relationship between generating CD8<sup>+</sup> T-cell responses to CMV and limiting CMV infection after HSCT (reviewed in [31]). In a seminal study, Reusser *et al.* reported that patients that generated a CMV-specific CD8<sup>+</sup> cytotoxic T-cell response within the first 3 months after HSCT did not develop CMV pneumonia [15]. Numerous additional studies have shown that adoptive therapy can restore CMV immunity, reduce the risk for CMV infection, reduce the need for antiviral therapy and treat infections that are resistant to antivirals (recently reviewed in [62]). Thus, transferring CMV-specific T cells to HSCT recipients has become an attractive therapeutic option to quickly restore CMV immunity and prevent CMV-related mortality.

In the development of this therapy, it is important to consider the fact that the repeated interactions between CMV-specific T cells and their antigen profoundly effect the T-cell compartment. Indeed, the process of ‘memory inflation,’ in which a subset of viral antigens accumulate and persist in large numbers, is thought to be the result of repeated antigen encounter by CMV-specific T cells [21,22,58,63–65]. Importantly, CMV-specific T cells are markedly altered by these repeated antigen encounters. Most CMV-specific CD8<sup>+</sup> T cells that undergo memory inflation downregulate an array of molecules associated with memory T cells, and express a phenotype consistent with terminally differentiated effectors, or senescent T cells (e.g., CD62L<sup>low</sup>, CD127<sup>low</sup>, CD27<sup>low</sup>, CD28<sup>low</sup>, CD57<sup>high</sup>, KLRG-1<sup>high</sup>, NKG2A<sup>high</sup>, previously reviewed in [66]). Importantly for adoptive therapies, these terminally differentiated CD8<sup>+</sup> T cells have a short half-life of ~45–60 days in mice [24] and humans [25], and have limited proliferative capacity [27,64]. Such cells might be expected to provide

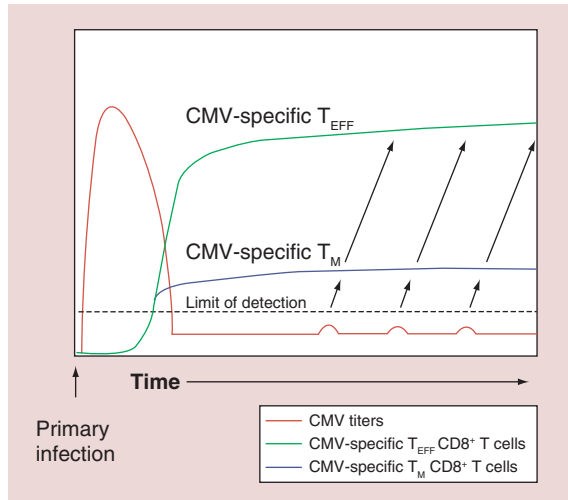
only transient protection in an adoptive recipient. Moreover, adoptive immunotherapy protocols have typically involved the expansion of CMV-specific T cells *in vitro*, followed by their infusion into immune-compromised patients without regards to differentiation status. Presumably, this antigen-driven T-cell expansion in culture further promotes their effector differentiation prior to adoptive transfer, thus potentially limiting their efficacy.

**Memory T cells retain superior expansive capacity**

Data generated by our group and the Oxenius lab has suggested that a rare population of memory T cells, with substantial proliferative and self-renewal capacity, was responsible for repeatedly producing short-lived effectors in response to viral reactivation in immune competent hosts [27,64] (Figure 1). Theoretically, such memory-like T cells could provide robust, sustained immunity after adoptive therapy. Indeed, multiple studies in humans and mice have found that small subsets of ‘inflationary’ CMV-specific T cells retain a memory-like (CD27<sup>hi</sup>, CD127<sup>hi</sup>, KLRG-1<sup>low</sup>) phenotype. Cells with the same specificities, and even individual T-cell clones, can be distributed across these terminally differentiated and memory-like phenotypes [23,27,64,67–68]. Our recent work showed that the memory-like subset of ‘inflationary’ T cells could support sustained effector populations in response to acute infection or reactivating virus, even after adoptive transfer [27]. Specifically, these memory-

like cells could: i) repeatedly produce new effector-like progeny after stimulation, ii) repeatedly produce new memory-like T cells that retained their proliferative capacity even after multiple stimulations (i.e., not undergo terminal differentiation after stimulation) and iii) sustain themselves in the complete absence of antigen and then respond to an infectious challenge or viral reactivation at a later time. All three of these properties would seem to make CMV-specific memory-like T cells an ideal subset for adoptive immunotherapy.

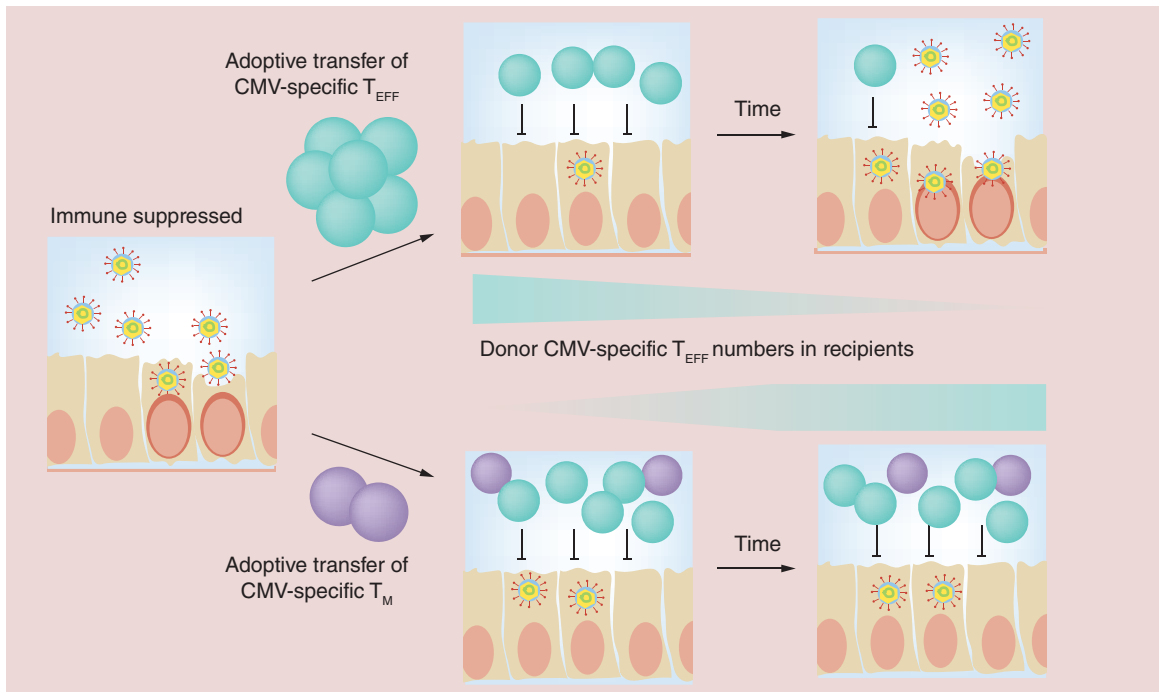
There is additional emerging evidence from humans and nonhuman primates that memory-phenotype T cells may be the ideal population for CMV-specific adoptive immunotherapy. First, the CMV-specific repertoire in bone marrow transplant recipients was dominated by transferred CMV-specific T cells that had been memory phenotype in the donor, and not those that were largely effector phenotype before the transplant [68]. These data suggested that the memory phenotype T cells were substantially more expansive than their effector counterparts. Second, CMV-specific T-cell lines derived from central memory cells displayed superior engraftment in a nonhuman primate model of adoptive immunotherapy, relative to T-cell lines derived from effector memory or effector T cells [69]. Together, these data indicate that memory-phenotype T cells maybe far more relevant for establishing long-term immunity after bone marrow transplantation (Figure 2). By extension, we might predict that a preponderance of effector or effector-memory phenotype CMV-specific T cells might identify patients at risk of CMV reactivation and disease. Careful analyses of the CMV-specific T-cell populations in the donor and recipient may help to stratify these individuals into those requiring more or less aggressive therapies. Moreover, approaches to isolate and expand CMV-specific T cells without promoting their terminal differentiation, may be extremely valuable for future adoptive immunotherapies.



**Figure 1. Model for the maintenance of overwhelming numbers of cytomegalovirus-specific T cells during persistent/latent infection in a healthy host.** After acute infection, CMV titers (red line) fall below the limit of detection. However, stochastic reactivation events stimulate proliferative memory-phenotype T cells (blue line) to divide and differentiate, producing effector T-cell progeny (green line) that are poorly proliferative and have a relatively short half-life. CMV: Cytomegalovirus; T<sub>EFF</sub>: Effector T cell; T<sub>M</sub>: Memory T cell.

**Part III: CMV & cancer: a new avenue for adoptive immunotherapy**

Work over the last several years has shown that HCMV proteins and nucleic acids can be found in samples of human glioblastoma (GBM), colon, breast and prostate cancers and rhabdomyosarcomas [70–74]. Significantly, MCMV infection was shown to induce rhabdomyosarcoma formation in mice heterozygous for the tumor suppressor p53 [74]. Although significant efforts have gone into elucidating this phenomenon, the correlation between CMV and cancer remains controversial, and results have varied by study (reviewed recently Dey *et al.* [75]). As yet, it is unclear whether the virus causes cellular transformation leading to tumors, exacerbates or accelerates tumor formation



**Figure 2. Adoptive T-cell therapies utilizing memory-phenotype T cells may more robustly sustain cytomegalovirus-specific immunity.** Our data have shown that memory-phenotype murine cytomegalovirus-specific T cells can persist in the absence of antigen, but expand robustly in response to viral reactivation, producing both effector and memory progeny. By contrast,  $T_{EFF}$  have a limited half-life and are poorly proliferative. In an adoptive immunotherapy setting, transferring  $T_{EFF}$  would be expected to provide transient protection that is lost over time as donor T cells decline (upper panels). By contrast, transferring memory-phenotype T cells would enable long-lasting protection that sustains T-cell numbers (lower panels). CMV: Cytomegalovirus;  $T_{EFF}$ : Effector T cell;  $T_M$ : Memory T cell.

or is merely a passenger making use of these rapidly dividing cells. Regardless of cause and effect, the fact that CMV can be found in these tumors has led to a recent push for targeting HCMV to treat GBM [76,77]. Clinical trials have demonstrated increased overall survival in GBM patients treated with Valganciclovir, a common antiviral used to treat CMV infection, in addition to standard therapy [78–80]. Along with this, there are several ongoing clinical trials studying the efficacy of adoptive T-cell therapy using CMV-specific T cells, or dendritic cells pulsed with CMV pp65 RNA to vaccinate GBM patients [76,77,81,82]. Promisingly, in these studies, CMV-specific T cells were able to infiltrate GBM tumors and DCs pulsed with a CMV-derived peptide from the viral pp65 were able to stimulate GBM killing via pp65-specific T cells. Most importantly, vaccination resulted in improved survival of the patients.

It is surprising that boosting CMV-specific T cells, or infusing *in vitro* expanded T cells could be effective in these individuals given that they were already CMV-positive and thus, likely had large populations of CMV-specific T cells before therapy. Indeed, Crough *et al.* demonstrated that GBM patients had normal frequencies of CMV-specific T cells

when compared with healthy donors [83]. Surprisingly however, unlike healthy donors, large proportions of CMV-specific T cells in GBM patients were dysfunctional. Importantly, upon stimulation *in vitro* in the presence of cytokines, T-cell function could be restored. Therefore, vaccination and *ex vivo* T-cell expansion may work by improving the quality of CMV-specific T cells, enabling these cells to kill infected tumors. How and why this might work are fascinating questions to be addressed, and it will be exciting to learn how this therapy progresses in the coming years.

#### **Part IV: CMV as a vaccine platform to promote continuous T-cell immunity** **CMV as a vaccine vector**

Even though CMV can cause significant morbidity in immune-compromised individuals, and can be found in a variety of human cancers, it has recently drawn interest as a potential vaccine vector because of its ability to induce memory inflation. ‘Inflationary’  $CD8^+$  T cells driven by CMV do not show signs of exhaustion in immune-competent people [26], as commonly seen in several other chronic infections [84]. CMV-driven T cells are also able to migrate into virtually all tissues

of the body at steady state [63,85]. Importantly, recombinant CMVs can be used to induce memory inflation of T cells specific for the recombinant antigens in both mice and nonhuman primates [86] and we are beginning to understand how the position of an antigen within the viral genome impacts the T-cell response [87]. Additionally, unlike many viruses, CMV is able to re-infect previously infected individuals [88], allowing for vaccination and boosting with CMV vectors even in CMV-seropositive people. Due to these traits, CMV-based vaccines are currently being developed for clinical use.

To date, CMV has been used as a vaccine vector in a few settings. MCMV was first tested for its ability to induce immunologic contraception using a recombinant virus expressing zona pellucida 3, an ovary antigen, in an attempt to control mouse populations in Australia [89]. Interestingly, this vaccine did not work by inducing zona pellucida 3-specific T cells, but rather antibodies that could sterilize the infected mice [90]. In contrast, infection with recombinant MCMVs expressing the MHC class I restricted viral epitopes from influenza nucleoprotein or LCMV-glycoprotein induced protection against Vaccinia virus expressing nucleoprotein or glycoprotein respectively that correlated with T-cell numbers in the vaccinated mice [86]. More recently, extensive studies have been performed with three recombinant Rhesus Macaque CMVs (RhCMVs) expressing the simian immunodeficiency virus (SIV) epitopes Gag, Retanef or Env [4–7]. In this work, prophylactic vaccination and boosting with these vectors induced protection and clearance of SIV in approximately half of rhesus macaques subsequently infected with SIVmac239. It is still unclear why half of the monkeys were fully protected and half were apparently unprotected. Interestingly, the RhCMV vaccine vectors stimulated SIV-specific CD8<sup>+</sup> T cells that were promiscuous and recognized uncommon SIV viral epitopes restricted by MHC II [6]. This effect seems to be related to the altered tropism of the viral vector compared with wild-type RhCMV. Whether these promiscuous T cell populations are essential for protection against SIV awaits further work. As these studies move toward clinical trials, it will be exciting to learn whether similarly constructed HCMV vectors will induce similarly targeted CD8<sup>+</sup> T cells.

These successes with CMV as a viral vaccine vector have led several groups to explore CMV-based vaccines for cancer. Although this work is at a much earlier stage, prophylactic and therapeutic vaccination with recombinant MCMV vectors have shown some efficacy in models of prostate cancer [91] and melanoma [92,93]. However, there are many remaining questions. One melanoma-targeting vaccine-induced antibodies with

therapeutic efficacy [92] while another induced tumor-specific CD8<sup>+</sup> T cells [93]. Whether one is better than the other is unknown. Interestingly, the vaccine promoting melanoma-specific CD8<sup>+</sup> T cells required an altered version of the gp100 tumor epitope to be expressed by the vaccine. Native gp100 antigen failed to induce a T-cell response [93]. Addressing whether this is a general problem with MCMV-based vectors inducing self-reactive T cells or whether this is limited to certain antigens will also require further study. Nevertheless, these recent successes using MCMV as an anticancer vaccine vector are encouraging.

Interestingly, there are data that CMV-driven NK cells and  $\gamma\delta$ -T cells may have anti-tumor effects [94,95]. Indeed, CMV reactivation has been associated with reduced risk for leukemia relapse in HSCT patients [96,97]. Thus, CMV vaccination or using CMV-stimulated NK cells and  $\gamma\delta$ -T cells in adoptive therapies may have general anticancer effects that could be exploited.

### The risks associated with CMV-based vectors & the use of crippled viruses

Most people in the world are already infected with CMV. Indeed, many people are already infected with multiple strains of CMV. Therefore, introducing a CMV-based vaccine into people would not be expected to pose additional risks that are not already inherent in the population. Nevertheless, as discussed in detail above, CMV is dangerous in immunocompromised patients, making therapeutic vaccination for late stage cancer patients possibly dangerous, especially if they are CMV-negative at the time of treatment. In addition, while a vaccine promoting immunity to a foreign pathogen like HIV would seem to pose minimal risk, the same cannot be said of immunity directed against a cancer (self) antigen. Increased safety could be achieved by using crippled viruses as the vaccine vector. We used a spread defective variant of MCMV that lacked the glycoprotein L ( $\Delta$ gL-MCMV) [60]. Glycoprotein L (gL) forms a heterodimer with the glycoprotein H (gH) within the viral envelope, and this heterodimer is essential for viral entry into cells. We produced the  $\Delta$ gL-MCMV on complementing cell lines that provided the gL protein in *trans*, thus producing virions that could go through one round of infection *in vivo*, but produce only noninfectious viral particles thereafter. Surprisingly, a single inoculation of  $\Delta$ gL-MCMV led to memory inflation of CD8<sup>+</sup> T cells and ‘inflationary’ populations that persisted for over a year. Likewise, a spread-defective variant of MCMV lacking the essential virion protein M94 was shown to persist in the lungs of immune competent mice for at least

1 year after infection [98]. Importantly, the spread-defective  $\Delta$ gL-MCMV was safe even for completely immune-deficient SCID mice [60], indicating that similarly spread-defective strains of HCMV may be safe, persistent vaccines.

### Conclusion & future perspective

CMV-specific T cells are at the core of multiple therapeutic approaches designed to control CMV, other pathogens and even cancer. The use of adoptive immunotherapy to target CMV and other opportunistic viruses is clearly a promising approach and a substantial body of research now stands as proof of principle. The work performed to date demonstrates that cultured antiviral T cells can be effective at limiting CMV reactivation and replication with limited side effects in humans. Ongoing and future work will be needed to optimize the recovery, expansion and infusion of antiviral T cells to generate the most robust and sustained responses. Emerging

clinical data and research in animal models, suggest that targeting the most proliferative memory-like T cells may improve the long-term benefits of these adoptive therapies. Thus, the selection and use of defined cell subsets may become more common in the near future. Recent work also suggests that targeting CMV by adoptive therapies or vaccination may be an effective way to target CMV-infected tumor cells. Clinical trials are ongoing with promising initial results and we would expect substantial progress to be made in the coming few years. The use of CMV as a vaccine vector is in its infancy and the first clinical trials are expected to be launched in the next several years. Much work remains to define the advantages and disadvantages of these vectors, as well as mechanistic underpinnings of success or failure in different disease settings. Although these therapies are at various stages of development and testing, they demonstrate the potential for CMV-specific T cells in a diverse group of settings.

#### Executive summary

##### Part I: cytomegalovirus reactivation & control

- Cytomegalovirus (CMV) reactivation is a constant event requiring continuous immune surveillance.
- CMV-specific T cells play a primary and fundamental role in blocking viral reactivation.
- Both CD4<sup>+</sup> and CD8<sup>+</sup> T cells contribute to the control of CMV.

##### Part II: adoptive immunotherapy to control CMV

- Long-term control of CMV in transplant patients depends on the recovery of CMV-specific CD4<sup>+</sup> and CD8<sup>+</sup> T-cell populations.
- CMV-specific T cells from healthy donors can be expanded or selected and infused into patients at risk of disease.
- CMV-specific T cells are largely effector-phenotype, poorly proliferative and short-lived as a result of their ongoing viral immune surveillance. It is unclear whether these effector-like cells contribute to the expansion of CMV-specific T cells or the sustained control of CMV in adoptive recipients.
- Small numbers of CMV-specific T cells with memory phenotype and function persist in infected hosts and these cells are superior at responding to CMV reactivation in an animal model.

##### Part III: CMV & cancer: a new avenue for adoptive immunotherapy

- CMV DNA and protein have been found in several human cancers.
- Although it is still unclear whether CMV contributed to the development of these tumors, CMV-specific immunotherapy may limit tumor growth by directing immunity to the infected tumor cells. The initial clinical trials have shown promise.

##### Part IV: CMV as a vaccine platform to promote continuous T-cell immunity

###### CMV as a vaccine vector

- Due to the enormous number of T cells elicited and sustained by CMV infection, CMV-based vectors are being developed as vaccines for infectious disease and cancer.
- The most developed CMV-based vaccines are being used to target HIV and these have shown profound efficacy in a nonhuman primate model of HIV infection.
- Recent work has highlighted the potential for CMV-based vectors in cancer vaccination.
- Crippled (single-cycle) versions of CMV may offer superior safety without sacrificing the sustained T-cell immunity elicited by wild-type vectors.

###### Conclusion

- Adoptive T-cell therapy offers clear benefits for patients at risk of CMV disease and methods will improve as various labs optimize the selection, recovery and expansion of antiviral T cells.
- Adoptive T-cell therapy or vaccination may also enable targeting of human cancers infected by CMV.
- CMV-based vaccines hold tremendous promise, but much work remains to determine whether this promise will be fulfilled.

**Financial & competing interests disclosure**

CM Snyder has a financial stake in UbiVac CMV, a company developing spread-defective CMV-based vectors for cancer vaccination. The authors have no other relevant affiliations or financial involvement with any organization or entity with a

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