

COMMENTARY



Intratumoral infection by CMV may change the tumor environment by directly interacting with tumor-associated macrophages to promote cancer immunity

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ABSTRACT

Cytomegalovirus (CMV) is a herpesvirus that induces an extremely robust and sustained immune response. For this reason, CMV has been proposed as a vaccine vector to promote immunity to both pathogens and cancer. However, exploration of CMV as a vaccine vector is at an early stage and there are many questions. Using a mouse melanoma model, we recently found that a CMV-based vaccine induced large populations of melanoma-specific T cells, but was not effective at slowing tumor growth unless it was injected directly into the tumor. These surprising results have led us to hypothesize that CMV may be adept at modulating the tumor micro-environment through its infection of macrophages. Importantly, injection of CMV into the growing tumor synergized with blockade of the PD-1 checkpoint to clear well-established tumors. Here, we discuss our results in the context of CMV-based vaccines for pathogens and cancer.

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Introduction

Effective therapeutic vaccines that elicit anti-tumor immunity are rare.¹ These failures are due to a lack of tumor associated antigen (TAA)-specific T cell infiltration into tumors, an immune suppressive environment, TAA-specific T cell dysfunction, previous immunity to viral vectors, and highly mutable tumors that may lose TAA or MHC expression.^{1,2} It has been hypothesized that cytomegalovirus (CMV) could be a vaccine vector that could overcome these deficiencies. CMV is a β -herpesvirus that establishes a systemic, but asymptomatic infection. Like all herpesviruses, CMV becomes latent in infected cells and persists for life. However, latency must be controlled by continuous, life-long immune surveillance. In the case of CMV infection, this immune surveillance leads to exceptionally large cellular and humoral immune responses,³ including gigantic populations of virus-specific CD8⁺ T cells, through a process known as “memory inflation”.^{4–10} The use of CMV as a vaccine vector, particularly for cancers, is still in its infancy, but some pre-clinical work has shown the potential of CMV-based vaccines for cancers after a systemic vaccination. Surprisingly, when we tested a similar CMV-based vaccine in a mouse model of melanoma we found no benefit until we directly infected tumors. In this review, we discuss our unexpected results and their implications for the use of CMV to modulate the tumor environment and promote anti-tumor immunity.

Memory inflation and the use of cytomegalovirus as a viral vaccine vector

CMV has several features that may make it amenable to being a viral vaccine vector. CMV is a large virus and is able to accommodate large genetic insertions in its genome.

The Mocarski laboratory showed that foreign DNA could be cloned into the viral immediate early 2 (IE2) locus without disrupting viral growth.¹¹ Therefore, recombinant antigens could be added to the CMV backbone to generate targeted immune responses. CMV can re-infect previously infected hosts, so natural CMV infections should not limit vaccination. Finally, CMV infections elicit robust immune responses.³ The Reddehase laboratory described the maintenance of large anti-viral CD8⁺ T cell populations that were sustained¹² or even enriched⁸ after the clearance of acute murine CMV (MCMV) infection. Subsequent work by the Klenerman laboratory demonstrated that persistent MCMV infection drove the slow accumulation of certain anti-viral CD8⁺ T cells, a process they termed “memory inflation”.¹³ These “inflationary” CMV-specific T cells migrate to almost any tissue in the body, including tumors,^{14–17} and are sustained by persistent antigen stimulation, which continuously promotes effector T cell differentiation.^{15,18} Despite virus persistence and repeated antigen stimulation, CMV-specific T cells do not become exhausted, even over decades of time in humans.^{17,19–21} The end result of this is that CMV-specific T cell populations can become the largest T cell populations in the circulation of healthy hosts, accounting for an average of 5% of all circulating T cells.²² Thus, in theory, CMV-based vaccines might generate robust, sustained immunity that does not need to be recalled, but is always in an effector state.

The first CMV-based vaccines

CMV was first proposed as a vaccine vector to promote immune contraception in wild mouse populations in

Australia.²³ Ultimately, this was tested with an MCMV encoding the zona pellucida 3 (ZP3) antigen, which sterilized female mice.²⁴ This effect was subsequently linked to antibodies targeting ZP3.²⁵ In addition, the Klenerman laboratory demonstrated that CD8⁺ T cell memory inflation could be elicited to recombinant antigens encoded in the CMV backbone and that these T cell responses could be protective against virus challenges.⁵ Together, these studies provided the “proof of concept” that CMV-based vaccines could promote protective immunity.

Since this initial work, CMV-vectored vaccines have been used to protect against Ebola,²⁶ respiratory syncytial virus (RSV),²⁷ and *Mycobacterium tuberculosis* (MTb)²⁸ in mouse models, and some interesting outcomes of have been described. In the RSV vaccine study, the intranasal route of vaccination was critical to park protective T cells in the lung parenchyma.²⁷ In contrast, the MTb vaccine provided resistance primarily through an NK cell response.²⁸ Most dramatically, a rhesus macaque CMV (RhCMV) encoding simian immunodeficiency virus (SIV) antigens was profoundly protective against SIV challenge in rhesus macaques.²⁹⁻³⁴ Strikingly, RhCMV vaccine vectors stimulated SIV-specific CD8⁺ T cells that were promiscuous and recognized non-canonical SIV viral epitopes restricted to MHC-II and HLA-E molecules, an effect that was traced to both evasion of class I MHC by the RhCMV vector as well as its altered tropism compared with wild-type vectors.^{10,29,30} Although much work remains to elucidate how each vaccine is providing protection and how the vaccine design can be manipulated to promote the most desired immune responses, these successes have sustained and propelled the field.

Cytomegalovirus as a viral vaccine vector for cancer

The successes with CMV vectors in the infectious disease setting led to its use as an anti-tumor vaccine. The Jarvis laboratory was the first to publish a study demonstrating efficacy of a CMV vector promoting cancer-specific immunity. In their study, systemic infection (via the intraperitoneal route - IP) with a recombinant MCMV vector expressing an immunogenic peptide from prostate-specific antigen (PSA) was able to induce memory inflation of PSA-specific CD8⁺ T cells in mice that expressed PSA as a self-antigen, and a subsequent delay in growth of a prostate tumor model expressing human PSA.³⁵ Interestingly, encoding the full-length PSA in the MCMV backbone was not as effective as encoding only the PSA₆₅₋₇₃ peptide, a result that correlated with increased PSA-specific CD8⁺ T cell responses after tumor challenge. In a second study, IP infection with MCMV expressing the melanoma protein Trp-2 induced prophylactic and therapeutic protection against B16 melanomas.³⁶ Surprisingly, the protective effect of this vaccine was antibody-dependent, and T-cell-independent. Importantly however, this vaccine worked when the vector was a “single-cycle,” spread-defective version of the vaccine that could only go through one round of infection. Thus, even a vaccine safe for immune compromised patients was effective. In another study, IP infection with MCMV expressing an altered form of the melanoma gp100 epitope (gp100^{25-33: E25K, S27P}) induced memory inflation of gp100-specific CD8⁺ T cells and subsequent prophylactic and therapeutic protection in a B16 lung

metastasis model.³⁷ Crucially, encoding the native form of gp100 in the viral backbone abrogated these effects presumably due to its failure to stimulate gp100-specific T cell responses.³⁷ Finally, a recent study has shown that MCMV encoding epitopes from the E6 and E7 proteins of human papillomavirus, delayed or prevented subsequent growth of the E6 and E7 expressing TC-1 tumor cells in a prophylactic setting.³⁸ Collectively, these experiments demonstrated that CMV-based cancer vaccines could be effective, but may work through diverse mechanisms depending on the antigen encoded in the genome. Thus, determining how to design vectors to promote the most desired anti-tumor immune responses is crucial to move the field forward, and this is an area of active investigation.³⁸⁻⁴⁴

Intratumoral MCMV infection

Keeping in line with the studies described above, we generated an MCMV-gp100 vaccine for melanoma using a modified gp100 peptide antigen (gp100^{25-33: S27P}), which promoted robust gp100-specific CD8⁺ T cell memory inflation.⁴⁵ Given that a similar MCMV-gp100 vaccine delayed the growth of B16 lung nodules³⁷ we expected to observe some impact on the growth of subcutaneously-implanted B16 melanomas. Surprisingly however, the IP route of vaccination did not delay the growth of subcutaneously-implanted B16 tumors, nor prolong mouse survival, despite promoting substantial numbers of tumor-localized gp100-specific T cells. Unsurprisingly, gp100-specific T cells recovered from the tumors were profoundly functionally impaired. We had previously shown that most “inflationary” T cells were confined to the blood and that many were stuck in the vasculature of the lungs,¹⁴ where they might have access to tumor nodules delivered by intravenous (IV) injection, possibly explaining the different outcomes. Thus, we wondered whether vaccination directly into the tumor (intratumoral - IT) would better recruit T cells to the most needed site. While the IT route of infection did not increase T cell recruitment to the tumors, it caused a striking delay in tumor growth (⁴⁵, summarized in Fig. 1). We do not believe that the virus was directly killing tumor cells (i.e. acting as an oncolytic virus) as the only infected cells that we could find in the tumor were tumor-associated macrophages (TAMs).⁴⁵ Additionally, CD8⁺ T cells were required for the therapeutic effect. Thus, we currently hypothesize that IT delivery of MCMV alters TAMs in such a way as to boost CD8⁺ T cells responses. Interestingly, the function of gp100-specific T cells in the tumor was still impaired after IT vaccination.⁴⁵ Thus, to improve the treatment, IT delivery of MCMV was combined with PD-1/PD-L1 checkpoint inhibition (Fig. 1, ⁴⁵). These therapies robustly synergized leading to tumor clearance and long-term protection in ~60% of the mice. Most surprisingly, the MCMV-gp100 vaccine was only marginally better than the wild-type MCMV at delaying tumor growth or synergizing with PD-L1 blockade. Thus, even an MCMV that did not specifically promote gp100-specific T cells was able to induce a nearly identical tumor growth delay.⁴⁵ We interpret these results to suggest that the major effect of MCMV was not to prime tumor-specific

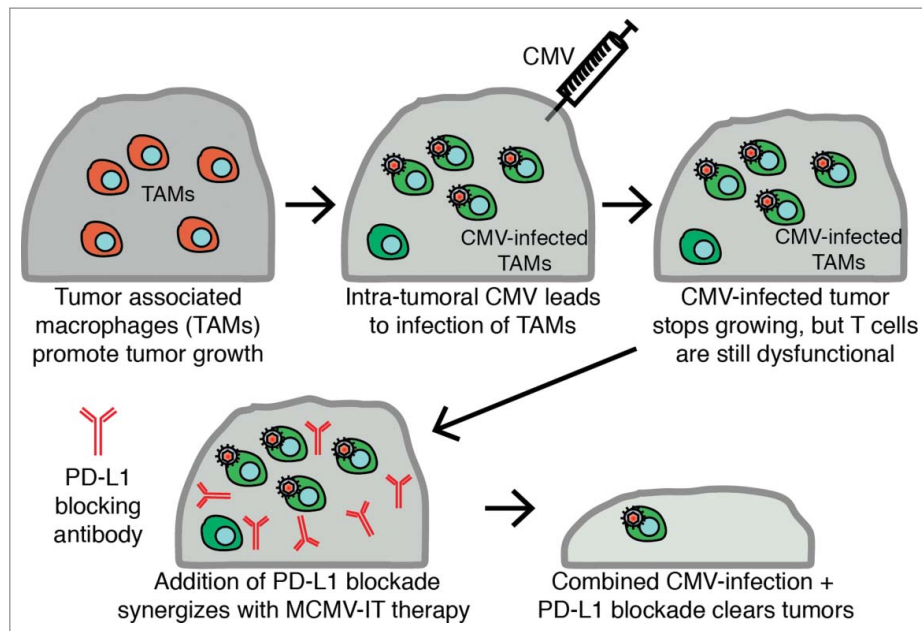


Figure 1. Schematic of Intratumoral CMV therapy. Tumor-associated macrophages (TAM) in tumors help to promote and drive tumor growth. After injecting CMV intratumorally, CMV infected TAMs and this correlated with a delay in tumor growth. Combining this intratumoral CMV infection of TAM with anti-PD-L1 therapy synergized to induce clearance of more than 60% of tumors and long-term protection.

T cells, but rather to shift the tumor to a more pro-inflammatory, anti-tumor microenvironment.

Intratumoral cancer therapies

Intratumoral administration of cancer therapeutics has been around for more than a century. Dr. William Coley first documented the use of IT therapy in the late 1800s when he isolated the bacteria that caused erysipelas and used it to treat solid tumors.⁴⁶ In 1975, successful treatment of a man with advanced melanoma using IT Bacille Calmette-Guérin (BCG) reignited the field.^{47,48} Chemotherapy, gene therapy, oncolytic viruses and immunotherapy have now been tested with varying success as IT therapies.^{46,47,49-53} IT chemotherapy and gene therapy can lessen their side effects such as non-specific cytotoxicity and severe systemic inflammation respectively. Although usually limited to easily accessible tumors, minimally invasive surgery has been used to administer IT therapies to unresectable tumors (e.g.⁵⁴).

Recent work has highlighted the potential for IT delivered oncolytic viruses. IT delivery of Herpes Simplex Virus T-VEC, which encodes GM-CSF, to metastatic melanoma became the first FDA approved oncolytic virus in the US in 2015.⁵¹ Other oncolytic HSVs that encode microRNA targets, tissue-specific promoters, or targeted attachment proteins are currently in development. In addition to T-VEC, the oncolytic adenovirus H101 is approved in China for IT administration in combination with chemotherapy for head and neck cancer.⁵¹ Additionally, the GM-CSF encoding vaccinia virus Pexavec and the coxsackievirus A21 drug Cavatek are both in clinical trials as oncolytic IT therapies.^{51,52} T-VEC, H101, Pexavec, and Cavatek all work primarily by directly infecting and lysing tumor cells when delivered IT. Our data suggest that MCMV did not work in this way and in fact, it may not have infected tumor cells much at all.

MCMV infection of tumor associated macrophages

Unlike oncolytic viruses, MCMV delivered IT seemed to primarily infect tumor associated macrophages (TAMs), which are crucial for tumor immune evasion and outgrowth and are now being targeted in several clinical trials.⁵⁵ Thus, we speculate that MCMV infection of TAMs may have altered TAMs to enable improved tumor-specific immune responses (Fig. 2). There is a clear correlation between tumor infiltration by macrophages and poor clinical prognosis in several cancers⁵⁵⁻⁵⁷ and depletion of macrophages with clodronate in mouse models has demonstrated the importance of macrophages in tumor progression and development.⁵⁸ Typically, TAMs are divided into pro-inflammatory ('M1-like') and anti-inflammatory ('M2-like'). Pro-inflammatory TAMs are thought to promote tumor-specific immunity that can eliminate tumor cells or hold tumors in equilibrium (Fig. 2). Aside from producing pro-inflammatory cytokines, and other inflammatory mediators like reactive oxygen and nitrogen species, pro-inflammatory TAMs also have high expression of MHC-I and II, enabling potent antigen presentation and activation of T cells.^{55,59} However, tumor progression and late stages of tumor growth are often associated with elevated numbers of anti-inflammatory, 'M2-like' macrophages.^{55,59} Anti-inflammatory, 'M2-like' TAMs produce high levels of IL-10 and TGF- β , lose the expression of pro-inflammatory T_H1 cytokines and MHC-II, express high levels of PD-L1, and contribute to poor CD8⁺ T cell responses.^{55,57,59} Furthermore, anti-inflammatory, 'M2-like' TAMs can release CCL22, causing T_{REG} infiltration, as well as CCL2, which causes an accumulation of TAMs, and are able to help induce and seed metastasis.^{55,57,59} Thus, TAMs are an important part of the tumor microenvironment (Fig. 2).

Macrophages are also important targets for CMV biology during normal viral infection (Fig. 2C). Both MCMV and

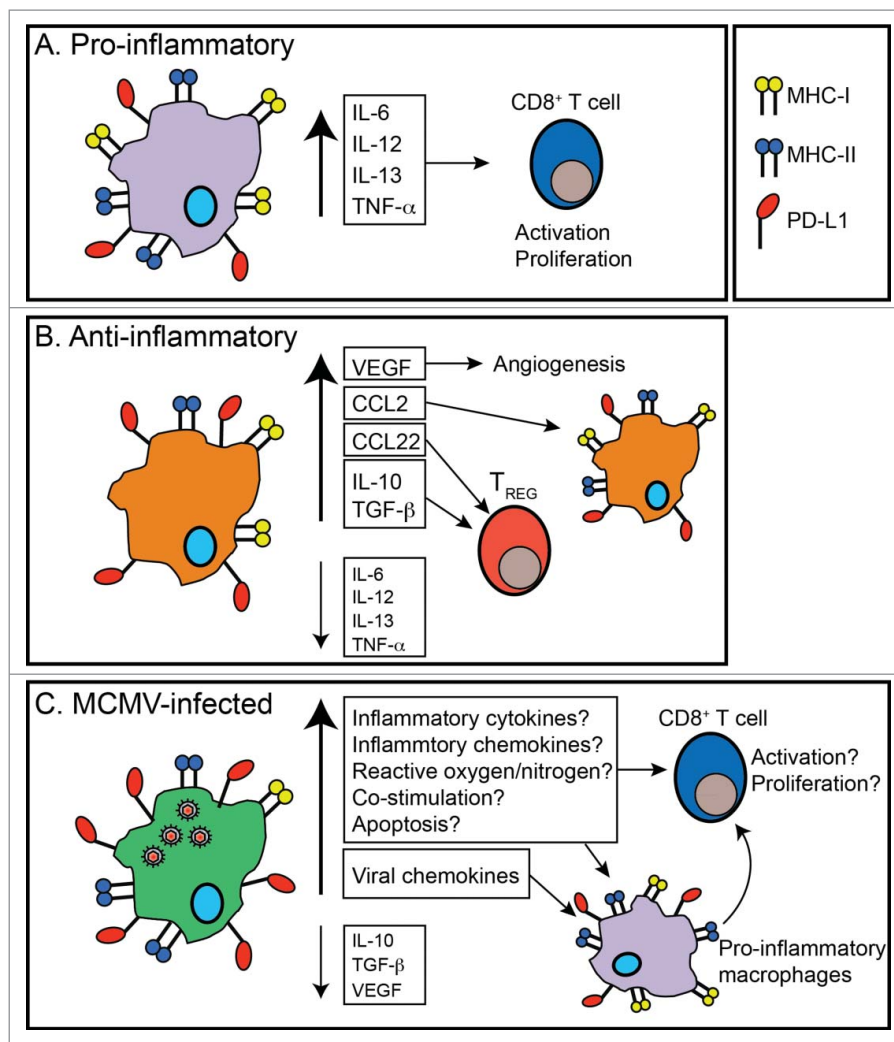


Figure 2. Tumor associated macrophage and changes induced by CMV infection. A) Pro-inflammatory tumor associated macrophages (TAM) express high levels of MHC-I, MHC-II and release pro-inflammatory cytokines. These attributes can contribute to T cell activation and proliferation, and subsequent anti-tumor effects. B) Tumor progression is associated with an accumulation of anti-inflammatory TAMs, which express lower levels of MHC-I, MHC-II and proinflammatory cytokines and increase expression of anti-inflammatory cytokines, leading to diminished T cell responses. Additionally, anti-inflammatory TAMs release VEGF, which induces angiogenesis, CCL2, which can induce the accumulation of anti-inflammatory macrophages, and CCL22, which can induce T_{REG} infiltration. Together, these attributes contribute to tumor progression. C) After IT infection with MCMV, TAM became infected. CMV infection of macrophages is known to promote a shift toward a more pro-inflammatory phenotype. Additionally, CMV encodes its own virally-encoded chemokines, which attract inflammatory macrophages. Thus, although more work is needed to define the changes in TAM, we hypothesize that MCMV infection of TAMs leads to increased T cell activation and proliferation and increased recruitment of pro-inflammatory macrophages to the tumor. Together, these attributes of TAM MCMV infection may shift the tumor microenvironment from anti-inflammatory to pro-inflammatory.

HCMV establish latency in macrophages, monocytes and progenitors of monocytes and dendritic cells.⁶⁰⁻⁷² Moreover, CMVs express several chemokines, which can directly or indirectly recruit inflammatory monocytes to the site of infection.⁷³⁻⁷⁹ In addition, HCMV promotes monocyte differentiation into immune-stimulatory, 'M1-like' macrophages^{80,81} and preferentially infects immune-suppressive 'M2-like' macrophages over 'M1-like' macrophages.⁸² After HCMV infection, macrophages release pro-inflammatory cytokines, have increased expression of toll-like receptors (TLRs) and improved MyD88 signaling,⁸⁰⁻⁸⁴ leading to increased T cell proliferation and function. Thus, MCMV infection of TAMs and recruitment of new pro-inflammatory TAMs, may shift the tumor microenvironment from immune suppressive to pro-inflammatory. MCMV infection of TAMs also may induce their apoptosis, leading to the delayed tumor growth. This, however, is unlikely in our model as there were large percentages of TAMs

present in MCMV IT infected tumors 5 d after the initial infection.⁴⁵ Additionally, MCMV infection may decrease the ability of TAMs to induce pro-angiogenic factors such as VEGF, leading to less vascularization of the tumor and slowed tumor growth. However, it must be noted that MCMV infection of macrophages may also increase VEGF production, leading to increased vascularization,⁸⁵ an effect that could promote, rather than inhibit, tumor growth. Thus, the specific interactions between CMV and the tumor environment must be dissected so that the desired outcomes can be promoted (Fig. 2C).

The interaction between MCMV and TAMs may also suggest additional avenues of synergy. For example, if MCMV IT therapy skews TAMs toward pro-inflammatory responses or depletes them, then synergy may be achieved by combining MCMV IT therapy with a drug like Paclitaxel, a common chemotherapy that enhances the infiltration of TAMs in breast cancer.⁸⁶ Likewise, since Cox-2 expression in the tumor

microenvironment correlates with anti-inflammatory TAMs, the combination of IT-MCMV and Cox2 inhibitors may be synergistic by extending the therapeutic effect beyond the directly infected cells, to further decrease anti-inflammatory TAMs.⁸⁷⁻⁸⁹

It is also possible that the PD-L1 blocking antibody was synergistic because it could alter TAMs. Indeed, PD-L1 expression was increased on both haematopoietic and non-haematopoietic cells after IT-MCMV infection.⁴⁵ Thus, it is possible that the anti-PD-L1 antibody may be depleting PD-L1^{hi} TAMs, as seen with other checkpoint inhibitors,⁹⁰⁻⁹² or altering TAMs by disrupting the PD-1/PD-L1 signaling, which has been linked to an increase in immune suppressive TAMs.^{87,88}

Finally, improving CMV's ability to kill macrophages may be ideal for improving the therapy. CMV delays apoptosis and programmed necrosis of infected macrophages *in vitro* through the expression of several genes.⁹³ Using viruses deficient in one or more of these genes might kill TAMs more quickly and have a larger effect if loss of TAMs is part of the therapeutic effect. In addition, CMVs lacking the genes that prevent cell death are much less pathogenic and therefore would be expected to be safer. Together, these approaches may offer ways to improve the efficacy and/or safety of MCMV IT therapy and may also offer a better understanding of the impact of MCMV on the tumor environment.

MCMV IT therapy's effect on CD8⁺ T cells

As CD8⁺ T cells were crucial for prolonged survival after MCMV IT therapy, and the therapy synergized with PD-L1 blockade, the effect of the on CD8⁺ T cells must be determined. It was surprising that IT delivery of both wild-type MCMV and MCMV-gp100 had nearly equivalent impacts on tumor growth and equivalent abilities to synergize with PD-L1 blockade. These data imply that IT delivery of MCMV improves the function of pre-existing tumor-specific T cells, or enables priming of new tumor-specific T cell responses. We favor the former hypothesis because IT delivery of wild-type MCMV did not promote expansion of Pmel-I transgenic (gp100-specific) T cells except in one animal.⁴⁵

Stimulation of the innate immune system by CMV

Although our data suggest that IT delivery of MCMV worked through the adaptive immune system, there is substantial data arguing for an effect of MCMV and HCMV infection on the innate immune system. Both MCMV and HCMV can promote robust NK cell activation, although our data suggest that NK cells were not crucial in the efficacy of IT MCMV infection in the B16 melanoma model.⁴⁵ Likewise, CMV has been shown to promote $\gamma\delta$ -T cells that are protective against CMV infection and can also cross-react with cancer cells, presumably due to similar expression of stress-induced ligands by infected cells and cancer cells.⁹⁴ Interestingly, early work from the Reddhead laboratory in a model of liver-adapted lymphoma showed that MCMV infection could induce apoptosis of the tumor cells in a manner that still remains unclear, but is not due to viral infection of the tumor cells, or the induction of T cells, B cells or NK cells.⁹⁵⁻⁹⁷ Thus, other innate immune activation and

alternative mechanisms to trigger tumor cell death cannot be ruled out at this time.

The Future of CMV IT therapy

MCMV IT therapy preferentially infected TAMs and induced CD8⁺ T cell-dependent improved survival of tumor bearing animals. The therapy synergized with anti-PD-L1 therapy, inducing tumor clearance and protection in over half the animals. Thus, MCMV IT therapy represents a potentially powerful anti-tumor treatment. However, there are many unknowns. The importance and effect of TAMs infection on the tumor microenvironment and the subsequent induction of CD8⁺ T cell dependent responses must be investigated to be able to improve the therapy and define the mechanisms of action. Additionally, we need to know whether the IT CMV therapy will remain effective with killed, replication-defective or spread-defective versions of CMV to develop a safe clinical reagent. It will be important to determine whether altering the immune response at one tumor site will have effects on distant tumor sites and metastases. Otherwise, the therapy would be restricted to tumors that are available for injection. Lastly, it is crucial to determine whether IT CMV therapy is effective in a multitude of different cancers, besides melanoma. Since macrophages may play distinct roles in different cancers, this type of therapy is unlikely to be equally effective in all tumor types. In sum, MCMV IT therapy represents a potentially potent therapy by acting directly on tumor-associated macrophages, but it must be better understood for future improvement.

Disclosure of potential conflicts of interest

C.M.S. has a financial interest in UbiVac CMV, for the development of spread-defective, CMV-based therapeutics.

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