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ONCOLYTIC VIROTHERAPY

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Abstract

Oncolytic virotherapy is an emerging treatment modality which uses replication competent viruses to destroy cancers. Advances in the past two years include preclinical proof of feasibility for a single-shot virotherapy cure, identification of drugs that accelerate intratumoral virus propagation, new strategies to maximize the immunotherapeutic potential of oncolytic virotherapy, and clinical confirmation of a critical viremic thereshold for vascular delivery and intratumoral virus replication. The primary clinical milestone was completion of accrual in a phase III trial of intratumoral herpes simplex virus therapy using talimogene laherparepvec for metastatic melanoma. Challenges for the field are to select 'winners' from a burgeoning number of oncolytic platforms and engineered derivatives, to transiently suppress but then unleash the power of the immune system to maximize both virus spread and anticancer immunity, to develop more meaningful preclinical virotherapy models and to manufacture viruses with orders of magnitude higher yields compared to established vaccine manufacturing processes.

REVIEW ARTICLE

Oncolytic viruses are therapeutically useful anticancer viruses that will selectively infect and damage cancerous tissues without causing harm to normal tissues¹. Each virus has a specific cellular tropism that determines which tissues are preferentially infected, and hence, what disease is caused. Rabies virus, for example, damages neurons, hepatitis B virus damages hepatocytes, HIV damages helper T lymphocytes and influenza virus damages airway epithelium. Many, if not most, naturally occurring viruses have a preferential, although nonexclusive, tropism for tumors and tumor cells. This probably has more to do with tumor biology than with virus biology since most tumors have evolved not only to avoid immune detection or destruction, but also to resist apoptosis and translational suppression, which are the key responses used by normal cells to limit a virus infection. Oncolytic viruses can kill infected cancer cells in many different ways, ranging from direct virus-mediated cytotoxicity through a variety of cytotoxic immune effector mechanisms. Conventional concepts of cell death (apoptosis, necrosis or autophagy) are generally inadequate to fully describe the complex cell killing scenarios encountered in virotherapy. This is because the oncolytic virus typically takes over and controls the molecular cell death machinery of the

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infected cancer cell, allowing death to occur only after available cellular resources have been maximally exploited for the synthesis and assembly of new viruses². In addition to the killing of infected cells, oncolytic viruses can mediate the killing of uninfected cancer cells by indirect mechanisms such as the destruction of tumor blood vessels, the amplification of specific anticancer immune responses or through the specific activities of transgene-encoded proteins expressed from engineered viruses¹.

Specific targeting of cancer cells is obviously the sine qua non for oncolytic virotherapy and can be achieved in several ways. Some viruses such as H1 autonomously replicating parvoviruses, reovirus, Newcastle Disease Virus, Mumps virus, Moloney leukemia virus have a natural preference for cancer cells, whereas such as measles, adenovirus, Vesicular Stomatitis Virus, vaccinia and Herpes Simplex Virus can be adapted or engineered to make them cancer-specific. Surface markers such as EGF receptor, Her2-neu, Folate receptor, Prostate Specific Membrane Antigen and CD20, and nuclear transcription factors PSA, hTERT, COX-2, osteocalcin expressed selectively by tumor cells can be targeted by using them as receptors for virus entry or as essential cofactors for viral gene expression^{3, 4}. Alternatively, oncolytic viruses can be engineered to exploit the defective antiviral defenses of tumor cells as explained below⁵. Normal cells respond to virus infection by downmodulating their metabolism and/or by undergoing apoptosis, thereby inhibiting virus propagation. Successful viruses use a variety of strategies to combat these innate immune responses, but become non-pathogenic when engineered or evolved to incapacitate their immune combat proteins. Examples include the VSV matrix protein, the NS1 protein of influenza virus, the C and V proteins of paramyxovirus family members, the HSV γ 34.5 protein and the proteins encoded in the E1 and E3 regions of the adenovirus genome. Interestingly, as the apoptotic and antimetabolic responses of tumor cells are generally deficient, attenuated viruses with defective immune combat proteins often retain their ability to propagate in tumor cells. An alternative way to 'target' viruses to cancer cells is to selectively eliminate their undesirable tropisms by engineering targets for brain, liver or muscle-specific microRNAs into their genomes such that the viral life cycle is selectively blocked in the relevant target tissue⁶.

Here we provide a critical overview of the current state of the field of oncolytic virotherapy research, emphasizing what we consider to be the most important recent advances and the main challenges going forward. The review is divided into three sections. The first section reviews the clinical oncolytic virotherapy experience to date and illustrates that the approach has genuine promise but that its full potential has yet to be realized. The subsequent sections address the two key stages of a successful oncolytic virotherapy treatment episode, both of which are truly hotbeds of preclinical research innovation: first, delivery of the virus to the tumor; and second, spread of the virus infection through the tumor. Optimizing the efficiency and accuracy of both of these critical processes will doubtless continue to challenge the field for years to come, but there have been many recent developments, several of which are already being translated to determine whether they can improve clinical outcomes.

Because of space constraints, citations have been limited to key manuscripts published since 2007. However, in some cases, we refer to seminal papers published before this time. Where

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multiple primary manuscripts address the same topic, because of space limitations, we have cited review articles. We apologize to those investigators whose work has not been cited and take full responsibility for these omissions.

Clinical development

The idea of using viruses to treat cancer first began to take hold In the 1950s when tissue culture systems and rodent cancer models were originally developed⁷. Hundreds of cancer patients were treated with impure oncolytic virus preparations (even infected body fluids) administered by almost every imaginable route⁸. The viruses were usually arrested by the immune system and did not affect tumor growth, but sometimes the infection took hold and tumors regressed, especially in immunosuppressed patients, although they frequently became sick or died when the infection spread to normal tissues. In one particularly promising study from Osaka University, tumor regressions were reported in 37 of 90 terminal cancer patients treated with a non-attenuated mumps virus⁹. But this work was not continued beyond the 1970s and the strains of mumps virus used for the work have since been lost (T. Asada, personal communication). The modern era of oncolytic virotherapy, in which virus genomes are engineered to enhance their anti-tumor specificity, can be traced to a 1991 publication in which a thymidine kinase (TK)-negative herpes simplex virus (HSV) with attenuated neurovirulence was shown to be active in a murine glioblastoma model¹⁰. Since that first application of virus engineering to an oncolytic HSV, the pace of clinical activities has accelerated considerably, with numerous ongoing or completed trials using oncolytic viruses belonging to at least 10 different virus families (Table 1), and a steady stream of new oncolytic viruses entering the clinical arena^{11–13}.

The clinical tolerability of oncolytic viruses has overall been excellent, even at today's highest feasible doses¹⁴. Even so, future oncolytic virus trials will likely use even higher doses as manufacturing yields are continually increasing due to a variety of technical advances, such as cell substrate optimization or the use of cell microcarriers and disposable wave bioreactors^{15–17}. Hence, it may be premature to judge whether effective oncolytic virotherapy will be devoid of serious toxicities at clinically effective doses. One unique safety risk is the concern that an oncolytic virus might spread from the treated patient and mutate to regain its pathogenic potential¹⁸. However, although virus shedding has sometimes been documented in urine or respiratory secretions, oncolytic virus transmission to contacts and carers has not yet been seen¹⁴.

Clinical efficacy—Evidence for efficacy of single-agent oncolytic virotherapy comes from two recent phase 1/2 clinical trials backed up by an number of quite compelling anecdotal reports^{7, 19–21}. In one trial, talimogene laherparepvec, formerly named OncoVEX, which is an oncolytic HSV coding for granulocyte/macrophage colony stimulating factor (GM-CSF) was administered by direct intratumoral injection to patients with metastatic malignant melanoma and led to complete regressions of injected and uninjected lesions in 8 of 50 treated patients¹⁹. This study remains the most compelling demonstration that intratumoral administration of an oncolytic virus can powerfully cross prime and amplify anticancer immunity. Perhaps because of its well-known susceptibility to immunotherapy,

In the second trial, an oncolytic vaccinia virus, JX594, also engineered to express GM-CSF, was administered intratumorally to patients with non-resectable hepatocellular carcinoma, leading to objective responses in 3 of 10 evaluable patients²⁰.

Trials combining oncolytic viruses such as reovirus, vaccinia and HSV with drugs or radiation are giving a high frequency of tumor responses^{21, 23–25}, but it is difficult to know whether the oncolytic viruses are contributing to these responses over and above the active anticancer drugs with which they are being combined. Only through randomized phase 3 trials can this critical question be answered.

So far, clinical trials have failed to provide a clear demonstration that direct viral lysis of infected cells is an important mechanism of tumor destruction¹⁴. Thus, the oncolytic paradigm, where a systemically administered virus spreads extensively at sites of tumor growth to cause tumor destruction, remains to be proven. Of major significance in relation to this point, it has recently been determined in a phase I clinical trial that intravenously administered JX594 was recoverable from tumor biopsies only when the viremic threshold dose of 10⁹ infectious units was exceeded²⁶. The critical insight gained from this study is that systemically administered oncolytic viruses can specifically target sites of tumor growth by extravasating from tumor blood vessels and replicating in the tumor. This is a concentration driven process and is therefore detectable only above a threshold virus dose. Direct oncolytic tumor destruction may therefore be tightly linked to the dose of virus administered which for many oncolytic viruses is limited primarily by manufacturing considerations.

Additional insights have been gained from the ongoing and completed clinical trials. First, since the clinical trial results often fall short of hopes and expectations, it is clear that better, more reliably predictive, preclinical models are needed. Specifically there is a need for orthotopic cancer models in immunocompetent animals that are not only susceptible to the oncolytic virus being evaluated, but also mirror the human pathogenesis of the viral infection. Current models are inadequate because they often lack an immune system (cultured cells and human xenograft models), or are not susceptible to the virus in question, although exceptions do exist (e.g. vaccinia).

A second additional insight is that iterative phase I clinical trials may become standard practice in the oncolytic virotherapy field. Conventional drugs are typically perfected before they enter clinical testing but oncolytic viruses are more akin to motor cars with multiple component parts, all of which are constantly subject to improvement, refinement and perfection through engineering efforts. Iterative phase I trials provide the only mechanism whereby the steady stream of new engineering modifications that only slightly change the product specification can be accommodated into the development pipeline.

Another important recent insight from the clinic is that it is clinically feasible to monitor virus spread by reporter transgene expression monitoring, a useful source of pharmacokinetic data which can be especially helpful during early stage clinical

development. The progress of an oncolytic virus infection can be monitored in rodents by post-mortem analysis of the changing biodistribution of virus-infected cells at multiple timepoints, but this is impractical in oncolytic virus-treated human subjects. Hence, little has been learned of the reasons for inferior outcomes of oncolytic virotherapy in humans versus rodents. Reporter genes have therefore been engineered into oncolytic virus genomes to facilitate repetitive, noninvasive determination of the number and location of virus-infected cells in the body ^{27, 28}.

When we administered an oncolytic measles virus encoding the soluble extracellular domain of carcinoembryonic antigen (MV-CEA) intraperitoneally to patients with refractory ovarian cancer, serum CEA monitoring studies suggested that the virus infected only a small number of tumor cells, and was not undergoing significant amplification in $vivo^{29}$. Reporter genes compatible with radioactive tracers have also been tested in humans. Oncolytic HSVs are amenable to PET imaging via the HSV TK, which phosphorylates specific positron-emitting substrates, trapping them inside the cell, as demonstrated in a clinical trial of HSV TK gene therapy for glioblastoma³⁰, although clinical validation for tracking the spread of a replication competent oncolytic virus is still awaited. The gene encoding the thyroidal sodium iodide symporter (NIS), which concentrates radioactive iodide, has been inserted into the genomes of several oncolytic viruses such as adenovirus, measles, VSV, HSV, and vaccinia have been used preclinically in conjunction with various radioisotopes $(^{125}I, ^{123}I, ^{124}I \text{ and } ^{99m}TcO_{4})$ to monitor *in vivo* spread³¹. This versatile NIS approach was recently validated in a clinical study in which ^{99m}TcO₄-based SPECT/computed tomography (CT) imaging was used to monitor the intratumoral spread of an oncolytic adenovirus coding for NIS³². In an approach known as radiovirotherapy, we demonstrated that it will also be feasible in the future to increase the potency of a NIS-expressing virus by administering ¹³¹I, which delivers high-energy beta particles, into the infected tumor³³.

Delivering oncolytic viruses to the tumor

Although several ongoing trials are emphasizing intratumoral delivery, systemic delivery will be absolutely required for treatment of metastatic cancer. The goal of systemic therapy is to exceed the 'viremic threshold' above which the virus nucleates a critical number of intratumoral infectious centers whose expansion and coalescence leads to tumor destruction. Current research is therefore focused on minimizing oncolytic virus sequestration in liver and spleen, evading neutralization by serum factors, targeting the viruses to the vascular endothelial cells lining tumor blood vessels and selectively enhancing their permeability.

Minimizing sequestration in the liver and spleen—Intravenously administered viruses are rapidly cleared from the circulation as a result of sequestration by the mononuclear phagocytic system (MPS) in the liver and spleen. Before clearance, they are typically coated (opsonized) with antibodies, complement, coagulation factors and/or other serum proteins that facilitate their recognition by splenic macrophages and hepatic Kupffer cells. These 'decorated' particles bind to receptors (e.g., Fcγ receptors, complement receptor 1 (CR1), CR3 or scavenger receptors) on macrophages/endothelial cells, resulting in receptor-mediated phagocytosis and accelerated clearance from the circulation³⁴. Some

viruses, for example adenoviruses, can bind directly to scavenger receptors on Kupffer cells inducing proinflammatory cytokines that can result in serious dose-limiting toxicities^{35, 36}.

Strategies to minimize sequestration include chemical modification of the coat proteins of the viruses by conjugation of biocompatible polymers, such as polyethylene glycol (PEG) and N-[2-hydroxypropyl]methaacrylamide (HPMA)^{37, 38}. Both PEG and HPMA are already used clinically to prolong the circulation times of proteins and liposomes and to reduce off target toxicities³⁹. Polymer coating can destroy virus infectivity, which can be restored by re-engineering receptor binding ligands onto the surface of the shielded particles⁴⁰. For example, PEGylation of VSV glycoprotein pseudotyped lentiviral vectors increased vector circulation half-life fivefold and dramatically inhibited complement inactivation⁴¹. PEGylated adenovirus 5 (Ad5) is cleared fourfold slower than unmodified Ad5⁴². The length of the PEG influences outcome; 20 kDa PEG but not 5kDa PEG can detarget oncolytic Ad5 from Kupffer cells and hepatocytes without inducing liver enzymes⁴³. HPMA-cloaked adenovirus vectors are also protected from neutralizing antibodies and have a prolonged circulatory half-life⁴⁴. An alternative approach to minimize sequestration of viruses (e.g. HSV) that are readily bound by IgM and complement proteins is to deplete these serum factors by pretreating with cobra venom factor or cyclophosphamide⁴⁵⁻⁴⁷.

Virus sequestration by the MPS is saturable⁴⁴. Sequestration by the liver and spleen can therefore be inhibited either by pre-conditioning to saturate MPS scavenger receptors or by poisoning the macrophages/endothelial cells. Predosing of mice with polyinosinic acid, which binds to scavenger receptors on endothelial cells or macrophages in the liver and spleen, can reduce MPS sequestration of adenoviruses⁴⁸. Clodronate-loaded liposomes can also deplete liver Kupffer cells and splenic macrophages of mice. Oncolytic adenoviral therapy has been combined with clodronate liposomes for depletion of Kupffer cells to enhance therapeutic outcome^{49, 50} Other MPS blocking strategies include preadministration of gadolinium chloride (GdCl3) or gamma globulins^{42, 51}. In one study, GdCl3 prolonged the circulatory half-life of an Ad5 vector, with a 100-fold difference in blood levels at 60 minutes⁵¹. Predosing with high doses of intravenous adenoviral particles is toxic to Kupffer cells, which decline substantially in numbers by 4 hours, greatly reducing MPS clearance of a second dose^{52, 53}.

Evading neutralization by serum factors—Many of the barriers viruses encounter following intravenous administration (e.g. neutralizing antibodies, inactivation by complement or scavenging by Kupffer cells) can be overcome by hiding oncolytic viruses inside carrier cells. Two approaches have shown promise in pre-clinical models: infusing *ex vivo* infected tumor cell lines⁵⁴ or using normal primary cells that can home to tumor beds⁵⁵. Permissive tumor cells have the advantage of being easy to propagate and genetically modified, are productive virus factories *in vivo* and could in theory be used as an 'off the shelf' product. In one study, we demonstrated that lethally irradiated myeloma cells infected with an oncolytic measles virus were therapeutically potent when administered intravenously to myeloma-bearing mice with protective titers of antimeasles antibodies⁵⁶.

Mesenchymal stem cells⁵⁷ (MSCs) are another cell type that have been used both preclinically and in small clinical trials⁵⁸ to deliver oncolytic viruses to tumor beds. MSCs

have been shown to preferentially engraft into solid tumors⁵⁹ and we recently have shown they could efficiently deliver oncolytic measles viruses to intraperitoneal ovarian cancer deposits in the presence of neutralizing antiviral antibodies⁶⁰.

Cellular carriers should ideally be combined with oncolytic viruses that will not kill the carrier before it has infiltrated into the tumor. Some viruses can piggyback on cells found normally in the circulation. Dendritic cells and T cells when admixed with reovirus, carried and delivered their oncolytic cargo even in the face of neutralizing antibodies^{55, 61}. VSV and measles virus can be delivered to tumor beds by loading onto T cells and when bound to these cells, VSV particles are protected from neutralizing antibodies^{62, 63}. Technology for routine isolation of assorted white cells from blood products is widely available clinically and thus may make the implantation of carrier cell approaches in the clinic more practical going forward.

Selectively increasing the permeability of tumor blood vessels—The EPR effect was first described in 1986 when it was shown that the leaky vasculature of tumors could be exploited to allow the passage of macromolecules from the lumenal side of the blood vessels into tumor tissues^{64, 65}. Leakiness is due to the presence of fenestrae (50–80 nm) and intercellular gaps between tumor endothelial cells (200–900 nm compared with 2–6 nm in normal blood vessels), facilitating extravasation of macromolecules, viruses and nanoparticles^{66–69}. However, poor lymphatic drainage and dense stromal tissue increase the interstitial fluid pressure in tumors impeding virus extravasation and diffusion.

Vascular permeability can be increased by preadministration of interleukin 2 (IL-2), tumor necrosis factor alpha (TNF- α), histamine or a bradykinin analog^{65, 68, 70}. Giving chemotherapy can reduce the intratumoral interstitial fluid pressure by killing tumor cells, thereby enhancing extravasation without directly impacting vessel permeability⁶⁸. In one recent study, a combination of vascular endothelial growth factor (VEGF) and metronomic doses of paclitaxel or cisplatin increased the vascular permeability of the tumor endothelium and improved the delivery of Sindbis vector to tumors⁷¹. Multiple injections of VEGF165 resulted in superior reovirus infection of proliferating tumor endothelium, thereby increasing therapeutic activity of in a syngeneic B16 melanoma model⁷². Systemic IL-2 accompanied by depletion of T regulatory cells (Tregs) also enhanced the extravasation of oncolytic viruses in B16 metastases in the lungs of mice⁷³.

Targeting the viruses to tumor vessel endothelium—In addition to being structurally different from normal vessels⁶⁸, the tumor vasculature is antigenically distinct^{74–76}. Targets visible from the lumenal side include antigens overexpressed on tumor endothelial cells (e.g., $\alpha_v\beta_3$ integrins, VEGF receptor 2, prostate-specific membrane antigen (PSMA), urokinase plasminogen activator receptor, phosphatidylserine, E-selectin, vascular cell adhesion molecule (VCAM), tissue factor, endosialin and endoglin (CD105)^{75, 76}. High-affinity protein or peptide ligands are available for targeting most of these endothelial markers. Other targets include structural elements that are exposed during vessel formation and remodeling, for example, laminin (targeted by L36 single chain Fv (scFv)⁷⁷) and fibronectin (targeted by the E19 scFv)⁷⁸. Importantly, most of the above mentioned vascular targets are not expressed exclusively on tumor blood vessel endothelium.

Chemical or genetic modifications to oncolytic viruses have been used to selectively target the tumor cell surface, de-target sensitive tissues or create dual target viruses to enhance both vascular targeting and tumor infection (transductional targeting). For instance, we previously genetically modified measles virus⁷⁹ to display a variety of polypeptide ligands on its surface facilitating infection of tumor cells overexpressing the targeted receptor³. Polymer coating has been used to mask natural attachment proteins, re-directing virus infection by chemically coupling therapeutic antibodies (e.g., cetuximab) that bind tumor cells⁸⁰.

A scFv against E selectin was conjugated onto polymer-coated adenoviral particles to enhance their binding to activated endothelial cells in inflamed areas or in tumors⁸¹. Oncolytic measles viruses expressing vascular targeting peptides, the amino terminal fragment of urokinase plasminogen activator, or cyclic RGD and echistatin, which bind to $\alpha_{\nu}\beta_{3}$ integrin receptor, were shown in our laboratory to infect tumor vessel endothelial cells *in vivo*^{82, 83}. Echistatin binds the $\alpha_{v}\beta_{3}$ receptor with 1000-fold higher affinity than cyclic RGD⁸⁴ and this is associated with enhanced ability of the virus to infect tumor vasculature. Recently, we also reported that VSV can naturally interact with tumor blood vessel endothelium in CT26 colorectal tumors in Balb/c mice⁸⁵. By 24 hours post intravenous infusion of VSV, vascular perfusion had shut down at the core of the tumors and VSV antigen was detected in the blood vessels. To improve transfer of oncolytic viruses from blood vessel lumen to tumor parenchyma, the concept of heterocellular fusion between endothelial cells and underlying tumor cells has also been explored⁸⁶. Adenoviral vectors encoding a fusogenic membrane glycoprotein driven by the human endothelial receptor tyrosine kinase promoter have been shown in vivo to trigger fusion between endothelial cells and epithelial cells, facilitating transendothelial virus penetration⁸⁶.

Intratumoral oncolytic virus spread

Mammalian cells have evolved to resist virus infections. A typical infection involves attacks on cellular defenses by viral gene products (virulence proteins), defensive parries by the host cell through the elaboration of anti-viral proteins and further counter attacks by the virus. Viral virulence genes encode proteins that suppress host defense systems, facilitate virus spread between cells and usurp cell metabolic processes. Oncolytic viruses are selected or engineered to be attenuated in normal tissues often by mutation or deletion of virus virulence genes^{75, 87}. Thus, an oncolytic virus entering a normal cell triggers the cellular anti-viral response but is unable to counterattack so the infection is quickly eliminated. The antiviral response involves production of proteins that counteract the virus by acting directly against the virus^{88, 89}, communicating with adjacent cells⁹⁰ or jump-starting apoptotic programs⁹¹. Interferons and their receptors are key players in this anti-viral response, reprogramming the physiological properties of infected and surrounding cells, inducing cell cycle arrest, providing anti-angiogenic signals, promoting apoptosis, inhibiting protein synthesis and activating the immune system.

In contrast to normal cells, the successful tumor cell has often eliminated/inactivated key gene products that have the dual role of controlling critical cell growth/death programs *and* aiding in resisting virus infections⁹². Because of these tumor-specific mutations, oncolytic

viruses—despite their defective virulence genes—can initiate productive infections in cancer cells. Occasionally, cancer cells are completely devoid of antiviral activity⁸⁷ but partial inactivation is more common resulting in only limited sensitivity to oncolytic virus therapy⁹³.

Enhancing oncolytic virus growth in tumors through genetic arming-

Incorporation of virulence genes from other virus strains or repairing previously attenuated/ deleted virulence genes can overcome residual antiviral responses found in some tumors^{93, 94}. Although this could compromise excellent safety profile of oncolvtic viruses, it is expected that the approach can be finetuned to enhance clinical oncolytic virus therapy. A related approach to improve oncolytic virus potency is to combine viruses with complementing virulence proteins⁹⁵. An interferon-sensitive oncolytic vesicular stomatitis virus (VSV) replicated efficiently in refractory tumor cells when they were co-infected with an oncolytic poxvirus encoding a secreted interferon antagonist. We achieved a similar 'ping pong' synergy effect using an engineered fusogenic VSV to accelerate the intratumoral spread (through cell fusion) of an oncolytic poxvirus⁹⁵. Because VSV has an RNA genome and vaccinia has a DNA genomes, the exchange of virulence genes leading to a pathogenic 'super virus' seems very unlikely. Clinical development of poxviruses encoding an interferon antagonist (B18R protein) is currently underway⁹⁶ and two interferon-responsive oncolytic viruses (Oncovex and Reolysin) are in advanced stage trials. If approved as single agents, these agents could be combined to create more effective oncolytic virus therapy regimens.

Chemical sensitizers to enhance oncolytic virus growth in tumor cells-

Another tactic to neutralize residual antiviral activities in oncolytic virus resistant cancers is through the use of small molecules. Several groups (including our own) have shown that histone deacetylase inhibitors (HDAC inhibitors) can suppress the residual IFN responsiveness of tumor cells, thereby increasing oncolytic virus potency without compromising specificity^{97–100}. We have identified additional compounds enhancing oncolytic virus growth in refractory tumor cells using high-throughput screening. The previously uncharacterized compound, 3,4-dichloro-5-phenyl-2,5-dihydrofuran-2-on, whose mechanism of action remains the subject of intense ongoing investigations, enhanced the growth of various oncolytic viruses on a spectrum of tumor cells by blunting their interferon¹⁰¹. Several chemically unrelated compounds were isolated in this screen but their cellular targets have not yet been defined. A similar screen with an attenuated herpes simplex virus (HSV) lacking ribonucleotide reductase identified two molecules (dipyridamole and dilazep) that inhibit the cellular equilibrative nucleoside transporter-1 (ENT1), thereby inducing cellular ribonucleotide reductase¹⁰².

Rapamycin potentiates the growth of several oncolytic viruses in rodent tumor models, mainly by disrupting the TORC1 [TOR (target of rapamycin) complex 1]-dependent production of interferon and/or disrupting the phosphotidylinositol 3-kinase AKT pathway^{103–105}. Cyclophosphamide has also been shown to improve oncolytic virus efficacy through several mechanisms. It dampens the innate antiviral response, slows the generation of anti–oncolytic virus neutralizing antibodies, may target T-regs and may affect tumor

vasculature enhancing oncolytic virus extravasation^{106–108}. Even so, combining drugs with viruses is not without risk and could promote off-target infections, compromising safety¹⁰⁷.

Improving virus spread in tumors—Some oncolytic viruses are particularly well equipped to spread within and between tumors. For instance, vaccinia virus generates multiple virus 'subspecies' adapted in different ways for efficient spread. The extracellular enveloped or 'cloaked' form (EEV) facilitates widespread dissemination and to some extent avoidance of neutralizing antibodies¹⁰⁹ whereas the cell associated (CEV) form has an actin tail that propels the virus into adjacent tumor cells¹¹⁰. Other viruses, such as oncolytic measles, spread by fusing infected with uninfected cells^{111, 112}. Engineering this capacity (cell fusion) into other virus platforms can improve therapy¹¹³ but can also lead to increased unwanted pathology¹¹⁴.

Movement of viruses through tumors can be impeded by dense intratumoral connective tissue^{115, 116}. Recently, it has been shown that losartan, an US Food and Drug Administration (FDA; Silver Spring, MD)-approved angiotensin II receptor antagonist/ antihypertensive agent can enhance the intratumoral spread of an oncolytic HSV by disrupting transforming growth factor beta 1 (TGF- β 1) signaling, which decreases collagen production, although several weeks of antifibrotic activity would be needed to impact clinical oncolytic virus outcomes¹¹⁷. Hyaluronan is a sulfated glycosaminoglycan and key component of the tumor extracellular matrix. Injecting hyaluronidase into tumors enhances the spread and efficacy of oncolytic adenoviruses¹¹⁸. A hyaluronidase-expressing oncolytic adenovirus demonstrated improved spread and activity in a human melanoma xenograft model¹¹⁹. Lastly, damage caused to tumors by cytotoxic agents, radiation or apoptosis inducers can lead to creation of voids and channels that facilitate virus spread¹¹⁶.

Engineering tumor selectivity into oncolytic virus backbones—Many of the earliest engineered oncolytic viruses were based upon the adenovirus backbone and were designed to take advantage of 'tumor-specific' promoter elements¹²⁰. For example, the telomerase reverse transcriptase promoter is inactive in essentially all adult somatic tissues but is robustly expressed in cancer cells¹⁰². An alternative strategy is to use a promoter element that targets both the cancer and an expendable adult tissue (e.g., prostate¹²¹). This approach was extended to HSV¹²² and more recently to replication-competent retroviral oncolytic vectors¹³. Expression profiling is providing new leads for promoters that could be used for oncolytic virus regulation¹²³. Poxviruses are not amenable to transcriptional targeting because they replicate entirely in the cell cytoplasm and regulate their transcription independently of the host cell transcriptional machinery.

Transductional targeting (discussed previously) can also be used to eliminate toxicities, particularly when the oncolytic virus binds an ubiqutious receptor. VSV was pseudotyped with the surface glycoprotein from a non-neurotropic lymphocytic choriomeningitis virus (LCMV) or retargeted measles virus, thereby eliminating its neurotoxicity without compromising its ability to infect and kill cancer cells^{124, 125}. Modification of the hypervariable loop of the adenovirus hexon protein ablates the ability of that virus to infect normal hepatocytes but not tumor cells¹²⁶.

Given the potential off-target effects of transcriptional and transductional targeting, other tropism modifying strategies are of interest. One exciting new strategy is the application of microRNA targeting to oncolytic viruses^{127, 128} which takes advantage of differential expression of certain microRNA species in tumor and normal tissues. Insertion of liver-specific microRNA binding sites in the 3' untranslated region (UTR) of the E1A gene of an oncolytic adenovirus eliminated its hepatotoxicity without destroying tumor cell killing activity^{128, 129}.

MicroRNA regulation of oncolytic virus tropism was first described in RNA viruses that cannot be controlled through transcriptional targeting 130-132. Particularly dramatic are the results with coxsackievirus A21 (CVA21), a very potent oncolytic virus in mice that also causes fatal myositis due to off-target infection of normal muscle. We demonstrated that inclusion of muscle-specific microRNA targets into the 3' UTR of CVA21 eliminated muscle toxicity but did not compromise anticancer activity¹³¹. One potential issue with this approach is that microRNA targets can mutate during oncolysis so it may be prudent to use a second selectivity strategy to minimize the chances of toxic escape variants arising during therapy. An oncolytic adenovirus was regulated by both transcriptional targeting (telomerase promoter) and microRNA targeting of the E1 gene¹³³. Dual targeting approaches may facilitate the generation of potent but highly specific oncolytic strains encoding wild-type virulence proteins. The positioning of microRNA targets is another critical determinant for their effectiveness in attenuating virus replication. Inclusion of microRNA targets in VSV can eliminate unwanted neurotoxicity in mice; however, it is successful only when positioned at the extreme end of the viral genome controlling the expression of the L or polymerase gene of the virus 130 .

Oncolytic virus replication can also be targeted by regulation of viral protein translation. The potently oncolytic chimeric poliovirus, PVS-RIPO (live-attenuated poliovirus type 1 (Sabin) vaccine containing an internal ribosome entry site (IRES) element from human rhinovirus type 2), lacked neurotoxicity because translation from the inserted rhinovirus IRES is selectively blocked in neurons¹³⁴. Translational control through the IRES element of another picornavirus, encephalomyocarditis virus, appears to play a role in its oncolytic specificity¹³⁵. The level and activation state of eukaryotic initiation factor 4E (eIF4E) contribute to the initiation and progression of a variety of cancers and some viruses actively promote eIF4E activation¹³⁶. Incorporation of complex 5[°] UTRs responsive to the levels of cellular eIF4E has therefore been used to target HSV¹²² and adenovirus¹³⁷.

Conditional manipulation of protein stability is also being used to regulate oncolytic virus expression. Fusion of a 'destabilizing domain' was used to create chimeric proteins that are inherently unstable¹⁴⁹¹³⁸. A cell-permeable synthetic small molecule ligand called Shield-1 ((S)-(R)-3-(3,4-dimethoxyphenyl)-1-(3-(2-morpholinoethoxy)phenyl)propyl 1-((S)-2-(3,4,5-trimethoxyphenyl)butanoyl)piperidine-2-carboxylate), can bind the destabilizing domain and reverse this instability allowing regulated production of imaging proteins (e.g., luciferase) or TNF- α in animal tumor models^{138–140}. This provides an attractive system for fine-tuned expression of therapeutic transgenes to control the spread of an oncolytic virus.

Controlling adaptive immunity and clearance of oncolytic viruses—When reviewing the history of the oncolytic virus field, it is notable that immunosuppressed patients have generally responded better to oncolytic virus therapy than those with an intact immune system, but this increased oncolytic activity was often associated with unacceptable toxicity⁷. Impairment of the adaptive antiviral immune response is therefore a double-edged sword but can be used to advantage, provided the virus is so specific for the tumor that it cannot damage normal tissues. Virus-targeting technologies have now advanced to the point where combining virotherapy with immunosuppressive drugs has become an appealing approach by which to enhance their antitumor activity. There are many immunosuppressive drugs to choose from in this regard, but cyclophosphamide is currently the most favored because it is potently toxic to both T and B lymphocytes, has direct antitumor activity, has been very widely used since 1949 for both cancer therapy and for immunosuppression, and is very reasonably priced¹⁴¹. Several preclinical studies have shown that cyclophosphamide can retard immune clearance of oncolytic viruses, enhance persistence of virus infection and prolong therapeutic efficacy¹⁴²; the approach is now being evaluated in clinical trials.

Enhancing antitumor immunity—Immune evasion by tumors, recognized as one of the 'Hallmarks of Cancer'", represents an important target for new cancer therapeutics¹⁴³. Tumors produce immunosuppressive cytokines (e.g., TGFβ) and recruit immune inhibitory cells (e.g., T-regs), thereby paralyzing the antitumor immune response^{144, 145}. Oncolytic viruses may uniquely combine tumor debulking activity (via direct tumor lysis and/or vascular attack) with potent activation of adaptive and innate immune responses. Targeted infection of the tumor leads to a localized inflammatory response triggering an immune storm directly within the malignancy, facilitating immune recognition of cancer specific neo-antigens¹⁴⁵. As discussed in the earlier section on clinical trial results, work with the oncolytic vaccinia virus JX-594 and more recently with Amgen's oncolytic herpes virus, talimogene laherparepvec, both armed with GM-CSF, suggests clinical benefit can be attained when localized oncolytic activity is coupled with immune cell recruitment^{19, 145, 146}.

The idea of an oncolytic vaccine combining virus-mediated tumor destruction with immune recognition of tumor antigens is attractive but requires careful orchestration as the activated immune system may prematurely suppress therapeutic virus replication¹⁴⁷. Recent work suggests that even limited infection of distant lymph node metastases may lead to enhanced therapeutic benefit. Bridle and colleagues¹⁵⁹ have shown that an oncolytic rhabdovirus expressing a tumor antigen can robustly boost a primed anti-tumor immune response, but only if given systemically when the virus can access both the tumor and distant lymph tissues^{145, 148}. Recentlxzy, rather striking results have been observed in tumor-bearing animals 'vaccinated' with an oncolytic rhabdovirus expressing a complex library of cDNAs encoding normal cellular antigens¹⁴⁹. Although it is unlikely that a library of viruses expressing thousands of unique sequences could become a therapeutic product, this work does suggest that future oncolytics that express a small number of carefully selected tumor antigens should be tested in the clinical setting. Towards that goal, three VSVs encoding melanoma specific antigens that induce IL-17 recall responses were selected from a library of VSV-cDNA, that when used in combination but not alone were as efficacious as the

parental complete VSV-cDNA library¹⁵⁰. Several groups are now combining adoptive cell therapy with oncolytic viruses reasoning that virus-mediated tumor cell destruction should enhance the activity of the transferred cells^{145, 151}.

Which virus for which indication?-Given that naturally occurring viruses have such widely differing structures, lifecycles and tropisms, resulting in a diversity of highly distinctive clinical manifestations, it would seem logical that each oncolytic virus would be ideally suited to a specific malignancy. It is therefore somewhat surprising that there are to date very few examples of this specific matching of a given oncolytic virus with a specific class of malignancy, and most of the oncolytic viruses currently in development show a relatively broad spectrum of antitumor activity, typically against both epithelial and hematological malignancies. Certain oncolytic viruses were initially developed with the expectation that they would be better suited to a given broad class of malignancies, but this has subsequently proven not to be the case. Thus, oncolytic adenoviruses were considered better suited for therapy of epithelial malignancies but are now showing activity against hematologic cancers^{152, 153}; herpes simplex viruses were developed originally for brain cancer therapy but are now showing promise in a variety of non-central nervous system tumors, including sarcomas and epithelial malignancies^{153, 154}; and measles viruses were originally considered an ideal candidate for hematologic malignancies but have also proven to have broad spectrum activity against epithelial malignancies and sarcomas¹⁵⁵.

Obviously, where viruses are engineered to target specific cell surface receptors or nuclear transcription factors, their utility is thereafter limited to tumors that express the relevant target, but to date there has been a definite preference for clinical translation of oncolytic viruses with a broader spectrum antitumor activity. This may be a consequence of the safety concerns being more difficult to address for fully retargeted viruses that stringently target a single type of tumor. Although it may seem counterintuitive that a virus engineered to restrict its host range might have greater pathogenic potential than the parent virus, several examples indicate the association of loss of pathogenic potential (attenuation) with the broadening of virus host range^{156, 157}. Thus, assumptions about safety and host range cannot be trusted and have to be tested experimentally in appropriate animal species to directly address this question.

It is apparent from the above discussion that safety considerations are ever-present in preclinical oncolytic virus studies and may also be drivers of the choice of virus for a given indication. Different viruses have differing toxicities, and genetically manipulating these viruses may result in unexpected toxicities, such as an instance in which insertion of the IL-4 gene into a murine poxvirus resulted in 100% lethality in pre-vaccinated animals that were previously completely immune to the wild-type virus¹⁵⁸. Natural and engineered virus tropisms, virus mutability and capacity for evolution, immunomodulatory/antiapoptotic and cytotoxic gene products, virus transmissibility, prevalence of antiviral immunity in the population and availability of drugs or antisera to eliminate unwanted or persistent infections are all important factors to be considered in the safety analysis of oncolytic viruses that are candidates for clinical translation.

CONCLUSIONS

Oncolytic viruses are structurally and biologically diverse, spreading through tumors and killing tumor cells by multiple mechanisms and with different kinetics. Because of their large size and immunogenicity they are constrained by physical barriers and by host immunity, but they can also cross-prime and amplify antitumor immunity, serving as a cancer immunotherapy. Overall, the field has been slow to develop but recent clinical trial data has been promising and a first-in-class USA approval is expected soon for a recombinant herpes simplex virus being tested in a randomised phase III clinical trial that recently completed accrual. This virus (Talimogene Laherparepvec, previously OncoVEX) is administered by intralesional injection to patients with metastatic malignant melanoma and spreads locally, cross-priming the antimelanoma immune response, but does not spread systemically to distant sites of tumor growth. Thus the talimogene study is primarily exploiting the oncolytic virus as tumor-debulking immunotherapy and does not clinically validate the 'oncolytic paradigm' where systemic and intratumoral spread of the infection lead to tumor debulking as a prelude to immune-mediated eradication of minimal residual disease. However, a more direct validation of the oncolytic paradigm may soon come from ongoing clinical trials testing intravascular OV delivery in immunotherapy-resistant tumors using, for example, reovirus, vaccinia, and measles viruses. And for the future, there is a long and growing list of new or improved versions of oncolytic viruses that have been ingeniously selected, engineered and honed for systemic therapy, several of which will doubtless, in the fullness of time, join the growing arsenal of clinically approved anticancer drugs.

Looking beyond the expected clinical approval of oncolytic viruses as single agents, there is enormous scope for the development of more complex protocols to achieve superior treatment outcomes. Preclinical studies provide a very strong basis for this assertion, demonstrating numerous synergistic interactions that can overcome the various barriers constraining oncolytic viruses, such as the use of cell carriers to optimize virus delivery⁶⁰ or of immunosuppressive drugs¹⁵⁹ to enhance their intratumoral spread. One particularly interesting prospect is that new drugs will be developed capable of potently suppressing the innate immune responses of virus infected cells. Elucidation of the intracellular signaling pathways of innate immunity has been progressing very rapidly in recent years, so the stage is now set for this important area of drug discovery.

But as the field comes closer to its lofty goal of a single shot virotherapy cure for cancer¹⁶⁰, it is very likely that we will encounter significant treatment-related toxicities. The minimal toxicity in clinical trials to date is often cited as a strength of the oncolytic virotherapy approach, but in the absence of rapid destructive intratumoral virus spread, which is the ultimate goal, it is hardly surprising the treatment has seemed innocuous. With increased potency and more reliable efficacy, toxicity will surely follow, and hence the need for ever more stringent virus targeting technology to ensure that the destructive power of these exciting new drugs is focused exclusively on the tumor.

The most important technical challenges that continue to attract the attention of the oncolytic research community are the optimization/enhancement of systemic virus delivery,

intratumoral virus spread and cross-priming of anticancer immunity. However, the harmonization of solutions to these problems is perhaps a greater challenge still, although definitely achievable¹⁶⁰. Suppressing immunity may increase intratumoral spread, but diminishes cross-priming of the anticancer immune response. Conversely, enhancing immunity may improve cross-priming but the price paid is to limit intratumoral virus spread, the basis of oncolytic tumor debulking. Many of the 'solutions' that have been developed to date have been analyzed in artificial model systems that are not powered to reveal the positive and negative consequences of a given modification to all aspects of the overall treatment paradigm. This points to another major challenge for the field which is to develop better model systems that really do reliably mirror the human oncolytic virotherapy scenario. Mouse xenograft models lack a functional immune system, and immunocompetent mouse tumor models are frequently misleading because the viruses being tested behave differently in mice and humans. Thus, many oncolytic viruses cannot infect mouse cells so they lack activity in syngeneic mouse models, while others preferentially infect mouse vs. human cells so their anticancer activity (and toxicity) is not transferrable to human trials. The use of oncolytic agents such as vaccinia virus which are capable of infecting mouse and human cells with equal efficiency is a potential solution to this problem and may prove to be an important factor for the acceleration of their clinical development. In addition the development of transgenic mouse models susceptible to "human specific" viruses remains an important goal.

The biggest overall challenges facing the field at the current time have less to do with the development of new technology solutions for virus delivery and spread than with how to get them clinically tested. There are so many elegant solutions available for hypothetical problems that it can be demoralizing for scientists to see their engineering efforts lost in the morass. Clinical testing of each new virus modification is simply not realistic because of the enormous amount of work and expense required – manufacture, pharmacology/toxicology testing, protocol development and regulatory approval - to move each new product into phase I trials. Take for example oncolytic adenoviruses where a PubMed search shows 100 publications in the past 10 years on almost as many unique adenovirus configurations representing multiple serotypes (for antibody evasion) with or without engineered fiber modifications (for transductional targeting), hexon modifications (to eliminate hepatic sequestration), polymeric coats (for shielding), gene deletions (for physiological targeting), transgene insertions (to combat innate immunity, enhance adaptive immunity, promote spread, increase cytotoxicity or facilitate noninvasive monitoring). Every one of these modifications can be classified as a new product and the modifications are coming so fast that an oncolytic virus that was state of the art a few years ago may today be considered archaic even before it has completed phase II clinical testing.

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Figure 1.

Barriers to efficient oncolytic virus delivery via the bloodstream (virus neutralization by serum factors, sequestration by the mononuclear phagocytic system or lack of extravasation) and solutions to circumvent them.



Figure 2.

Factors constraining intratumoral virus spread (host innate or acquired immunity and extracellular matrix) and solutions to circumvent them.



Figure 3. A timeline of milestones in the development of oncolytic virotherapy to improve virus specificity, potency, delivery and spread

SPECIFICITY: (1) **Translational targeting**: 1991, Engineering of a replication-competent HSV attenuated for neurovirulence for glioma treatment¹⁰. (2) **Transcriptional targeting**: 1997, targeting of HSV using albumin promoter/enhancer for hepatoma cells¹⁶¹ and Ad using Prostate specific antigen (PSA) promoter for prostate cancer cells¹⁶², (3)

Transductional targeting: 2005, targeting entry and cytopathic effects of oncolytic measles virus by display of single-chain antibody on the virus attachment protein, (4) **MicroRNA targeting**: 2008, to control unwanted toxicity of picornarvirus¹³¹ and vesicular stomatitis virus¹³² while retaining antitumor activity, (5) **DNA shuffling**, 2008, Mixing a pool adenoviral serotypes and passaging the pools under conditions that invite recombination between serotypes to generate tumor selective virus¹⁶³.

POTENCY: (1) **Prodrug activation**: 1998, an oncolytic adenovirus expressing cytosine deaminase and HSV-Tk designed to work in combination with 5-FC and Ganciclovir, (2) **Proapoptotic genes**: 2000, introduction of the adenovirus death protein (ADP) into an oncolytic adenovirus to enhance its cytotoxicity¹⁶⁴, (3) Immune stimulation: 2001, oncolytic HSV encoding IL-12 and GM-CSF to recruit T lymphocyte-mediated antitumor immune response¹⁶⁵, (4) **Radioisotope**: 2004, an oncolytic measles virus encoding the human

sodium iodide symporter (NIS) which concentrates beta-emitting (radiovirotherapy) and gamma-emitting isotopes (imaging)³³, (5) 2006, **Matrix degrading proteins**: adenovirus encoding relaxin protein to enhance virus intratumoral spread¹⁶⁶, (6) **Shuffling:** 2008, Mixing a pool adenoviral serotypes and passaging the pools under conditions that invite recombination between serotypes to generate more potent adenovirus ColoAd1¹⁶³.

DELIVERY & SPREAD: (1) **Immune suppressive drugs**: 1999, Addition of cyclophosphamide to combat and innate and adaptive antiviral immunity to enhance intratumoral spread of HSV, (2) **Cell carriers**: 2006, use of cytokine induced killer cells to deliver oncolytic vaccinia virus to tumor, resulting in synergistic antitumor activity¹⁶⁷, (3) **Shielding**: 2008, Polymer coating and retargeting of oncolytic adenovirus for ovarian cancer to enhance viral pharmacokinetics⁴⁰, (4) **Infectious Nucleic Acid**: 2011, delivery of oncolytic picornarvirus using infectious nucleic acid (RNA) to successfully achieve sustained viremia and tumor regression.

CLINICAL TRIALS: (1) **Activity**: 2009, Phase II trial with intralesional injection of oncolytic HSV, OncoVEX (talimogene laherparepvec), in melanoma patients. 26% complete response (8 out of 50), with durability in both injected and uninjected lesions including visceral sites¹⁹. Undergoing Phase III evaluation. (2) **Viremic Threshold**: 2011, Intravenous delivery of JX-594, oncolytic vaccinia virus, in patients with metastatic tumor, demonstrating the need for a viremic threshold to be reached for efficient virus delivery to tumors²⁶.

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Table 1

A listing of current and recently completed oncolytic virotherapy trials. A search was performed on www.clinicaltrials.gov and the clinical trial database of the Journal of Gene Medicine (http:// www.wiley.com/legacy/wileychi/genmed/clinical/).

Virus	Name	Modifications	Phase	Tumor	Route	Combination	Site	Status* (PubMed reference)
Adenovirus	Oncorine (H101)	E1B-55k-E3-	Π	SCCHN	Ш	Cisplatin	Multileft	Completed, PMID: 14693057
			Ш	SCCHN	Ш	Cisplatin	Multileft	Completed, PMID: 15601557
	Onyx-015	E1B-55k-E3B-	Ι	Lung Mets	N		Mutlileft	Completed, PMID: 11420638
			Ι	Glioma	Intracavity		Mutlileft	Completed, PMID: 15509513
			Ι	Ovarian Ca	II		Mutlileft	Completed, PMID: 11896105
			I	SCCHN	IT		Multileft	Completed, PMID: 10741699
			Ι	Solid tumors	IV	Enbrel	Mary Crowley	Completed, PMID: 17704755
			Ι	Sarcoma	Ш	Mitomycin-C Dox, cisplatin	Mayo Clinic	Completed, PMID: 15647767
			II/I	PanCa	Ц	Gemzar	UCLA	Completed, PMID: 12576418
			Ш	CRC	IV		Mutlileft	Completed, PMID: 12697873
			П	Hepatobiliary	Ш		Montefiore	Completed, PMID: 12576437
			Ш	CRC, PanCa	IA		Mutlileft	Completed, PMID: 12414631
			Ш	SCCHN	Ш		Multileft	Completed, PMID: 11208818
			Ш	SCCHN	Ш	Cisplatin, 5-FU	Multileft	Completed, PMID: 10932224
			Ш	CRC	N	5-FU/leucovorin	Stanford	Completed, PMID: 15803147
	CG7060	PSA control	Ι	Prostate Ca	Ц	RT	Johns Hopkins	Completed, PMID: 11606381
	CG7870/CV787	rat probasin-E1A hPSA-E1B	II/I	Prostate Ca	N	ı	Multileft	Completed, PMID: 16690359
		H3+	II/I	Prostate Ca	N	Docetaxel	Mary Crowley	Terminated, 2005
	CG0070	E2F-1, GM-CSF	III/II	Bladder Ca	Intracavity	ı	UCSF	Not yet open, PMID: 16397056
	Telomelysin	hTERT	Ι	solid tumors	П	ı	Mary Crowley	Completed, PMID: 19935775
	Ad5-CD/TKrep	CD/TK	Ι	Prostate Ca	Ш	5-FC & GCV	Henry Ford, Detroit	Completed, PMID: 12208748
			Ι	Prostate Ca	Ц	5-FC+GCV+RT	Henry Ford, Detroit	Completed, PMID: 14612551
	Ad5-D24-RGD	RGD, Delta-24	Ι	Ovarian Ca	IP	ı	UAB	Completed, PMID: 20978148
			Ι	Glioma	Ц	ı	MD Andersen	Recruiting
			ΠЛ	Glioma	П	ı	Erasmus Medical left	Recruiting
	Ad5-SSTR/TK-RGD	SSTR, TK, RGD	I	Ovarian Ca	IP	GCV	UAB	Active, PMID: 16397056
	CGTG-102	Ad5/3, GM-CSF Delta-24	II/I	Solid tumors	Ш	ı	Baylor	Not open, PMID: 20664527

Virus	Name	Modifications	Phase	Tumor	Route	Combination	Site	Status* (PubMed reference)
			I	Solid tumors	IT/IV	Metronomic CTX	Docrates Hospital Helsinki	Recruiting
	INGN-007 (VRX-007)	wtE1a, ADP	Ι	Solid tumors	TI		Mary Crowley	Not open, PMID: 19197324
	ColoAd1	Ad3/11p	II/I	CRC, HCC		'	PsiOxus	Not open, PMID: 18560559
Coxsackie Virus (CVA21)	CAVATAK		Ι	Melanoma	Ш		Viralytics	Completed
			II	Melanoma	TI	,	Viralytics	Recruiting
			Ι	SCCHN	Ш		Viralytics	Terminated
			Ι	Solid tumors	N		Viralytics	Recruiting
Herpes Simplex Virus	Talimogene Laherparepvec (OncoVEX)	GM-CSF	Ι	Solid tumors	Ш	1	Multileft	Completed, PMID: 17121894
		ICP34.5(-)	Ш	Melanoma	Ш	ı	Multileft	Completed, PMID: 19915919, 19884534
		ICP47(-)	III	Melanoma	Ш		Multileft	Active
		Us11↑	II/I	SCCHN	Ш	RT, Cisplatin	Multileft	Completed, PMID: 20670951
	G207	ICP34.5(-), ICP6(-) LacZ(+)	II/I	Glioma	П	I	U of Alabama	Completed, PMID: 18957964, 10845725
		I	Glioma	IT	RT	U of Alabama	Completed	_
	G47Delta	From G207, ICP47-	Ι	Glioma	Ц	ı	Tokyo Hospital	Recruiting, PMID: 11353831
	HSV 1716 (Seprehvir)	ICP34.5(-)	Ι	Non-CNS solid tumors	ŢŢ	ı	Cincinnati	Recruiting
		Ι	SCCHN	IT	ı	U of Glasgow	Completed, PMID: 18615711	
		Ι	Glioma	IT	ı	U of Glasgow	Completed, PMID: 15334111, 11960316	
		Ι	Melanoma	IT	ı	U of Glasgow	11229673, 2001	
		Ι	Mesothelioma	IP	ı	UK	not active	
	HF10	HSV-1 HF strain	Ι	Solid tumors	Ш	ı	Multileft	Recruiting
		Ι	Pancreatic Ca	IT	ı	Nagoya University	Completed, PMID: 21102422	
		Ι	Breast Ca	IT	ı	Nagoya University	Completed, PMID: 16865590	
		Ι	SCCHN	IT	ī	Nagoya University	Completed, PMID: 16923721	
	NV1020		Ι	CRC liver mets	IA	-	MSKCC	Completed, PMID: 19018254
Measles Virus (Edmonston)	MV-CEA	CEA	I	Ovarian Ca	Ш	'	Mayo Clinic	Completed, PMID:20103634
			Ι	Glioma	П	ı	Mayo Clinic	Recruiting
	SIN-VM	NIS	Ι	Myeloma	N	CTX	Mayo Clinic	Recruiting
			I	Ovarian Ca	II	,	Mayo Clinic	Recruiting

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Virus	Name	Modifications	Phase	Tumor	Route	Combination	Site	Status* (PubMed reference)
			III	Mesothelioma SCCHN	ПР		U of Minnesota/Mayo Clinic Mayo Clinic	Recruiting Not open
Newcastle Disease Virus	NDV-HUJ PV701 MTH-68/H NV1020			Glioma Solid tumors Solid tumors Solid tummors	IV IV Inhalation IV		Goldyne Savad Inst Ottawa Hospital UCRI Multileft	Completed, PMID: 16257582 Completed, PMID: 16638865 Completed, PMID: 8275514 Completed, PMID: 11980996
Parvovirus	H-1PV	-	II/I	Glioma	IT/IV		University Hospital Heidelberg	Recruiting, PMID: 20299703
Poliovirus (Sabin)	PVS-RIPO	IRES	I	Glioma	п	ı	Duke	Recruiting, PMID: 20299272
Reovirus (Dearing)	Reolysin	,	ПЛ	Glioma	П		Multileft	Completed, PMID: 18253152
			Ι	Peritoneal Ca	Ъ		Ohio State	Recruiting
			I ·	Solid tumors	2	-	Multileft	Completed, PMID: 18981012
			II	Solid tumors CRC	2 2	CIX FOLFIRI	Multileft Multileft	Recruiting Recruiting
			Π	Sarcoma	N		Multileft	Completed
			Π	Melanoma	N		Multileft	Suspended
			Π	Ovarian, Peritoneal Ca	N	PTX	Mutlileft	Recruiting
			Π	Pancreatic Ca	N	PTX, CBDCA	Multileft	Recruiting
			Π	SCCHN	N	PTX, CBDCA	Multileft	Not recruiting
			Π	Melanoma	N	PTX, CBDCA	U of Texas	Recruiting
			Π	Pancreatic Ca	IV	Gemzar	U of Texas	Recruiting
			Π	Lung Ca	N	PTX, CBDCA	Multileft	Recruiting
			Ш	SCCHN	IV	PTX, CBDCA	Multileft	Recruiting
Seneca Valley Virus	NTX-010		п	Small Cell Lung Ca	IV	ı	NCCTG multileft	Recruiting, PMID: 17971529
Retrovirus	Toca 511	CD	II/I	Glioma	ш	5-FC	Multileft	Recruiting, PMID: 16257382
Vaccinia (Wyeth strain)	JX-594	GM-CSF TK(-)	I	CRC	IV	ı	South Korea	Recruiting
			I	Solid tumors	N		Multileft	Completed
			I	HCC	IT		Busan, South Korea	Completed, PMID: 18495536

Virus	Name	Modifications	Phase	Tumor	Route	Combination	Site	Status* (PubMed reference)
			Ι	Pediatric solid tumors	П		Cincinnati	Recruiting
			Ι	Melanoma	IT		Busan, South Korea	Completed, PMID: 21772252
			II/I	Melanoma	П	I	Multileft	Completed. PMID: 10505851
			Π	HCC	IT	ı	Multileft	Not recruiting, data analysis
			IIB	HCC	N		Multileft	Recruiting
			II/I	CRC	IV/IT	Irinotecan	Multileft	Recruiting
			Π	CRC	IT		Ottawa Hospital	Not yet recruiting
Vaccinia (Western Reserve)	vvDD-CDSR	TK-, VGF- LacZ, CD Somatostatin R	Ι	Solid tumors	VI/TI	I	U of Pittsburgh	Recruiting PMID: 15336655
Vaccinia (Lister)	GL-ONC1 (GLV-h68)	Renilla Luciferase GFP, β-gal	Ι	Solid tumors	N		Royal Marsden	Recruiting, PMID: 21779374
		b-glucoronidase	II/I	Peritoneal Carcinomatosis	IP		University Hospital Tuebingen	Recruiting
			II/I	SCCHN	IV	RT,Cisplatin	Moores UCSD Cancer left	Recruiting
Vesicular Stomatitis Virus (Indiana)	VSV-hIFNβ	IFN beta	Ι	нсс	П	ı	Mayo Clinic	Recruiting

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