

Therapeutic Cancer Vaccines: Current Status and Moving Forward

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Concurrent with US Food and Drug Administration (FDA) approval of the first therapeutic cancer vaccine, a wide spectrum of other cancer vaccine platforms that target a diverse range of tumor-associated antigens is currently being evaluated in randomized phase II and phase III trials. The profound influence of the tumor microenvironment and other immunosuppressive entities, however, can limit the effectiveness of these vaccines. Numerous strategies are currently being evaluated both preclinically and clinically to counteract these immunosuppressive entities, including the combined use of vaccines with immune checkpoint inhibitors, certain chemotherapeutics, small-molecule targeted therapies, and radiation. The potential influence of the appropriate patient population and clinical trial endpoint in vaccine therapy studies is discussed, as well as the potential importance of biomarkers in future directions of this field.

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This article reviews the current state of the design, development, and clinical applications of therapeutic cancer vaccines. The numerous previous failures of cancer vaccines may be attributed to the fact that not all vaccines are created equal (there has been a clear evolution in vaccine design) and/or to issues in clinical trial design and patient population. This article covers the diversity of therapeutic cancer vaccine platforms and the wide range of tumor-associated antigens being targeted; the current state of clinical trials that test therapeutic cancer vaccines, including the results of recent randomized multicenter phase II and phase III studies; issues to be considered in future clinical trials; the profound influence of the tumor microenvironment and immunosuppressive factors in inhibiting optimal vaccine efficacy and potential strategies to combat these phenomena; combinatorial therapies, including the use of cancer vaccines with immune stimulants, inhibitors of immune suppression, certain chemotherapeutic agents, small-molecule targeted therapies, hormones, and radiation; the potential use of biomarkers to select those patients who are most likely to benefit from vaccine therapy; and other future directions in this field. This article is not designed to comprehensively review all preclinical and clinical studies of therapeutic cancer vaccine, but instead, to delineate the factors and concepts that are currently being evaluated to develop vaccines as therapeutics for a wide range of human cancers and cancer stages, either as monotherapies or as combination therapies.

Spectrum of Current Therapeutic Cancer Vaccine Platforms

Many diverse vaccine platforms have now been evaluated in phase II and/or phase III clinical trials, including injection of peptides or

proteins in adjuvant, injection of recombinant viruses or other recombinant microorganisms, delivery of killed tumor cells, or delivery of protein- or peptide-activated dendritic cells (DCs) to the patient (Table 1). Each of the 14 platforms in Table 1 has strengths and weaknesses that can be influenced by the particular tumor-associated antigen (TAA) that is targeted, the disease and disease stage that is targeted, the clinical trial endpoint, and whether the vaccine is evaluated in combination with an immune stimulant, an inhibitor of immune suppression, or another mode of cancer therapy.

DCs are the most potent antigen-presenting cells (APCs) (43). Numerous phase II studies have now evaluated the use of peptide- or protein-pulsed, or viral vector-infected, DCs to treat patients with prostate cancer, colorectal cancer, melanoma, glioma, and other cancers (Tables 1 and 2) [eg, references (44,53)]. The Sipuleucel-T vaccine (41), which was recently approved by the US Food and Drug Administration (FDA) for the therapy of asymptomatic metastatic castrate-resistant prostate cancer (mCRPC), consists of APCs from peripheral blood mononuclear cells (PBMCs) that have been incubated with prostatic acid phosphatase (PAP, a prostate antigen) fused to granulocyte macrophage colony-stimulating factor (GM-CSF). This vaccine regimen consists of leukaphereses to purify PBMCs from the patient and processing in a central facility where the PAP fusion protein is added to the APCs; these cells are then reinfused to the patient for the purpose of conferring immunity; this process is repeated three times at biweekly intervals. One drawback of DC and/or APC platforms is that they require leukapheresis(es) and cell culture processing of PBMCs, and thus a limited number of vaccinations can be used.

Vaccines based on peptides from TAAs, which are usually administered in an adjuvant and/or with an immune modulator,

Table 1. Spectrum of current vaccine platforms in phase II/III clinical studies*

Vaccine platform	Example	Cancer type	Selected references
Peptides/proteins			
Peptide	gp100 (modified), MUC-1 (Stimuvax), HER2/neu	Melanoma, lung	(1–9)
Protein	MAGE-A3, NY-ESO	Melanoma	(10)
Antibody	Anti-idiotype	Lymphoma	(11–14)
Glycoproteins	sTn-KLH	Melanoma	(15,16)
Recombinant vectors			
Poxvirus	rV, rF-PSA-TRICOM (Prostvac)	Prostate	(17–29)
<i>Saccharomyces cerevisiae</i> (yeast)	yeast-ras	Pancreatic	(30,31)
<i>Listeria</i>	Listeria-mesothelin	Pancreatic	(32)
Alpha- and adenoviruses	adeno-CEA, alpha-CEA	Carcinoma	(33)
Tumor Cells			
Autologous	adeno-CD40L, colon (BCG)	CLL, colon, melanoma	(34–36)
Dendritic cell/autologous tumor cell fusions		Myeloma	(37)
Allogeneic	GVAX (+GM-CSF)	Pancreatic	(38–40)
Dendritic cells/APCs			
APC–protein	Sipuleucel-T (PAP-GM-CSF)	Prostate	(41,42)
Dendritic cell–peptide	Glioma peptides	Glioma, melanoma	(43–45)
Dendritic cells–vector infected	rV, rF-CEA-MUC1-TRICOM (Panvac-DC)	Colorectal	(46,47)

* APC = antigen-presenting cell; BCG = Bacillus Calmette–Guerin adjuvant; CD40L = CD40 ligand; CEA = carcinoembryonic antigen; CLL = chronic lymphocytic leukemia; gp100 = glycoprotein 100; GM-CSF = granulocyte macrophage colony-stimulating factor; MAGE-A3 = melanoma-associated antigen 3; MUC-1 = mucin 1; NY-ESO = New York esophageal carcinoma antigen 1; PAP = prostatic acid phosphatase; PSA = prostate-specific antigen; rF = recombinant fowlpox; rV = recombinant vaccinia; sTn-KLH = sialyl-Tn–keyhole limpet hemocyanin.

are generally cost-effective and have the advantage that the investigator knows exactly which epitope to evaluate in terms of patients' immune responses (1). However, they also have a potential drawback because they target only one epitope or a few epitopes of the TAA. It is generally believed that for a cancer vaccine to be optimally efficacious, it must induce antigen-specific CD8+ cytolytic

Table 2. Spectrum of current and potential therapeutic cancer vaccine targets*

Target type	Examples	Selected references
Oncoprotein	point mutated: ras, B-raf, frame shift mutations, undefined unique tumor mutations; HER2/neu, MUC-1 C-terminus, p53	(1,7,8,48,49)
Oncofetal antigen	CEA, MUC-1	(2–4,19,26)
Cancer–testes	MAGE-A3, BAGE, SEREX-defined, NY-ESO	(10,50–52)
Tissue lineage	PAP, PSA, gp100, tyrosinase, glioma antigen	(5,6,24,25,27,41,44,53)
Stem cell/EMT	Brachyury, SOX-2, OCT-4, TERT, CD44 ^{high} /CD24 ^{lo} , CD133 ⁺	(54–62)
Viral	HPV, HCV	(63,64)
Glycopeptides	sTn-KLH	(15,16)
Antiangiogenic	VEGF-R	(65,66,67)
B-cell lymphoma	Anti-id	(11–14)

* BAGE = B melanoma antigen; CEA = carcinoembryonic antigen; EMT = epithelial–mesenchymal transition; gp100 = glycoprotein 100; HCV = hepatitis C virus; HPV = human papillomavirus; MAGE-A3 = melanoma-associated antigen-3; MUC-1 = mucin 1; NY-ESO = New York esophageal carcinoma antigen 1; OCT-4 = octamer-binding transcription factor 4; PAP = prostatic acid phosphatase; PSA = prostate-specific antigen; SOX-2 = (sex determining region Y)-box-2; sTn-KLH = sialyl-Tn–keyhole limpet hemocyanin; TERT = telomerase reverse transcriptase; VEGF-R = vascular endothelial growth factor receptor.

T cells (CTLs), which are responsible for tumor cell lysis, and antigen-specific CD4+ “helper” T cells, which provide cytokines to enhance CTL activity. Some polypeptide vaccines potentially contain both CD4 and CD8 epitopes; for example, Stimuvax contains both kinds of epitopes for mucin 1 (MUC-1), which is found as a cell surface proteoglycan associated with several tumor types (2–4). Protein-based vaccines are more costly than peptide-based vaccines, but they usually also contain both CD4 and CD8 epitopes. Many peptide- and protein-based vaccines are used as part of a DC vaccine platform.

Anti-idiotype vaccines are directed against specific antibodies found on the surface of B-lymphoma cells (11–13). They have the advantage of targeting a unique tumor-specific antigen. A disadvantage is that their generation and production are quite labor intensive in that, to date, each anti-idiotype vaccine has been patient specific. However, some researchers have shown that these patient-specific vaccines can be produced in less than a month [reviewed in (12)].

The most evaluated viral-based vectors are from the poxviridae family. They include vaccinia, modified vaccinia strain Ankara (MVA), and the avipoxviruses (fowlpox and canarypox; ALVAC). Poxviruses have the ability to accept large inserts of foreign DNA, and thus can accommodate multiple genes. Intracellular expression of the transgene(s) allows for processing of the tumor antigen by both the class I and class II major histocompatibility complex (MHC) pathways (17). Because poxvirus replication and transcription are restricted to the cytoplasm, there is minimal risk to the patient (or host) of insertional mutagenesis. It has been shown in preclinical studies that when the transgene for a TAA is inserted in vaccinia or MVA, it becomes more immunogenic, most likely because of the Toll-like receptors (TLRs) and other properties of the virus that induce a local inflammatory response. This same property of the non-avian poxviruses, however, leads to virus neutralization by the host immune response and limits their use to one, or at most two, vaccination(s). Recombinant avipoxviruses can

be used multiple times; they will induce antiviral immune responses, but they are not neutralizing because their “late” viral coat proteins are not produced in mammalian cells (18–20).

Other viruses can also be used as vectors for TAAs. Alphaviruses such as Venezuelan Equine Encephalitis (VEE) virus are attractive vectors because, once they have infected the host, they replicate RNA in the cytoplasm and express high levels of a transgene (33). Recombinant adenovirus vectors are easy to engineer and have shown utility as vaccines and gene therapy agents (34), but clinical evaluation has been hindered by high levels of preexisting antiviral immunity. Newer variants of adenoviruses are being developed and evaluated that may potentially be less immunogenic.

Bacteria and yeasts have shown some promise as vaccine vectors in preclinical studies and may also serve as vectors for immunizing cancer patients. Heat-killed recombinant *Saccharomyces cerevisiae* is inherently nonpathogenic, can be easily propagated and purified, and is very stable. Recombinant yeasts have been shown to activate maturation of human DCs and to present both class I and class II epitopes of transgenes (30). Surprisingly, it appears that these vectors can be administered multiple times without eliciting host-neutralizing activity (31). Attenuated recombinant *Listeria monocytogenes* (Lm) bacteria have also been shown to target DCs, and, like viral and yeast vectors, they stimulate both innate and adaptive immune responses (32).

Although DNA vaccine platforms have shown promise in preclinical studies (65,66), early clinical trials have been disappointing. Their exact mode of action is not known at this time. However, new constructs and methods of administration may enhance their utility.

The use of whole tumor–cell vaccines has the advantage of presenting the patient’s immune system with a range of both known and undefined TAAs as immunogens. However, this same property also potentially diminishes the relative level of expression of a particular TAA or group of TAAs and its presentation and processing by APCs. The use of a killed whole tumor–cell vaccine is usually accompanied by an immune stimulant such as GM-CSF, Bacillus Calmette–Guerin adjuvant (BCG), or CD40 ligand (CD40L).

Autologous tumor cell vaccines have a great advantage because they present the unique set of TAAs, such as particular point mutations or fusion gene products, from a given patient’s own tumor (35). Because this technology depends on the availability of tumor biopsies, it is feasible for only some tumor types and stages. In one variation of this technique, DCs and autologous tumor cells are fused together before immunization of the patient (37). DC–tumor cell fusions combine the unique properties of whole tumor–cell vaccines with the enhanced antigen-presenting power of DCs.

Alternatively, allogeneic whole tumor–cell vaccines, which typically contain two or three established and characterized human tumor cell lines of a given tumor type, may be used to overcome many logistical limitations of autologous tumor–cell vaccines. The GVAX vaccine platforms (38,39,40), which contain allogeneic pancreatic, prostate, or breast tumor cells, are a testament to the ability to provide such a vaccine for multicenter evaluation.

Spectrum of Vaccine Targets

The validity of a vaccine target will depend on the ability of a tumor cell to process the TAA in the context of MHC for T-cell

or for B-cell recognition, the level of expression, the relative specificity of the TAA for tumor vs normal adult tissue, and the degree of inherent “tolerance” to the given TAA (68,69). Common targets include oncoproteins, oncofetal antigens, differentiation-associated proteins, and viral proteins, among others (Table 2). The ideal target is a somatic point mutation that helps to drive the neoplastic process. Clinical trials are underway to test vaccines that target the relatively few *RAS* mutations found in colorectal and pancreatic cancer. However, large numbers of tumor-associated mutations among the various exons of *TP53* make generating the large number of possible mutant p53 vaccines somewhat prohibitive. Similarly, it is difficult to target the wide array of frameshift mutations and unique mutations that occur in individual tumors. Non-mutated oncoproteins that have served as vaccine targets include overexpressed HER2/neu (ERBB2), p53, and the carboxy terminus of MUC-1 (48,49).

Numerous cancer vaccine trials have targeted a class of antigens categorized as oncofetal antigens, such as carcinoembryonic antigen (CEA), underglycosylated MUC-1, tumor-associated glycopeptides (15,16,19,68), and “cancer–testes” antigens defined by serological expression cloning (SEREX) immunodetection such as melanoma-associated antigen (MAGE-A3) and B melanoma antigen (BAGE) (10,50–52,70). These antigens are overexpressed in many tumor types and to a lesser extent in some normal adult tissues. Numerous trials have targeted “tissue lineage” antigens that are overexpressed in tumors and normally expressed in a non-vital organ, such as PAP, prostate-specific antigen (PSA), and the melanoma-associated antigens glycoprotein 100 (gp100) and tyrosinase. The recent FDA approval of the Gardasil vaccine targeting the human papillomavirus (HPV) (63,64,67) for the prevention of cervical cancer renders HPV an attractive target for cervical cancer therapy, as does targeting the hepatitis C virus for liver cancer therapy.

The most provocative potential targets for vaccine therapy are molecules that are associated with cancer “stem cells” and/or the epithelial–mesenchymal transition (EMT) process, both of which are associated with drug resistance. Drivers of EMT are also associated with tumor cell extravasation and intravasation to the metastatic site. Recent studies have described the plasticity of so-called “cancer stem cells” and the similarities between cells undergoing EMT and the acquisition of “stem-like” characteristics (54).

EMT and cancer “stemness” are associated with the proteins (sex determining region Y)-box-2 (SOX-2) and octamer-binding transcription factor 4 (OCT-4) and with carcinoma cells that are CD44^{high} and CD24^{low} and/or are CD133⁺ (55–59). Each of these gene products is currently being evaluated for immunogenicity in terms of generating human T-cell responses in vitro, but some of them also have a relatively broad range of expression in some normal adult tissues. The T-box transcription factor Brachyury has recently been identified as a major driver of EMT (60). It has been shown to be selectively expressed on both primary and metastatic lesions of several carcinoma types. T-cell epitopes have been identified on the Brachyury protein that have the ability to generate human T cells capable of lysing a range of human carcinoma cells (61). Vaccines are currently being developed to target gene products associated with EMT and cancer cells with stem-like characteristics. Preclinical studies have also shown the potential of vaccines

that target molecules involved in tumor angiogenesis, such as the vascular endothelial growth factor receptor (VEGF-R) (65,66).

A potent means to enhance immunogenicity of a self-antigen is by altering specific amino acids of TTAs to develop enhancer agonist epitopes, which are designed to enhance binding either to the MHC or to the T-cell receptor, resulting in higher levels of T-cell responses and/or higher avidity T cells (71–73). For example, the gp100 melanoma vaccine contains an enhancer agonist epitope, the PROSTVAC vaccine contains a PSA enhancer agonist epitope, and the PANVAC vaccine contains enhancer agonist epitopes for both CEA and MUC-1.

Current State of Cancer Vaccine Clinical Trials

Many vaccines were initially tested in patients with metastatic disease who had undergone multiple previous therapies. Clinical studies have now shown that patients will generally respond better to vaccines when they have been treated with fewer previous chemotherapeutic regimens and when a longer time has elapsed since their last chemotherapy (21,22). In many other forms of cancer therapy that kill tumor cells when ample doses of the therapeutic agent are delivered, the limiting factor is usually host toxicity. By contrast, cancer vaccines have demonstrated minimal toxicity, but their success is limited by the number of host-induced effector cells induced by the vaccine vs immunosuppressive entities. Consequently, patients with large tumor burdens would be even less likely to respond to vaccines than to chemotherapy.

Prostate Cancer as a Prototype for Vaccine Therapy

Although the vast majority of previous and ongoing vaccine trials have been conducted in patients with metastatic melanoma, several characteristics render prostate cancer a prototype disease for the evaluation of therapeutic cancer vaccines (23). First, time is often required to generate a sufficient immune response to curtail tumor growth, and prostate cancer is generally an indolent disease that may not lead to metastatic disease or death for over a decade or more. Second, prostate cancer cells express a variety of well-characterized TAAs. Third, the serum marker PSA can be used to identify patients with minimal tumor burden and those responding to therapy. Last, a well-defined nomogram, the Halabi nomogram (74), can be used at presentation of metastatic disease to predict a patient's probable response to standard of care chemotherapy and/or hormone therapy.

After early clinical trials of the Sipuleucel-T vaccine demonstrated its safety, larger trials were conducted in patients with minimally symptomatic mCRPC. Although a pair of small phase III trials failed to meet their primary endpoint of improved disease progression, there was evidence that Sipuleucel-T prolonged survival in mCRPC (42). A larger phase III trial (41) was then conducted in which more than 500 patients were enrolled and overall survival (OS) was set as the primary endpoint. In this trial, again, no change in time to cancer progression was seen; however, OS was improved in the vaccine arm (25.8 vs 21.7 months; $P = .032$; Figure 1, A). In April 2010, the FDA approved Sipuleucel-T for the treatment of minimal or non-symptomatic mCRPC.

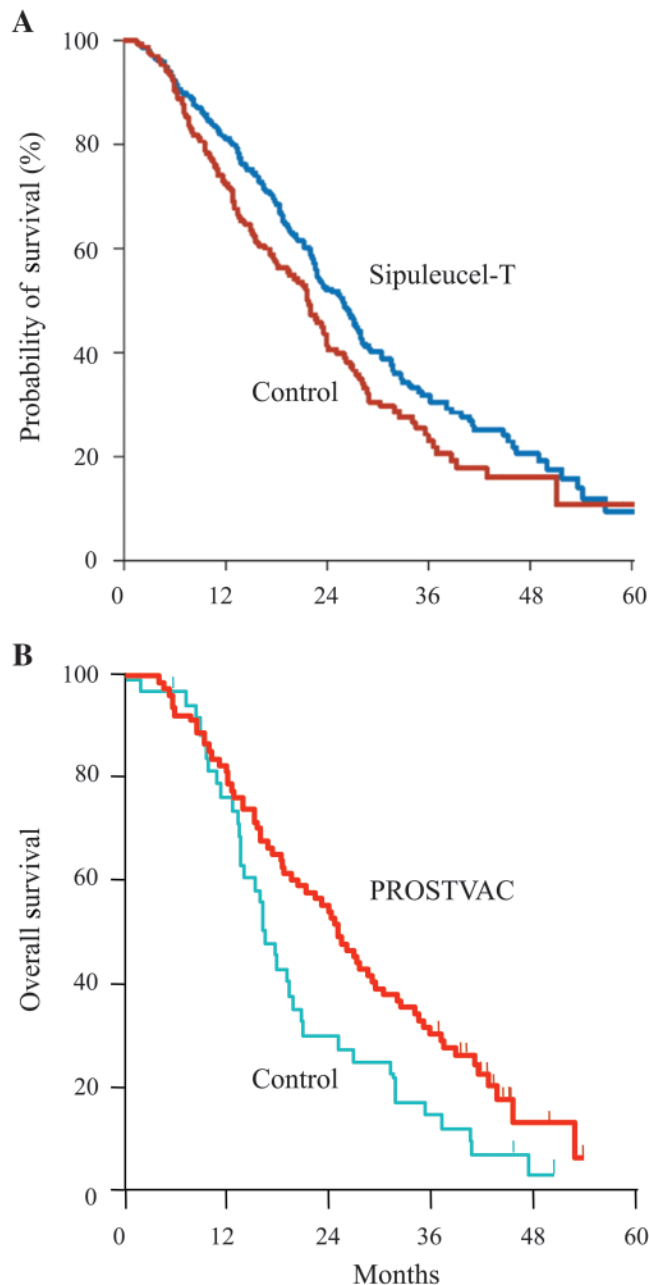


Figure 1. Overall survival (OS) in patients with metastatic castrate-resistant prostate cancer using cancer vaccines. **A)** Results of the primary efficacy analysis of treatment with Sipuleucel-T as compared with placebo control. Sipuleucel-T did improve patients' OS (hazard ratio for death = 0.78; 95% confidence interval = 0.61 to 0.98; $P = .03$) [adapted with permission from reference (41)]. The placebo control consisted of cultured antigen-presenting cells (APCs) from leukapheresis, without prostatic acid phosphatase-granulocyte macrophage colony-stimulating factor (PAP-GM-CSF) antigen. Per the trial protocol, the control group could receive cryopreserved APCs with antigen upon disease progression. **B)** OS of a randomized, placebo-controlled 43-center trial of PROSTVAC vaccine consisting of recombinant vaccinia and fowlpox vectors containing transgenes for prostate-specific antigen, B7.1, intercellular adhesion molecule 1, and lymphocyte function-associated antigen-3 (PSA-TRICOM) in patients with metastatic castrate-resistant prostate cancer; the trial compared PROSTVAC vaccine vs empty vector. There was an OS advantage of 8.5 months (OS = 25.1 vs 16.6 months; $P = .006$) and a 44% reduction in death in the vaccine arm [adapted with permission from reference (24)].

A second prostate cancer vaccine has also been evaluated in the same population of men with mCRPC. This “off-the-shelf” platform (PROSTVAC) consists of a recombinant vaccinia virus priming vaccination and multiple fowlpox booster vaccinations. Each vector contains transgenes for PSA and three costimulatory molecules (CD80, intercellular adhesion molecule 1, and lymphocyte function–associated antigen-3 that are collectively designated TRICOM) (72,75). A 43-center randomized placebo-controlled phase II trial enrolled 125 minimally symptomatic mCRPC patients (24). Similar to the experience with Sipuleucel-T, treatment with PROSTVAC did not alter time to progression; however, it improved median OS relative to a placebo, the control vector (OS = 25.1 vs 16.6 months, $P = .006$; Figure 1, B). Over the course of follow-up, 44% fewer patients who received PROSTVAC died compared with the control cohort (hazard ratio [HR] = 0.56) (24). The median OS in a second PROSTVAC single-arm phase II study (25) was 26.6 months, which was similar to that observed in the PROSTVAC arm of the larger PROSTVAC randomized trial.

Another point to consider in the treatment of mCRPC and other cancers is the patients’ quality of life. The Sipuleucel-T and PROSTVAC vaccines compared favorably with the chemotherapeutic agents and hormonal therapies that are FDA approved for mCRPC, especially because there were fewer serious adverse events seen with these vaccines than with the other agents (Table 3).

Phase III Studies

In addition to Sipuleucel-T, two different vaccine platforms have also recently demonstrated positive results in phase III trials. One phase III trial (5) compared patients with locally advanced stage III or IV melanoma who were treated with a modified gp100 peptide vaccine in Montanide ISA-51 adjuvant plus standard high-dose interleukin 2 (IL-2) vs such patients who were treated with high-dose IL-2 alone (5). The serious adverse events in both groups were consistent with IL-2 therapy. The group that was treated with the gp100 peptide vaccine plus IL-2 demonstrated a statistically significant improvement in objective clinical response rate compared with the group that was treated with IL-2 alone (16% vs 6%, $P = .03$). Patients treated with the peptide vaccine plus IL-2 had a slightly longer progression-free survival (2.2 vs 1.6 months; $P = .008$). The median OS was also longer in the vaccine plus IL-2 group than in the IL-2-only group (OS = 17.8 vs 11.1 months; $P = .06$). This was

the first phase III trial to demonstrate a clinical benefit for a peptide vaccine in melanoma. The findings in this trial were not observed in three previous independent phase II clinical trials that were reported together elsewhere (6). Those studies did not detect a benefit for the combination treatment, although the clinical trials were not powered to assess a progression-free endpoint, unlike the larger phase III trial.

A phase III study that used an anti-idiotypic vaccine in patients with follicular lymphoma has also shown evidence of patient benefit (11,12). In this study, patients were required to have previously untreated advanced-stage follicular lymphoma; they were treated on-trial with standard chemotherapy before vaccination. Those patients who achieved a complete remission were randomly assigned to receive Id–keyhole limpet hemocyanin (KLH) plus GM-CSF vs unconjugated KLH plus GM-CSF as a control. After a median follow-up of 56.6 months, the median time to relapse for the Id-KLH plus GM-CSF arm was 44.2 months vs 30.6 months for the KLH plus GM-CSF control arm, suggesting a benefit associated with use of this vaccine ($P = .045$; HR = 1.6). These results were in contrast to the negative results observed in two other phase III studies that have used recombinant idiotypic vaccine for follicular lymphoma (12,14). One of several key differences in the positive study compared with the two negative phase III trials of related idiotypic vaccine products was the low tumor burden required for all vaccinated patients on the positive study, although other issues may have contributed to survival outcomes (11–13).

Other vaccines have shown mixed results in phase III trials. For example, a phase III trial (35) using autologous tumor cells plus BCG showed a survival advantage in patients with Dukes stage B2-C3 colon cancer (33.3% deaths in the vaccine group compared with 56.5% in the control group; $P = .039$) and was ineffective in Dukes C and D patients.

There are several ongoing phase III trials employing different vaccine platforms in a range of human cancers. These include the use of Stimuvax, a vaccine targeting liposomal MUC-1 peptide, in non–small cell lung cancer (NSCLC) (2,76); Lucanix, an allogeneic whole tumor–cell vaccine containing a transforming growth factor-beta 2 (TGF- β 2) antisense transgene, in NSCLC (77); and vaccination with the MAGE-A3 protein, in stage III melanoma and in NSCLC (78). The PROSTVAC phase III trial was initiated in 2011 in patients with mCRPC. There are also

Table 3. Overall survival in patients with metastatic castrate-resistant prostate cancer who were treated with vaccines or other therapeutic modalities*

Agent	Type of therapy	Stop treatment AE, %	Improvement in median OS, mo	Hazard ratio	Reduction in death rate, %
Docetaxel	Chemotherapy	11	2.4	0.76	24
Cabazitaxel	Chemotherapy	18	2.4	0.70	30
Abiraterone	Hormone	19	3.9	0.66	34
Sipuleucel-T	Vaccine	1.5	4.1	0.78	22
PROSTVAC†	Vaccine	~2	8.5	0.56	44

* AE = adverse event; OS = overall survival; PROSTVAC = a recombinant vaccinia virus priming vaccination and multiple fowlpox booster vaccinations. Each vector contains transgenes for prostate-specific antigen and three costimulatory molecules (CD80, intercellular adhesion molecule 1, and lymphocyte function–associated antigen 3) that are collectively designated TRICOM.

† Results of a 43-center randomized placebo-controlled phase II trial.

numerous ongoing randomized phase II studies that employ most of the vaccine platforms noted in Table 1.

It is emphasized that in virtually all of the therapeutic cancer vaccine trials reported to date, there have been extremely low levels of toxicity—mostly limited to grade I and II levels. Moreover, although many of the target TAAs are also expressed at some levels in selected normal adult tissues, virtually no evidence of autoimmunity has been observed, with the exception of vitiligo resulting from the administration of some melanoma vaccines (36,45,79).

Issues in Vaccine Clinical Trial Design

Recent randomized phase II and III vaccine trials have revealed two major issues to be considered in clinical trial design: appropriate patient population and clinical trial endpoint. As described above, patients who have received fewer regimens of chemotherapy and who have had a longer time since their last chemotherapy appear to generally respond better to vaccines. There are now several examples of greater vaccine efficacy in patients with low grade or more indolent disease.

A clinical trial employing the PANVAC vaccine platform (ie, sequential administration of recombinant vaccinia and recombinant fowlpox vectors delivering the TAAs CEA, MUC-1, and TRICOM) in colorectal cancer patients showed some evidence of patient benefit (26), but no evidence of clinical benefit in patients with large volume liver metastases. These findings are in contrast to recent results from a trial to evaluate the same vaccine in colorectal cancer patients following metastasectomy (surgical removal) of liver or lung metastases. In this multicenter trial (46,47), 74 patients who had no evidence of disease after resection and completion of their physician-determined perioperative chemotherapy were vaccinated with PANVAC (ie, with vaccine alone or with vaccine-modified DCs). Data from a prospectively registered, comparable contemporary control group of colorectal cancer patients who had undergone metastasectomy were also available (46,47). The 2-year relapse-free survival was similar in all groups: 50% for the DC-PANVAC group, 56% for the PANVAC group,

and 55% for the contemporary control group. However, the 2-year OS was 95% for the vaccinated group and 75% for the contemporary control group; after approximately 40 months of follow-up, 67 of 74 (90%) of the vaccinated patients survived vs approximately 47% OS in the contemporary control group; the data for 3- to 5-year survival data of colorectal cancer patients after metastasectomy in five other trials ranges from 28% to 58% (80–85). A randomized trial is necessary to confirm these results. It is of interest, however, that this is yet another example of a vaccine trial that shows little or no evidence of an improvement in relapse-free survival, yet has an apparent benefit in OS (69).

Another example of the advantage of the use of vaccines in patients with more indolent disease is seen in the use of the Halabi nomogram in patients presenting with mCRPC (74). This nomogram was developed to predict survival using data from more than 1100 mCRPC patients who were treated with chemotherapy and hormone therapy. When this patient population was treated on trial with docetaxel, the standard of care, the nomogram accurately predicted survival both for patients predicted to live less than 18 months after therapy was initiated and for those who were predicted to live longer than 18 months (Table 4) (25). In a contemporary trial at the same institution in which the same patient population was treated with the PROSTVAC vaccine, the Halabi nomogram accurately predicted survival for those patients with a Halabi-predicted survival (HPS) of less than 18 months. However, the actual survival for patients with more indolent disease far exceeded that predicted [HPS of 20.9 months vs actual survival >37.3 months; see reference (25) and Table 4].

In patients who are treated with traditional cytotoxic agents, it is widely believed that improved time to progression is a prerequisite for improved OS. A recent study (86) evaluated tumor regression and growth rates in four chemotherapy trials and one vaccine trial in patients with mCRPC. Figure 2 illustrates the growth rate constants defined in that study. Cytotoxic agents affect the tumor only during the period of administration; soon after the drug is discontinued, because of drug resistance or toxicity, antitumor activity ceases and the growth rate of the tumor increases (Figure 2, A, line b). With vaccine therapy, the mechanism of action and kinetics

Table 4. Predicted survival of patients with metastatic castrate-resistant prostate cancer using the Halabi nomogram vs actual survival*

	Docetaxel therapy (n = 22)			PROSTVAC (n = 32)		
	All patients	Patients with HPS <18 mo	Patients with HPS ≥18 mo	All patients	Patients with HPS <18 mo	Patients with HPS ≥18 mo
Predicted survival by Halabi score, mo	16.5	13.0	21.0	17.4	12.3	20.9
Actual median overall survival, mo	15.5	15.4	16.9	26.6	14.6	Not reached (8 of 15 patients alive at 37.3 mo)
Difference, mo	-1.0	2.4	-4.1	9.2	2.3	≥16.4
Patients surviving longer than predicted by Halabi nomogram	11 (50%) of 22	8 (62%) of 13	3 (33%) of 9	22 (69%) of 32	10 (59%) of 17	12 (80%) of 15†

* HPS = Halabi nomogram–predicted survival; PROSTVAC (also known as PSA-TRICOM) = prostate-specific antigen, CD80, intercellular adhesion molecule 1 and lymphocyte function-associated antigen 3. Adapted from (25) with permission.

† P = .035.

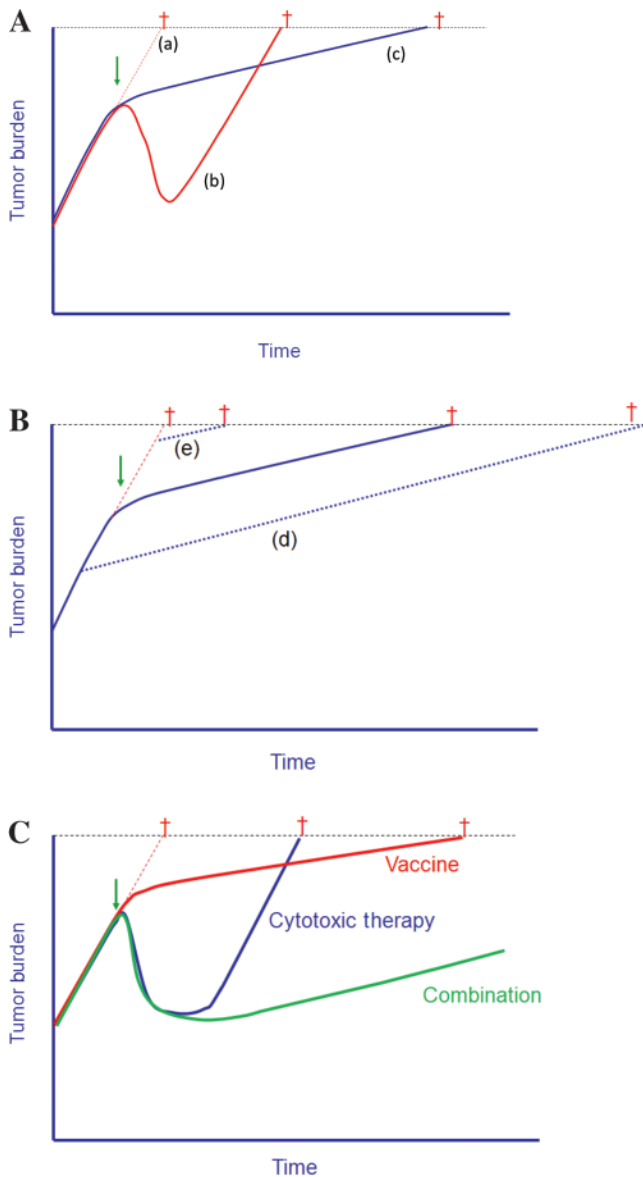


Figure 2. Tumor growth rates following chemotherapy vs vaccine therapy [adapted from data in references (86–88)]. **A)** Average tumor growth rates and time to death in patients with metastatic prostate cancer, from five clinical trials [four with chemotherapy and one with PROSTVAC vaccine (also known as PSA-TRICOM)]. Growth rate of tumor if no therapy is initiated (**line a**). With the use of chemotherapy, there was an initial tumor reduction, but the growth rate of tumors at relapse was similar to the initial tumor growth rate before therapy (**line b**). With PROSTVAC, there was a reduction in tumor growth rate following vaccine therapy (**line c**). Thus, for patients who received vaccine therapy with little if any tumor reduction (among whom there was virtually no increase in time to progression), an increase in overall survival was observed. **Dagger** denotes time to death. **B)** This phenomenon could potentially be enhanced if vaccine therapy is initiated earlier in disease progression or in patients with low tumor burden metastatic disease (**line d**) but would have minimal effect in patients with large tumor burden (**line e**). **C)** Predictions of enhanced overall survival if patients are treated with both vaccination and chemotherapy.

of clinical response appear to be very different (86). Therapeutic vaccines do not directly target the tumor; rather, they target the immune system. Immune responses often take time to develop and can potentially be enhanced by continued booster vaccinations; any

resulting tumor–cell lysis can lead to cross-priming of additional TAAs, thus broadening the immune repertoire (a phenomenon known as antigen cascade or epitope spreading). This broader, and perhaps more relevant, immune response may also take some time to develop. Although a vaccine may not induce any substantial reduction in tumor burden, vaccines as monotherapy have the potential to apply antitumor activity over a long period, resulting in a slower tumor growth rate (Figure 2, A, line c). This deceleration in growth rate may continue for months or years and, more importantly, through subsequent therapies. This process can thus lead to clinically significant improved OS, often with little or no difference in time to progression and a low rate of, or lack of, objective response [Figure 2, A, line c; reference (69)]. Thus, treating patients with a vaccine when they have a lower tumor burden, as compared with a greater tumor burden, may result in far better outcomes (Figure 2, B, line d vs line e). It is hypothesized that the combined use of vaccine and cytotoxic therapy (Figure 2, C) may result in both tumor regression (via the cytotoxic therapy) and reduced tumor growth rate (via vaccine therapy) (86,87,88). These concepts will be discussed below. Thus, early clinical trials with vaccine may have been terminated prematurely with the observation of tumor progression before sufficient vaccine boosts could be administered. This phenomenon has actually led to modifications in how vaccine clinical trials are now designed and to “new immune response criteria” for immunotherapy (89).

In some ways, our progress has been paralyzed by the insistence by some on the paradigm that only after tumor lesions are reduced using classic Response Evaluation Criteria in Solid Tumors (RECIST criteria) (90) can a therapeutic modality be considered beneficial. Indeed, randomized clinical studies with small-molecule targeted therapeutics, as well as with different immunotherapeutic agents, have sometimes demonstrated increased survival with minimal, if any, reduction in tumor burden or time to progression. In addition to the Sipuleucel-T and PROSTVAC vaccine trials described above, a phase III trial leading to FDA approval, in which patients with metastatic melanoma were treated with ipilimumab (an anti-cytotoxic T-lymphocyte antigen-4 monoclonal antibody), demonstrated a statistically significant advantage in survival without a statistically significant difference in time to progression (91). Patients with unresectable stage III or IV melanoma were randomly assigned to receive ipilimumab plus gp100 peptide, ipilimumab alone, or gp100 alone. The median OS was 10.0 months among patients receiving ipilimumab plus gp100, as compared with 6.4 months among patients receiving gp100 alone (HR for death = 0.68; $P < .001$). The median progression-free survival was 2.76 months in the ipilimumab plus gp100 group, 2.86 months in the ipilimumab-alone group, and 2.76 months in the gp100-alone group.

Influence of the Tumor Microenvironment and Immunosuppressive Factors

One of the major reasons for the limited success of therapeutic cancer vaccines to date is likely to be the negative influence of the tumor microenvironment and other immunosuppressive factors (92–95). Preclinical studies have shown that the interstitial pressure within a large tumor mass diminishes diffusion of macromolecules,

such as antibodies, and effector cells, such as T cells (96,97). Most solid tumors also lack T-cell costimulatory molecules. Thus, when activated T cells, especially those of relatively low avidity directed against self-antigens, bind to tumors lacking costimulatory molecules, they are anergized and lose lytic capacity. Similarly, it has been shown in preclinical models of chronic viral infection that T cells chronically exposed to viral antigen can become exhausted (98,99). The inhibitory co-receptor programmed death 1 (PD-1) has been shown to be present on such exhausted T cells (100).

The tumor microenvironment has been shown to contain a range of immunosuppressive immune cell types including CD4⁺ regulatory T cells (Tregs), myeloid derived suppressor cells (MDSCs), suppressor CD8⁺ T cells, tumor-associated macrophages (TAMs), and regulatory natural killer (NK)/NKT cells (101–109). Analysis of PBMCs from patients with several types of cancer has also shown increased levels of MDSCs and Tregs, as well as an increased suppressive function of Tregs on effector T cells (101,110–112). These suppressive cells, tumor cells, and other cells in the tumor microenvironment can also release into the microenvironment a number of soluble immunosuppressive factors, including TGF- β , IL-10, indoleamine-pyrrole 2,3 dioxygenase (IDO), and VEGF (106,113,114). Strategies to combat these immunosuppressive entities with combination therapies are described below. The tumor microenvironment can also influence the phenotype of tumor cells. For example, it has been shown (62) that IL-8 can drive carcinoma cells to the EMT phenotype and toward more stem-like characteristics.

As with any phenotypic analysis of cell types within a tumor mass, many tumors display antigenic heterogeneity, which is reflected both at the level of antigen expression and at the level of the antigen–peptide MHC complex. Tumor cells within masses have also been shown to have defects in MHC–peptide complexing machinery (115–119). Analysis of cloned populations of carcinoma cells has shown that this antigenic heterogeneity is an extremely “plastic” property, in which cloned cells will often revert to express a more heterogeneous antigenic population upon passage. In preclinical models, a tumor expressing CEA in only some of its cells can be cured with a CEA-based vaccine (20). CEA must be present in the vaccine and in some tumor cells for cures; however, analysis of the tumor masses in regressing tumors showed the presence of higher levels and greater avidity of T cells to other TAAs (120). When cured animals were subsequently challenged with CEA-negative tumor cells, they would also reject these tumors (120). This antigen cascade phenomenon has also been seen in clinical trials in which vaccinated patients develop higher levels of T-cell responses to TAAs compared with responses seen to the antigen found in the vaccine (1,7,121–124). This phenomenon is reported in some trials to be associated with patient benefit (7,121,124).

Immune Combination Therapies

Preclinical studies have clearly demonstrated that vaccine combined with a wide range of immune stimulants, or inhibitors of immune suppression, will greatly enhance antitumor responses. A major problem in the field of vaccine therapy, however, is the availability

of many of these other agents. A National Cancer Institute workshop in July 2007 listed 12 such agents (125,126), and most are still not available for use in vaccine combination. Many of these agents were evaluated as monotherapies to maximally tolerated dose in phase I trials and were found to have little or no activity and subsequently abandoned. Preclinical studies have demonstrated that when some of these agents are used far below the maximally tolerated dose with vaccine, and/or at lower doses locally with vaccine, they can greatly enhance vaccine efficacy (127,128).

Potential immune stimulants include cytokines such as IL-2, IL-15, IL-7, GM-CSF, and interferon (IFN) (129–140). IL-2 has the disadvantage of enhancing Tregs along with effector cells (131,141). IL-15, now in phase I, has the advantage of enhancing effector T-cell response and not Tregs (130,135,136). GM-CSF has been used successfully in the GVAX and other vaccine platforms (39,40,142). However, the use of GM-CSF is a dual-edged sword in which higher levels may be immunosuppressive (25). IFNs have been shown to enhance immune responses and to enhance the expression of TAAs and MHC on tumor cells (143–145). Antibody–cytokine fusions, such as hu14.18-IL2, have shown promise in patients with refractory neuroblastoma (146) and other similar immunocytokines are currently in development. A range of TLRs has been shown to enhance vaccine efficacy in preclinical studies, but most are being evaluated clinically as monotherapies. Classic adjuvants such as incomplete Freund’s, ISCOMs, Detox, and chitosan provide both a depot effect for the protein or peptide vaccine (or cytokine) and a level of immune stimulation in regional lymph nodes (147–150). Clinical trials will need to carefully evaluate both the doses and scheduling of all of these immune stimulants with vaccine.

Perhaps, the most promising area of immune vaccine combination therapy is that of immune checkpoint inhibitors and inhibitors of immunosuppressive entities. Ipilimumab has been shown in several preclinical models to enhance the avidity of T cells and to enhance antitumor effects in combination with vaccines such as CEA-TRICOM recombinant poxviruses (91,151–153). A recent clinical trial of the PROSTVAC vaccine in combination with ipilimumab showed evidence of enhanced survival in patients with mCRPC compared with what had been seen in previous trials with PSA-TRICOM as monotherapy (154); a randomized trial is thus warranted. The recent FDA approval of ipilimumab for metastatic melanoma (91) will undoubtedly make vaccine combination therapies with this agent more feasible with a range of vaccine platforms. Ongoing clinical studies with anti-PD-1 and anti-PD-1 ligand 1 monoclonal antibodies as monotherapies are also yielding promising results (155–159). These agents should also enhance clinical vaccine efficacy.

Other agents that will reduce or eliminate immune suppressive factors on cells are also being evaluated. Monoclonal anti-CD25-diphtherial toxin (Ontak) has been shown to reduce Tregs and to enhance vaccine efficacy (160,161). Both monoclonal and small-molecule inhibitors to TGF- β are also currently in clinical evaluation as monotherapies, but their full potential may be realized in vaccine combination therapy (162–165).

Very few cancer chemotherapy or small-molecule targeted therapy regimens employ one agent—and at times, they may use three or more agents. Each of the diverse vaccine platforms

described in Table 1 can potentially activate the host adaptive and innate immune response differentially, and the same or different TAAs may be targeted (see Table 2), making the use of combination vaccine therapy quite an attractive possibility. A recent pre-clinical study (166) that used two diverse recombinant CEA vaccine platforms (poxviral and yeast) in mice demonstrated that each vector processed TAA epitopes differentially and activated a different T-cell repertoire and cytokine profile, resulting in enhanced antitumor activity. Evidence for the potential of combination vaccine therapies is also seen in the preclinical and clinical results using diversified prime-boost strategies with vaccinia and fowlpox vectors (18,27), and the only acquired immune deficiency syndrome (AIDS) prevention vaccine to show some evidence of promising results used a diversified recombinant avipox prime-protein boost regimen (167).

Vaccines in Combination with Other Therapeutic Modalities

It was commonly assumed that chemotherapy and vaccines do not mix. However, preclinical studies have demonstrated, and early clinical evidence is emerging, that multiple forms of therapy can be used concurrently with cancer vaccines or after vaccine therapy, with additive or synergistic effects. Some of these phenomena are summarized in Table 5.

Certain chemotherapeutic agents can enhance vaccine-mediated T-cell killing by several distinct mechanisms. Oxaliplatin and anthracyclines such as doxorubicin will induce what has been termed “immunogenic tumor cell death,” which results in enhanced cross-priming of TAAs by DCs and subsequent activation of T cells (168–171). Agents such as docetaxel have been shown to suppress MDSCs and to increase the expression of TAAs, peptide-MHC complexes, adhesion molecules, and death receptors such as FAS on the surface of tumor cells and thus render them more susceptible to vaccine-induced T-cell killing (172,173); this same phenomenon has been observed when tumor cells have been exposed to external beam radiation, radiolabeled monoclonal antibodies, and bone-seeking radionuclides (174–176). Clinical trials

that have used PROSTVAC vaccine plus the taxane docetaxel in patients with prostate cancer (28) have been completed, and others are in progress that are testing PANVAC vaccine in combination with docetaxel in patients with metastatic breast cancer (177). A trial that uses PROSTVAC vaccine in combination with a bone-seeking chelated radionuclide in patients with prostate cancer metastatic to the bone is also in progress (178). Low-dose paclitaxel (another taxane) has also been shown to enhance DC function in preclinical studies (179).

As a consequence of the homeostatic proliferation of immune cells following certain chemotherapeutic regimens such as cisplatin with vinorelbine, different immune subsets have been shown to differentially recover, leading to enhanced effector to regulatory T-cell ratios (180). Clinical studies with cyclophosphamide (used at relatively low doses) have been shown to diminish the number of Tregs (40). Certain chemotherapeutic agents, such as lenalidomide, have been termed “immunostimulatory,” and have been shown to stimulate T-cell proliferation and to enhance IL-2 and IFN- γ (181,182).

Preclinical studies have now demonstrated that certain small-molecule targeted therapeutics have the ability to enhance vaccine-mediated T-cell lysis of tumors. Both a BCL-2 inhibitor (183) and the tyrosine kinase inhibitor sunitinib (184–186) have been shown to enhance the ratio of TAA-specific T cells to regulatory cells, resulting in enhanced vaccine efficacy. A B-raf inhibitor has been shown to enhance the expression of TAAs on melanoma cells containing a *BRAF* mutation, resulting in enhanced T-cell lysis of tumor in vitro (187). The mTOR inhibitor, rapamycin, has been shown to enhance IL-12 production and the generation of memory CD8+ T cells (188,189). Vaccines are also now being used with monoclonal antibodies such as the HER2/neu inhibitor trastuzumab, resulting in enhanced HER2/neu-specific immune responses in breast cancer patients (1).

Hormonal therapy that is used in the treatment of several different stages of prostate cancer has been shown to induce thymic regeneration and the induction of naive T cells (190–192); it is at this interval that vaccine therapy may be most effective (29,193). A clinical study has demonstrated increased levels of infiltrating T cells in prostate cancer biopsies post- (vs pre-) hormonal therapy (194). Randomized clinical trials that employ PROSTVAC vaccine in combination with either nilutamide or flutamide hormone therapy in patients with non-metastatic prostate cancer have been completed and are ongoing (195,196).

In all of the above vaccine combination therapies, the dose and scheduling of the combining agent, the appropriate patient population, and the clinical endpoint are very important. There are many anecdotal reports and several publications that imply that patients who have received vaccine therapy, and whose tumors have nevertheless progressed, often undergo unexpected clinical responses when subsequent therapies are administered (28,197,198). This phenomenon, however, has not been validated prospectively or in a randomized trial. The Eastern Cooperative Oncology Group has recently initiated a multicenter randomized clinical trial in patients with metastatic prostate cancer to prospectively evaluate this phenomenon (199). Patients will receive either docetaxel or 2 months treatment with the PROSTVAC vaccine followed by docetaxel; OS will be the primary endpoint.

Table 5. Vaccines in combination with other therapeutic modalities*

Modality	Mechanism of action to enhance vaccine efficacy
Chemotherapy	“Immunologic” tumor cell death Alterations in tumor cell phenotype Enriched effector:regulator cell ratios
Radiation	Alteration in tumor cell phenotype
Small-molecule targeted therapeutics	Alterations in tumor cell phenotype Enriched effector:regulator cell ratios
Hormonal therapy	Thymic regeneration and induction of naive T cells
Monoclonal antibodies	Enhanced ADCC
Imids† (lenalidomide)	Stimulate T-cell proliferation

* ADCC = antibody-dependent cell-mediated cytotoxicity.

† Imids are a novel class of immunomodulators.

Biomarkers

The most common biomarker used in vaccine therapy trials has been the immune response of patients to TAAs post- (vs pre-) vaccination. Most trials have analyzed antibody responses to TAAs and/or analyzed PBMCs for CD8⁺ and/or CD4⁺ responses to the TAA in the vaccine using enzyme-linked immunosorbent spot- or fluorescence-activated cell sorting-based assays for cytokine production, or by binding of a peptide tetramer complex to the surface of T cells. Preclinical studies have shown, however, that the level of cytokine production by a T cell is not always associated with its lytic capacity. Because of the limited availability of samples, however, few studies have actually been able to measure the lytic capacity of T cells. Unfortunately, tumor biopsy specimens, the more appropriate site to obtain TAA-specific T cells, are unavailable in many trials.

More recently, a more comprehensive analysis of immune cell subsets from PBMCs has occurred in some studies including, in addition to T-cell responses, analyses of Tregs, MDSCs, NK, DCs, and antibodies to TAAs (9,200). Ratios of effector to regulatory cells have also been analyzed (25). Numerous studies have used analyses of multiple serum cytokines and chemokines.

What have emerged from these bioassay studies to date, at times, are associations between clinical outcome and an immune assay(s); however, these results are far from having identified any one assay as a “surrogate” for clinical responses. Potential reasons for this may be that PBMC analyses may not reflect which immune cells are actually at the tumor site; few studies have actually analyzed the “antigen cascade” phenomenon, where the true biomarker may be a T-cell population directed against a TAA not in the vaccine but generated via cross-priming; the diversity of the immune responses among individuals may not allow any one marker or set of markers as a surrogate for clinical response; and only recently has survival benefit emerged as a prominent endpoint in many vaccine studies, and adequate samples may not yet be available for comprehensive correlative analyses with survival.

Future Directions

Ongoing and future studies with therapeutic cancer vaccines will involve combination therapies with a range of therapeutic modalities as detailed here, including combinations of vaccines. Clinical trial designs will involve patients with more indolent metastatic disease and treatment in the adjuvant or neoadjuvant settings. There will also be less reliance on strict RECIST criteria, with more trials employing the new “immune response criteria” (89) and survival as endpoints when vaccine is used as a monotherapy.

As the long-term safety profiles of therapeutic vaccines are established, they most probably will be used in patients with a high risk of cancer, such as in patients with high-grade prostatic intraepithelial neoplasia who are at risk for the development of prostate cancer and familial adenomatous polyposis patients who are at risk for the development of colorectal cancer. Celecoxib has been used for the treatment of familial adenomatous polyposis (201), and preclinical studies have demonstrated that antitumor effects are greatly enhanced when CEA-TRICOM vaccine is combined with celecoxib (202).

Numerous studies have demonstrated that analysis of the immune infiltrate in colorectal and other cancer biopsies before chemotherapy can serve as a very strong independent prognostic indicator for survival (203–209). This phenomenon has yet to be exploited to define which patients may respond best to vaccine therapy. Analyses of PBMC immune cell subsets for detailed HLA typing may also vbetter reveal which patients are most likely to respond. Finally, more detailed analyses of mutations and unique tumor gene products, along with a further delineation of cancer cell stemness and the EMT phenotype, may also reveal new targets for vaccine therapy.

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