



# Vaccine Therapies for Cancer: Then and Now

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## Abstract

There are strong biologic and preclinical rationales for the development of therapeutic cancer vaccines; however, the clinical translation of this treatment strategy has been challenging. It is now understood that many previous clinical trials of cancer vaccines used target antigens or vaccine designs that inherently lacked sufficient immunogenicity to induce clinical responses. Despite the historical track record, breakthrough advances in cancer immunobiology and vaccine technologies have supported continued interest in therapeutic cancer vaccinations, with the hope that next-generation vaccine strategies will enable patients with cancer to develop long-lasting anti-tumor immunity. There has been substantial progress identifying antigens and vaccine vectors that lead to strong and broad T cell responses, tailoring vaccine designs to achieve optimal antigen presentation, and finding combination partners employing complementary mechanisms of action (e.g., checkpoint inhibitors) to overcome the diverse methods cancer cells use to evade and suppress the immune system. Results from randomized, phase 3 studies testing therapeutic cancer vaccines based on these advances are eagerly awaited. Here, we summarize the successes and failures in the clinical development of cancer vaccines, address how this historical experience and advances in science and technology have shaped efforts to improve vaccines, and offer a clinical perspective on the future role of vaccine therapies for cancer.

## Key Points

Clinical translation of vaccine therapies for cancer has been challenging due to the complexity of cancer immunology and optimal vaccine design.

Advances in vaccine technology and understanding of cancer immunology support continued investigation of vaccine-based treatment strategies for cancer.

## 1 Introduction

Successful cancer immunotherapy ultimately requires tumor cell engagement by cytolytic effectors (T cells and antibodies) capable of specifically recognizing unique or aberrantly expressed tumor-associated antigens (TAAs) or tumor-specific antigens (TSAs). While some patients with cancer spontaneously generate sufficient levels or function of antigen-specific T cells with the potential to generate impressive anti-tumor activity, the majority do not. One approach to ensure an adequate level and function of immune effectors is through therapeutic cancer vaccination. This form of active immunotherapy aims to generate anti-tumor immune responses directed against TAAs or TSAs [1, 2]. The idea of vaccination against cancer has a long history and was initially built on the observation that some tumors spontaneously regress in patients experiencing an acute infection [3]. More than a century ago, Dr. William Coley leveraged this observation to develop a rudimentary anti-cancer immune therapy consisting of heat-inactivated bacteria [3, 4]. How a non-specific innate immune response against bacterial products could translate into a specific anti-tumor immune response was explained subsequently by the discovery that antigen-presenting cells (APCs) (dendritic

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cells [DCs]) could acquire immunogenic tumor-derived peptides released during the innate immune response. These peptides could then be used to activate anti-tumor T cells with cognate receptors [5]. This led to the hypothesis that use of tumor-derived antigens, if delivered to the immune system in a sufficiently immunogenic context (a “vaccine”), would, due to the preferential targeting of cancer cells, enable relatively safe and yet effective treatments for cancer, capable of inducing long-lasting immunity [6].

Despite this encouraging foundation, and although cancer vaccines have been the subject of intense preclinical and clinical investigation in a variety of malignancies over the past 40 years, the successful clinical translation from bench to bedside has been slow, with only two therapeutic cancer vaccines (sipuleucel-T and talimogene laherparepvec [T-VEC]) having gained regulatory approval in the United States or European Union and numerous negative phase 3 studies leading to product discontinuations. However, the interest in therapeutic cancer vaccination remains high for several reasons. First, the clinical efficacy of checkpoint inhibitors and the identification of tumor-antigen-specific T cells in treated patients now provide evidence that patients are able to prime tumor-reactive T cells and that this likely occurs spontaneously in the minority of cancer patients responding to checkpoint blockade monotherapy. Second, the identification of checkpoint-expressing T cells and checkpoint ligand-expressing tumor cells after cancer vaccine therapies suggests that combination therapies incorporating vaccines and checkpoint inhibitors may be effective, as demonstrated in preclinical studies [7–10]. Third, negative studies have provided lessons for the field moving forward, which are being applied in current trials and will be also used in future investigations. The main lessons, as recently reviewed by Hollingsworth and Jansen (2019), include the need for antigens and vaccine designs that elicit greater immunogenicity (particularly through optimal presentation of tumor antigens by professional DCs [6]) as well as combination treatment strategies to overcome multiple mechanisms of tumor-mediated immunosuppression [11]. Fourth, a deeper understanding of major histocompatibility complex (MHC)-antigen binding has evolved to allow for better vaccine design and selection of appropriate antigens. Fifth, the ability to preferentially induce type 1 anti-tumor immunity versus the more common type 2 tumor supportive immunity has increased. Finally, although limited efficacy has been observed with the therapeutic cancer vaccine sipuleucel-T, its approval provided clinical validation of the therapeutic vaccination concept, which remains scientifically sound.

Given how recent advances may transform the track record of cancer vaccines, there is a need to summarize these developments and how they will affect the future role of vaccines. This review describes the successes and failures in the clinical development of cancer vaccines, addresses how this historical experience and advances in science and

technology have shaped efforts to improve vaccines (e.g., through optimizing antigen presentation by professional APCs), and offers a clinical perspective on the future role of vaccine therapies for cancer.

## 2 Historical Overview of Cancer Vaccines

Several types of cancer vaccines have been developed that vary depending on the form of the delivered antigen used in the vaccine: proteins or synthetic peptides of cancer antigens, cell-based delivery of tumor antigen (e.g., modified tumor cells, DCs loaded with tumor antigens), and DNA/RNA coding for cancer antigens (e.g., plasmids, RNA, viral vectors) (Fig. 1).

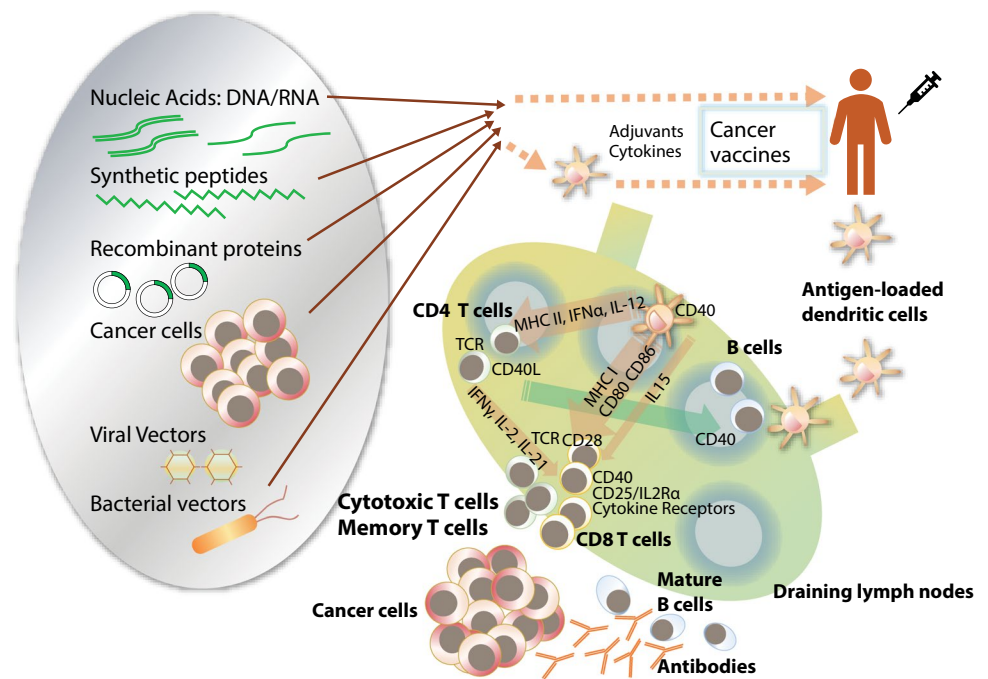
### 2.1 Peptide- and Protein-Based Vaccines

Peptide-based vaccines are relatively easy to manufacture, but combination with potent immune adjuvants is often needed to boost immunogenicity, and the number of people who may benefit from a given peptide vaccine is restricted by human leukocyte antigen (HLA) haplotype [13]. Several phase 3 studies investigating early peptide-based vaccines have not demonstrated clinical benefit despite demonstrating some induction of immune responses against TAAs or TSAs (Table 1) [11]. Explanations for lack of clinical benefit may lie in the properties of the peptides and adjuvants used, and early peptide vaccines may have been inherently inadequate for promoting antigen presentation and generating potent and durable anti-tumor immunity [6, 60–62].

A limitation of many early peptide vaccines was the use of short peptides (< 15 amino acids), including the minimal-length epitopes required to target cytotoxic lymphocytes (CTLs) but not T helper cells [6]. Short peptide epitopes are loaded onto non-professional APCs, including T cells and B cells [6, 63]. Yet, non-professional APCs circulate to non-inflamed lymphoid organs and do not deliver costimulatory signals to optimally prime and activate CTLs, thereby promoting tolerization [63]. Furthermore, cross-presentation of short peptides by professional APCs (DCs) is not as efficient or long lasting as that for synthetic long peptides [64]. Unfortunately, vaccines based on whole proteins (including idio-type vaccines) have also been largely unsuccessful in the clinic (Table 1). This may be due to the fact that the processing and presentation of whole proteins by DCs is inferior when compared with that for shorter peptides [65].

Overall, these results have provided rationale for the development of improved peptides such as synthetic long peptides with optimized immunogenicity, alternative peptide-delivery platforms such as nanoparticles, and more potent vaccine adjuvants.

**Fig. 1** Diverse therapeutic cancer vaccine platforms have a common mechanism of action [12]. [Figure reproduced from Maeng H et al. *F1000Res*. 2019 <https://doi.org/10.12688/f1000research.18693.1>. Licensed under CC BY 4.0.] CD cluster of differentiation, *IFN* interferon, *IL* interleukin, *IL2R $\alpha$*  IL-2 receptor alpha, *MHC* major histocompatibility complex, *TCR* T cell receptor



## 2.2 Cellular Vaccines

Commonly studied types of cell-based cancer vaccines include DCs loaded with tumor (neo)antigens, modified autologous cancer cells, and allogeneic tumor cell lines. Cell-based vaccines were among the initial types of therapeutic cancer vaccines tested. The first therapeutic cancer vaccine approved by the United States Food and Drug Administration was sipuleucel-T, a vaccine consisting of autologous peripheral blood mononuclear cells, including DCs, loaded with the prostatic acid phosphatase antigen fused with granulocyte-macrophage colony-stimulating factor (GM-CSF; an immune-cell activator). The approval of sipuleucel-T in 2010 for metastatic castration-resistant prostate cancer was based on results from the phase 3 IMPACT trial (NCT00065442) showing that treatment with sipuleucel-T significantly improved overall survival (OS) compared with placebo (median 25.8 vs. 21.7 months; hazard ratio [HR] 0.78;  $p = 0.03$ ; Table 2) [66]. These results and subsequent approval provided an early clinical validation of the therapeutic cancer vaccine concept. Real-world analyses suggest that sipuleucel-T remains an effective treatment option in the current treatment landscape, which includes androgen-receptor signaling pathway inhibitors (ASPIs). A retrospective cohort analysis of men with metastatic castration-resistant prostate cancer ( $N = 6044$ ; January 2013–December 2017) found that treatment with sipuleucel-T as first-line therapy or any-line therapy was associated with improved OS compared with treatment with ASPIs alone [68].

In contrast, cellular vaccines based on autologous tumor cells have not had the same success in several pivotal trials, as they either did not meet their primary endpoints or were

discontinued early because of clinical futility (Table 1). One possible explanation for this lack of success is the presence of immunosuppressive factors in the irradiated tumor cells or tumor cell lysates used for these vaccines [69, 70].

## 2.3 Genetic Vaccines

Viruses or plasmids can act as vectors for DNA or RNA encoding TAAs [11–13]. Viruses represent a promising platform for vaccines, as virus DNA or RNA may activate DCs by triggering pattern recognition receptors [11, 71].

As monotherapy, virus vector vaccines have not yet demonstrated consistent clinical benefit as demonstrated with the experiences with PROSTVAC and PANVAC (Table 1). For example, although a virus vector vaccine (PROSTVAC-VF) demonstrated OS benefit (but not progression-free survival [PFS; primary endpoint] or response) in a randomized phase 2 study of patients with metastatic castration-resistant prostate cancer [72], this positive signal was not validated in a subsequent phase 3 study (Table 1) [11, 16]. The investigators on the phase 3 study speculated that either PROSTVAC-VF did not generate sufficient immune responses as a single agent (possibly due to the choice of antigen or disease setting) or immunity was hampered by an immunosuppressive microenvironment [16]. To address these considerations, clinical trials of PROSTVAC-VF in combination with checkpoint inhibitors [11] and/or other cancer vaccines are ongoing (NCT02933255, NCT04020094, NCT03532217, and NCT03315871). However, a recent randomized phase 2 study found that addition of a viral

**Table 1** Select unsuccessful pivotal trials for therapeutic cancer vaccines

Vaccine platform type	Autologous or allogeneic	Product/com-pound name	Antigen(s)	Antigen ranking <sup>a</sup>	Identifier (phase, name)	Patient population	Regimens	Findings	Possible reason(s) for lack of trial success	Ref.
Viral vector	Allogeneic	PANVAC™-VF (falimarev)	MUC1, CEA	2, 13	NCT00088660 (phase 3)	Metastatic (stage IV) pancreatic cancer; already failed prior Gem	PANVAC™-VF + GM-CSF vs. BSC or palliative chemotherapy	Primary endpoint (OS) not met	Tumor burden (inappropriate population for vaccine monotherapy)	[15]
Viral vector	Allogeneic	PROSTVAC-V/F	PSA	22	NCT01322490 (phase 3, PROSPECT)	Asymptomatic/minimally symptomatic mCRPC	PROSTVAC ± GM-CSF vs. placebo	Primary endpoint (OS): Placebo: 34.3 months PROSTVAC: 34.4 months (HR comparison with placebo: 1.01 [95% CI 0.84–1.20, $p = 0.47$ ]) PROSTVAC + GM-CSF: 33.2 months (HR comparison with placebo: 1.02 [95% CI 0.86–1.22, $p = 0.59$ ])	Insufficient immune response or negative regulatory influences in the TME Ineffective as monotherapy Phase 2 false positive (underpowered for OS comparison) Potential for prolonged OS in the control arm relative to expected due to increasing availability of multiple life-extending treatments since the study was designed	[16]
Viral vector	Allogeneic	CMB 305	NY-ESO-1	10	NCT02609984 (phase 2, IMDZ-C232)	NY-ESO-1 + soft tissue sarcoma	CMB305 + atezolizumab vs. atezolizumab	Primary endpoints (OS, PFS), CMB305 + atezolizumab vs. atezolizumab: OS: 18.2 months vs. 18.0 months PFS: 2.8 months vs. 1.6 months	Imbalances in patient/disease characteristics: combination arm had more advanced disease and more prior lines of chemotherapy	[17]
Cell-based (tumor cell)	Allogeneic	Belagenpumatucel-L (Lucamix™)	-	-	NCT00676507 (phase 3)	Stage III/IV NSCLC; stable disease following frontline, platinum-based chemotherapy	Belagenpumatucel-L vs. placebo	Primary endpoint (OS), belagenpumatucel-L vs. placebo: median 20.3 months vs. 17.8 months (HR 0.94; 95% CI 0.73–1.20; $p = 0.59$ ); at second interim analysis, study was terminated for futility	Study design (late enrollment after induction therapy; single-agent therapy) Study did not require prior radiation within 6 months of randomization, which may have improved OS	[18]

Table 1 (continued)

Vaccine platform type	Autologous or allogeneic	Product/com-pound name	Antigen(s)	Antigen ranking <sup>a</sup>	Identifier (phase, name)	Patient population	Regimens	Findings	Possible reason(s) for lack of trial success	Ref.
Cell-based (tumor cell)	Allogeneic	GVAX®	-	-	NCT00089856 (phase 3, VITAL-1)	Metastatic, hormone-refractory prostate cancer	GVAX® vs. docetaxel + prednisone	Median survival (GVAX® vs. docetaxel + prednisone): 20.7 months vs. 21.7 months ( $p = 0.78$ ) Study terminated based on futility analysis showing < 30% chance of meeting primary endpoint Terminated early from lack of therapeutic effect	-	[11, 19]
Cell-based (tumor cell)	Allogeneic	GVAX®	-	-	NCT00133224 (phase 3; VITAL-2)	Taxane-naïve, metastatic, hormone-refractory prostate cancer patients with pain	GVAX® + docetaxel vs. docetaxel and prednisone	OS (GVAX® + docetaxel vs. docetaxel and prednisone): 12.2 months vs. 14.1 months ( $p = 0.01$ ) Accrual and treatment with GVAX® stopped because of IDMC recommendation Terminated early from lack of therapeutic effect	-	[11, 19]
Cell-based (tumor cell)	Allogeneic	Canvaxin™ (Cancer Vax)	-	-	NCT00052130 (phase 3, MMAIT-III)	Completely resected stage III melanoma	Canvaxin™ + BCG vs. BCG	Based on DSMB recommendation, study was terminated (low probability demonstrating significant improvement in Canvaxin™ containing treatment arm)	Population heterogeneity (burden of disease, heterogeneity of disease, immunological response)	[20, 21]

Table 1 (continued)

Vaccine platform type	Autologous or allogeneic	Product/com-pound name	Antigen(s)	Antigen ranking <sup>a</sup>	Identifier (phase, name)	Patient population	Regimens	Findings	Possible reason(s) for lack of trial success	Ref.
Cell-based (tumor cell)	Allogeneic	Canvaxin™ (CancerVax)	–	–	NCT00052156 (phase 3, MMAIT-IV)	Completely resected stage IV melanoma	Canvaxin™ + BCG vs. BCG	Primary endpoint (OS), Canvaxin™ + BCG vs. BCG: median 38.6 months vs. 34.9 months (HR 1.04; 95% CI 0.80–1.35; <i>p</i> = 0.77)	High survival in both treatment arms may be a result of selection bias; beneficial effect of metastasectomy, and/or use of BCG in control treatment arm	[20, 22]
Cell-based (tumor lysate)	Allogeneic	Melacine (theraccine)	–	–	– (phase 3)	Resected, intermediate-thickness, node-negative melanoma	After surgery: Melacine + DETOX adjuvant therapy vs. no further treatment	Primary endpoint (DFS, OS), vaccine vs. no treatment DFS: 107/300 events (tumor recurrences or deaths) vs. 114/300 (HR 0.92; Cox-adjusted $P_2$ = 0.51) OS: Not mature at time of publication	Study design (inadequately powered to detect small, clinically meaningful differences; methodology for staging regional nodes) Population heterogeneity	[23]
Cell-based (tumor lysate)	Allogeneic	Melacine (theraccine)	–	–	– (phase 3)	Stage IV melanoma with $\geq 1$ measurable lesion(s), ECOG PS 0–1	Melacine vs. DTIC, cisplatin, BCNU, and tamoxifen	Median survival (melacine vs. chemotherapy): 9.4 months vs. 12.3 months ( <i>p</i> = 0.16)	–	[24]
Cell-based (tumor cell)	Allogeneic	Algenpantucel-L (HyperA-cute® platform)	–	–	NCT01072981 (phase 3, IMPRESS)	Surgically resected pancreatic cancer, stage I or II (per AJCC)	Algenpantucel-L + SOC (Gem ± 5FU chemoradiation) vs. SOC alone	Primary endpoint (OS), algenpantucel-L + SOC vs. SOC: 30.4 months vs. 27.3 months; primary endpoint was not achieved	–	[25]



Table 1 (continued)

Vaccine platform type	Autologous or allogeneic	Product/com-pound name	Antigen(s)	Antigen ranking <sup>a</sup>	Identifier (phase, name)	Patient population	Regimens	Findings	Possible reason(s) for lack of trial success	Ref.
Cell-based (virus-augmented tumor cell)	Allogeneic	VMCL	–	–	–	Cutaneous melanoma; stage IIB or stage III (per AJCC); regional nodal involvement without evidence of systemic metastatic disease	VMCL vs. observation	<p>Median RFS: In eligible patients: 6.9 vs. 3.6 years (HR 0.86; 95% CI 0.70–1.07; <math>p = 0.17</math>)</p> <p>In ITT patients: 6.98 vs. 4.37 years (HR 0.89; 95% CI 0.72–1.09; <math>p = 0.27</math>)</p> <p>Median OS: In eligible patients: 12.6 vs. 7.3 years (HR 0.81; 95% CI 0.64–1.02; <math>p = 0.07</math>)</p> <p>In ITT patients: &gt;8.45 vs. 7.34 years (HR 0.83; 95% CI 0.67–1.04; <math>p = 0.11</math>)</p>	Better survival of control treatment arm in phase 3 study compared with phase 2 study	[26]
Cell-based (virus-augmented tumor cell)	Allogeneic	VMO	–	–	(phase 3)	Stage II melanoma (per IUAC) with positive lymph nodes	VMO vs. control (vaccinia virus alone)	<p>Median disease-free interval, VMO vs. control: 38.0 months vs. 37.0 months (<math>p = 0.99</math>)</p>	Population heterogeneity (potential differences for male vs. female patients)	[27]
Cell-based (virus-augmented tumor cell)	Allogeneic	VMO	–	–	(phase 3)	Stage III melanoma (per AJCC)	VMO vs. control (vaccinia virus alone)	<p>Median disease-free interval, VMO vs. control: 20.7 months vs. 26.9 months (<math>p = 0.61</math>)</p> <p>Median OS, VMO vs. control: 50.2 months vs. 41.3 months (<math>p = 0.79</math>)</p> <p>Median OS, 10-year follow-up, VMO vs. control: 7.71 years vs. 7.95 years (<math>p = 0.70</math>)</p>	Population heterogeneity (retrospective subset analysis showed that a subgroup of men may have a survival advantage with VMO)	[28, 29]

Table 1 (continued)

Vaccine platform type	Autologous or allogeneic	Product/com-pound name	Antigen(s)	Antigen ranking <sup>a</sup>	Identifier (phase, name)	Patient population	Regimens	Findings	Possible reason(s) for lack of trial success	Ref.
Cell-based (RNA elec-tropo-rated DC)	Autologous	Rocapuldencel-T (AGS-003)	—	—	NCT01582672 (phase 3, ADAPT)	Newly diagnosed meta-static RCC	Rocapuldencel-T + standard therapy vs. standard therapy alone	Primary endpoint (OS), rocapuldencel-T + standard therapy vs. standard therapy alone: median 27.7 months vs. 32.4 months (unadjusted HR 1.10; 95% CI 0.83–1.46; adjusted HR 1.06; 95% CI 0.79–1.40)	Insufficient long-term follow-up/potential delayed treatment effect	[30]
Cell-based (DC)	Autologous	Peptide-loaded DC vaccine	Several MHC class I- and class II-restricted peptides (9 or 10 mer) <sup>b</sup>	Includes 44, 8, 20, 16, 14	— (phase 3)	Metastatic (stage IV) melanoma	DC vaccine vs. DTIC	Primary endpoint (OR): < 6% in both treatment arms Following first interim analysis, study was prematurely closed (recommendation of external Data Monitoring and Safety Board because of extremely low probability of reaching study goals)	Population heterogeneity (significant differences in subgroups defined by performance status and HLA haplotype)	[31]



Table 1 (continued)

Vaccine platform type	Autologous or allogeneic	Product/com-pound name	Antigen(s)	Antigen ranking <sup>a</sup>	Identifier (phase, name)	Patient population	Regimens	Findings	Possible reason(s) for lack of trial success	Ref.
DNA vaccine	Allogeneic	Allovecin-7® (velimogene aliplasimid)	HLA-B7 and $\beta_2$ microglobulin	–	NCT00395070 (phase 3)	Recurrent stage III or stage IV melanoma	Allovecin-7® vs. chemotherapy alone (DTIC or TMZ)	No significant improvement in objective response rate at $\geq 24$ weeks (primary endpoint) or OS (secondary endpoint); based on this outcome, Allovecin program has been terminated	–	[32–34]
Ganglioside	Allogeneic	GM2-KLH (GMK)	GM2	–	EORTC 18961 (phase 3)	Stage II melanoma	GM2-KLH/QS-21 vs. observation after resection of primary tumor >1.5 mm	Primary endpoint (RFS), GM2-KLH/QS-21 ( $n = 627$ ) vs. observation ( $n = 627$ ): Second interim analysis: 135 events vs. 132 events (HR 1.00; 98% CI 0.75–1.34; $p = 0.99$ ); trial was stopped for futility Final analysis: 205 vs. 204 events (HR 1.03; 98% CI 0.84–1.25; $p = 0.81$ )	Vaccination schedule (i.e., impact of multiple vaccinations may be deleterious)	[35]

Table 1 (continued)

Vaccine platform type	Autologous or allogeneic	Product/com-pound name	Antigen(s)	Antigen ranking <sup>a</sup>	Identifier (phase, name)	Patient population	Regimens	Findings	Possible reason(s) for lack of trial success	Ref.
Gangli- oside	Allogeneic	GM2-KLH (GMK)	GM2	–	Intergroup trial E1694/S9512/ C509801 (phase 3)	Resected stage IIb/III melanoma (per AJCC)	GM2-KLH/ QS-21 vs. HDI therapy	Primary endpoint (RFS), GM2-KLH/ QS-21 vs. HDI therapy: In eligible patients: 151/389 (39%) events vs. 98/385 (25%) events (HR 1.47; 95% CI 1.14–1.90; log-rank one-sided $p < 0.05$ in favor of HDI [ $p$ = 0.0015]) In ITT population: HR 1.49 ( $p < 0.05$ in favor of HDI [ $p$ = 0.00045]) $P$ values for RFS crossed protocol- specified lower boundary, resulting in study termination Primary endpoint (OS), GM2-KLH/ QS-21 vs. HDI therapy: In eligible patients: (HR 1.52; 95% CI 1.07–2.15; log-rank one-sided $p = 0.01$ in favor of HDI) In ITT population: HR 1.38 ( $p = 0.02$ )	–	[36]
Protein (anti- idio- typic anti- body)	Allogeneic	Abagovomab	CA-125 (cleaved and released domain of MUC16)	–	NCT00418574 (phase 3, MIMOSA)	Stage III–IV epithelial ovarian, primary peritoneal, or fallopian tube cancer in first complete clini- cal remission	Abagovomab vs. placebo	Primary endpoint (RFS), abagovomab ( $n = 593$ ) vs. pla- cebo ( $n = 295$ ): 374 recurrence events vs. 180 recurrence events (HR 1.10; 95% CI 0.92–1.32; $p = 0.30$ )	Study design (combina- tion therapies and multi- antigen approaches remain reasonable approaches to study)	[37]

Table 1 (continued)

Vaccine platform type	Autologous or allogeneic	Product/com-pound name	Antigen(s)	Antigen ranking <sup>a</sup>	Identifier (phase, name)	Patient population	Regimens	Findings	Possible reason(s) for lack of trial success	Ref.
Protein (anti-idiotypic anti-body)	Allogeneic	BEC2	GD3	40	NCT00037713 (phase 3, SILVA)	Limited-disease SCLC	BEC2/BCG vs. observation	Primary endpoint (OS), BEC2/BCG vs. observation: median 14.3 vs. 16.4 months (HR 1.12; 95% CI 0.91–1.37; $p = 0.28$ )	Study design (imbalance in patient characteristics; choice of adjuvant or anti-idiotypic approach, single-antigen approach)	[38]
Protein (idiotype)	Autologous	MyVax (GTOP-99)	Patient-specific idiotype conjugated to KLH (recombinant DNA technique)	7	NCT00017290 (phase 3)	Previously untreated, advanced-stage (Ann Arbor stage III or IV) FL	MyVax + GM-CSF vs. control (KLH + GM-CSF)	Primary endpoint (PFS), MyVax vs. control: HR 0.98, 95% CI 0.72–1.33; $p = 0.89$	Insufficient humoral immune response (immune response observed in 41% of patients)	[39]
Protein (idiotype)	Autologous	Mitumprotimut-T (Specificid)	Patient-specific idiotype conjugated to KLH (recombinant DNA technique)	7	– (phase 3)	Treatment-naïve or relapsed/refractory CD20+ FL, WHO grade 1–3; candidate for rituximab	Mitumprotimut-T + GM-CSF vs. placebo (GM-CSF)	Primary endpoint (TTP), mitumprotimut-T vs. placebo: 9.0 months vs. 12.6 months (HR 1.38; 95% CI 1.05–1.82; $p = 0.02$ )	Imbalance in FLIPI risk groups Product (antigen and/or adjuvant selection) Insufficient immune response/inhibitory immune microenvironment	[40]

Table 1 (continued)

Vaccine platform type	Autologous or allogeneic	Product/com-pound name	Antigen(s)	Antigen ranking <sup>a</sup>	Identifier (phase, name)	Patient population	Regimens	Findings	Possible reason(s) for lack of trial success	Ref.
Protein (idiotype)	Autologous	BiovaxID®	Patient-specific idiotype conjugated to KLH (hybridoma technique)	7	NCT00091676 (phase 3)	Advanced stage FL in first remission (CR or CR unconfirmed) after chemotherapy	BiovaxID® + GM-CSF vs. control (KLH + GM-CSF)	<p>Primary endpoint (DFS), BiovaxID® + GM-CSF vs. control: For all 177 randomly assigned patients (includes 60 patients who did not receive vaccination): 23.0 months vs. 20.6 months (HR 0.81; 95% CI 0.56–1.16; <math>p = 0.26</math>)</p> <p>For the 117 patients who received vaccination: 44.2 months vs. 30.6 months (HR 0.62, 95% CI 0.39–0.99; <math>p &lt; 0.05</math> [<math>p = 0.047</math>])</p>	Study design (control treatment arm [KLH + GM-CSF vs. placebo]) Product (tumor Ig isotype may influence immunogenicity of vaccine)	[41]
Protein	Allogeneic	THERATOPE®	STn	56	NCT00003638 (phase 3)	MBC; previously received chemotherapy and had CR, PR, or no disease progression	STn-KLH vs. KLH	<p>Primary endpoints (TTP and OS), STn-KLH vs. KLH: TTP: 3.4 months vs. 3.0 months (Cox <math>p = 0.35</math>) OS: 23.1 months vs. 22.3 months (Cox <math>p = 0.92</math>)</p>	Study design (KLH as control arm rather than no treatment) Tumor burden (advanced metastatic disease) Treatment duration (continued vaccination beyond primary progression may have been advantageous)	[42]

Table 1 (continued)

Vaccine platform type	Autologous or allogeneic	Product/com-pound name	Antigen(s)	Antigen ranking <sup>a</sup>	Identifier (phase, name)	Patient population	Regimens	Findings	Possible reason(s) for lack of trial success	Ref.
Protein	Allogeneic	GSK2132231A	Recombinant MAGE-A3	8	NCT00796445 (phase 3, DERMA)	Resected, MAGE-A3-positive, stage III melanoma	GSK2132231A + AS15 vs. placebo	<p>Primary endpoint (DFS), GSK2132231A + AS15 vs. placebo: median 11.0 months vs. 11.2 months (HR 1.01; 95% CI 0.88–1.17; <math>p = 0.86</math>)</p> <p>In patients with potentially predictive gene signature: median 9.9 months vs. 11.6 months (HR 1.11; 95% CI 0.83–1.49; <math>p = 0.48</math>)</p>	<p>Product (choice of antigen, immunostimulant)</p> <p>Insufficient/absent immune response</p> <p>Target population (too advanced for antigen-specific immunotherapeutic alone)</p>	[43]
Protein	Allogeneic	GSK1572932A	Recombinant MAGE-A3	8	NCT00480025 (phase 3, MAGRIT)	Resected, MAGE-A3-positive, NSCLC	GSK1572932A + AS15 vs. placebo	<p>Primary endpoint (DFS), GSK1572932A + AS15 vs. placebo: median 60.5 months vs. 57.9 months (HR 1.02; 95% CI 0.89–1.18; <math>p = 0.74</math>)</p> <p>In patients who did not receive chemotherapy: median 58.0 months vs. 56.9 months (HR 0.97; 95% CI 0.80–1.18; <math>p = 0.76</math>)</p> <p>In patients with potentially predictive gene signature: not evaluated, as predictive gene signature could not be identified</p>	<p>Initial positive treatment effect in phase 2 trial may be a result of limited sample size and/or unnoticed imbalances across treatment groups</p>	[44]

Table 1 (continued)

Vaccine platform type	Autologous or allogeneic	Product/com-pound name	Antigen(s)	Antigen ranking <sup>a</sup>	Identifier (phase, name)	Patient population	Regimens	Findings	Possible reason(s) for lack of trial success	Ref.
Protein	Allogeneic	G17DT (Inse-gia)	Gastrin-17	–	NCT0044031 (phase 3)	Untreated with locally advanced, or recurrent, or metastatic pancreatic cancer	G17DT + Gem vs. placebo + Gem	Primary endpoint (OS), G17DT + Gem vs. placebo: Gem: 178 days vs. 201 days (HR 1.10; $p = 0.10$ )	Population heterogeneity (anti-G17 antibody levels correlated with OS)	[21]
Synthetic peptide (length: 25 mer)	Allogeneic	Tecemotide (L-BLP25, StimuVax)	MUC1	2	NCT00409188 (phase 3, START <sup>c</sup> )	Unresectable stage III NSCLC	Tecemotide vs. placebo	Primary endpoint (OS), tecemotide vs. placebo: 25.8 months vs. 22.4 months (adjusted HR 0.89; 95% CI 0.77–1.03; $p = 0.11$ )	Population heterogeneity (possibly more favorable effect in patients receiving concurrent as opposed to sequential chemoradiotherapy) Clinical hold potentially resulted in understimated treatment effect	[45, 46]
Synthetic peptide (length: 25 mer)	Allogeneic	Tecemotide (StimuVax; L-BLP25)	MUC1	2	NCT00925548 (phase 3, STRIDE)	ER-positive and/or PgR-positive, inoperable, locally advanced, recurrent, or metastatic BC in post-menopausal women	Tecemotide + hormonal therapy vs. hormonal therapy	Sponsor permanently terminated trial following clinical hold	Safety concerns	
Synthetic peptide (16 mer)	Allogeneic	GV1001	hTERT	23	ISRCTN4382138 (phase 3, Telo-Vac)	Locally advanced or metastatic pancreatic cancer; ECOG PS 0–2	Chemotherapy alone (Gem and capecitabine); chemotherapy with sequential GV1001 + GM-CSF; chemotherapy with concurrent GV1001 + GM-CSF	Primary endpoint (OS): Chemotherapy alone vs. sequential GV1001 + GM-CSF: median 7.9 months vs. 6.9 months (HR 1.19, 98.25% CI 0.97–1.48; $p = 0.05$ ) Concurrent GV1001 + GM-CSF: median 8.4 months (HR 1.05; 98.25% CI 0.85–1.29; $p = 0.64$ ; overall log-rank of $\chi^2_{df=4.3}$ ; $p = 0.11$ )	Nature of disease (early metastasizing, rapidly progressive may limit time to develop immune response; dense stromal reaction may impede/restrict synergistic potential of chemotherapy and GV1001)	[21, 47–49]

Table 1 (continued)

Vaccine platform type	Autologous or allogeneic	Product/com-pound name	Antigen(s)	Antigen ranking <sup>a</sup>	Identifier (phase, name)	Patient population	Regimens	Findings	Possible reason(s) for lack of trial success	Ref.
Synthetic peptide (16 mer)	Allogeneic	GV1001	hTERT	23	NCT00358566 (phase 3, Pri-movax)	Advanced, unresectable pancreatic cancer; ECOG PS 0–1	Gem alone vs. Gem with sequential GV1001 + GM-CSF	Preliminary data showed no survival benefit in GV1001 group vs. chemotherapy alone	–	[47, 49]
Synthetic peptide (14 mer)	Allogeneic	Rindopepimut (CDX-110)	EGFRvIII	5	NCT01480479 (phase 3, ACT IV)	Newly diagnosed, EGFRvIII-expressing glioblastoma	TMZ + rindopepimut + GM-CSF vs. control (TMZ alone)	Primary endpoint (OS), TMZ + rindopepimut + GM-CSF vs. control: Study was terminated at the second interim analysis for futility In the ITT population: median 17.4 months vs. 17.4 months (HR 0.89, 95% CI 0.75–1.07; $p = 0.22$ ) In the MRD population: median 20.1 months vs. 20.0 months (HR 1.01; 95% CI 0.79–1.30; $p = 0.93$ )	Patients in control treatment arm fared better in this study than matched control datasets Study design (control arm [KLH vs. inactive placebo]; TMZ use [treatment-induced lymphopenia may reduce immunotherapy efficacy]; vaccine started after radiotherapy vs. as early as possible) Product (single antigen rather than multi-peptide vaccine or other combination approaches)	[50, 51]
Synthetic peptide	Allogeneic	Elpamotide	VEGFR2	70	UMIN00002500 (phase 2/3, PEGASUS-PC)	Locally advanced or metastatic pancreatic cancer	Elpamotide + Montanide™ ISA 51 VG vs. placebo (saline + Montanide™ ISA 51 VG)	Primary endpoint (OS), elpamotide + Montanide™ ISA 51 VG vs. placebo: median 8.36 months vs. 8.54 months (HR 0.87; 95% CI 0.49–1.56; H-F $p = 0.92$ )	Population heterogeneity (subgroup analyses suggested that patients with strong injection site reactions may benefit from the vaccine, but these patients were limited in number) Study design (use of Montanide™ ISA 51 VG in control arm; single vs. multiple tumor targets)	[52]



Table 1 (continued)

Vaccine platform type	Autologous or allogeneic	Product/com-pound name	Antigen(s)	Antigen ranking <sup>a</sup>	Identifier (phase, name)	Patient population	Regimens	Findings	Possible reason(s) for lack of trial success	Ref.
Synthetic peptides (lengths: 9 mer; 9 mer; 9 mer)	Allogeneic	Peptide vaccine	Tyrosinase, gp100, MART-1/melan-A	20, 16, 14	ECOG E4697 study	Locally advanced (stage III) and/or stage IV melanoma with no evidence of disease after complete surgical resection	Peptide vaccine + Montanide™ ISA 51 ± GM-CSF vs. placebo ± GM-CSF	Secondary objective (RFS), peptide vaccine vs. placebo in HLA-A2+ patients: median 11.5 months vs. 9.8 months (HR 0.96; 95% repeated CI 0.74–1.23; <i>p</i> = 0.71)	Population heterogeneity (sites of metastases, baseline immune status) Insufficient immune response or lack of relevance of immune response Product (adjuvant selection, administration)	[53]
Synthetic peptide (length: 9 mer)	Allogeneic	Nelipepimut-S (E75; Neuvax™)	HER2	6	NCT01479244 (phase 3, PRE-SENT)	T1–T3, node-positive BC with low to intermediate HER2 expression	Nelipepimut-S + GM-CSF vs. GM-CSF	Primary endpoint (DFS), nelipepimut-S ( <i>n</i> = 376) vs. placebo ( <i>n</i> = 382): 37 recurrence events vs. 24 recurrence events; no significant difference in DFS events (HR 1.564; 95% CI 0.96–2.55; <i>p</i> = 0.07)	Study design (protocol-specified annual imaging instead of clinical assessment per ASCO guidelines hastened interim analysis [clinical significance of image-only recurrence events unclear])	[54, 55]
Protein-peptide complex	Autologous	Vitespen (HSPPC-96, Oncophage)	gp96-peptide complex	–	NCT00033904 (phase 3)	RCC at high risk of recurrence after nephrectomy	Vitespen vs. observation	Primary endpoint (RFS), vaccine vs. observation: 37.7% (136/361) vs. 39.8% (146/367) (HR 0.92; 95% CI 0.73–1.17; <i>p</i> = 0.51)	Study design (higher than expected number of patients with metastatic disease) Population heterogeneity (more targeted recruitment may have allowed enrollment of patients with earlier stage disease and better prognosis)	[56]

Table 1 (continued)

Vaccine platform type	Autologous or allogeneic	Product/com-pound name	Antigen(s)	Antigen ranking <sup>a</sup>	Identifier (phase, name)	Patient population	Regimens	Findings	Possible reason(s) for lack of trial success	Ref.
Protein-peptide complex	Autologous	Vitespen (HSPPC-96, Oncophage)	gp96-peptide complex	–	NCT00039000 (phase 3)	Stage IV melanoma, with expected resectability of some/all lesions to obtain $\geq 7$ g of cancer	Vitespen vs. physician's choice (DTIC/TMZ and/ or IL-2 and surgery)	Primary endpoint (OS), vaccine vs. physician's choice: HR 1.16; 95% CI 0.69–1.71; $p = 0.32$	Study execution (49% success rate for vaccine production based on suggested minimum threshold of four administrations in animal models)	[57]
Protein-peptide complex	Autologous	Vitespen (HSPPC-96, Oncophage)	gp96-peptide complex	–	NCT01814813 (phase 2)	Surgically resectable, recurrent glioblastoma	Vitespen + bevacizumab vs. bevacizumab alone	Primary endpoint (OS), vaccine + bevacizumab vs. bevacizumab alone at interim analysis: 7.5 months vs. 10.7 months (HR 2.06; 95% CI 1.18–3.60; $p = 0.008$ ); study terminated for futility	Population heterogeneity (exploratory analyses showed that patients with earlier stages of disease may have benefited from the vaccine)	[58]
Oncolytic virus	Allogeneic	pexastimogene devacirepvec (Pexa-Vec)	–	–	NCT02562755 (phase 3)	Advanced HCC without prior systemic therapy	Pexa-Vec vs. sorafenib	Interim futility analysis determined that the primary objective (OS) was not likely to be met	Difficult-to-treat population Immunosuppressive environment (liver) Imbalance between arms in salvage therapies received	[59]

**Table 1** (continued)

AJCC American Joint Committee on Cancer, ASCO American Society of Clinical Oncology, BC breast cancer, BCG bacillus Calmette–Guérin, BCNU carmustine, BSC best supportive care, CEA carcinoembryonic antigen, CI confidence interval, CR complete response, DC dendritic cell, DETOX detoxified Freund's adjuvant, DFS disease-free survival, DSMB Data and Safety Monitoring Board, DTIC dacarbazine, ECOG PS Eastern Cooperative Oncology Group performance status, EGFRvIII epidermal growth factor receptor variant III, EORTC European Organization for Research and Treatment of Cancer, ER estrogen receptor, FL follicular lymphoma, FLIP Follicular Lymphoma International Prognostic Index, Gem gemcitabine, GM-CSF granulocyte-macrophage colony-stimulating factor, HCC hepatocellular carcinoma, HDI high-dose interferon- $\alpha$ -2b, HER2 human epidermal growth factor receptor 2, H-F Harrington-Fleming test, HLA human leukocyte antigen, HR hazard ratio, HSPPC-96 heat shock protein peptide complex-96, hTERT human telomerase reverse transcriptase, IDMC independent data monitoring committee, Ig immunoglobulin, IL-2 interleukin-2, ITT intent-to-treat, IUAC International Union Against Cancer, KLH keyhole limpet hemocyanin, MAGE melanoma-associated antigen, MBC metastatic breast cancer, mCRPC metastatic castration-resistant prostate cancer, MHC major histocompatibility complex, MRD minimal residual disease, MUC1 mucin 1, NSCLC non-small-cell lung cancer, NY-ESO-1 New York esophageal squamous cell carcinoma 1, OR objective response, OS overall survival, PFS progression-free survival, PgR progesterone receptor, PR partial response, PSA prostate-specific antigen, RCC renal cell carcinoma, RFS recurrence-free survival, SCLC small-cell lung cancer, SOC standard of care, STn sialyl-Tn, TME tumor microenvironment, TMZ temozolomide, TTP time to progression, VEGFR2 vascular endothelial growth factor receptor 2, VMCL vaccinia melanoma cell lysate, VMO vaccinia melanoma oncolysate, WHO World Health Organization

<sup>a</sup>Antigen ranking based on Table 3 of Cheever, et al. *Clin Cancer Res.* 2009;15(17):5327–37[14]

<sup>b</sup>MHC class I- and II-restricted peptides included MAGE-1, MAGE-3, tyrosinase, gp100 analog, and MART-1/melan-A analog

<sup>c</sup>Discontinuation of clinical development program for tecemotide monotherapy in stage III NSCLC also included phase 3 START2 (NCT02049151) and INSPIRE (NCT01015443) trials.

vaccine to checkpoint inhibition did not yield a survival benefit in soft tissue sarcoma [17]. There is very limited clinical investigation of the PANVAC vaccine for pancreatic tumors (NCT00669734).

Like viral platforms, plasmid vector-based vaccines have intrinsic adjuvant immunogenicity because the nucleic acid itself may be immunostimulatory, triggering an innate immune response [73]. Other advantages of this platform are enhanced stability, ease of manufacturing, induction of intracellular antigen expression, and if full-length genes are utilized, there is no HLA restriction. Despite these advantages, there has been limited late-stage investigation of DNA or RNA vaccines to date. The DNA vaccine Allovectin-7<sup>®</sup>, which contains DNA sequences for HLA-B7 and  $\beta_2$ -microglobulin, did not improve objective response rate or OS compared with chemotherapy in patients with advanced melanoma in a phase 3 trial, leading to termination of the development program. There have since been numerous improvements in DNA and RNA delivery technologies (e.g., electroporation) and more modern nucleic acid vaccines have been tested in early-stage studies, as described in Sect. 3.

## 2.4 Other Types of Cancer Vaccines

In contrast to the therapeutic cancer vaccines described previously, some vaccines used in the treatment of cancer do not deliver defined tumor antigens to generate anti-tumor immunity, but nevertheless, generate an immune response. For example, intravesical immunotherapy (i.e., vaccination) with *Mycobacterium bovis* bacillus Calmette–Guérin (BCG) is approved for the treatment of certain types of bladder cancer. The mechanism of action of BCG immunotherapy has not been well understood, but recent data indicate that BCG improves the activation and exhaustion status of tumor-specific T cells [74].

Another cancer vaccine approach involves strategies to modify or inflame tumor cells by intratumoral administration of oncolytic viruses. In 2015, the oncolytic viral vaccine T-VEC, a herpes virus genetically modified to express GM-CSF [13], was licensed for the treatment of patients with unresectable melanoma. The approval of T-VEC was based on the phase 3 OPTiM trial (NCT00769704) demonstrating that a higher proportion of patients treated with T-VEC versus GM-CSF had a durable clinical response ( $\geq 6$  months continuously and beginning within the first year; 16.3% vs. 2.1%, respectively;  $p < 0.001$ ; Table 2). There was also a trend for improved OS in the T-VEC group (23.3 vs. 18.9 months;  $p = 0.051$ ) [67]. The mechanism of action for oncolytic viral vaccines such as T-VEC is different to that of other notable cancer vaccines. This form of “in situ” vaccination results in the killing of tumor cells by the virus and the release of tumor antigens [75, 76]. These

**Table 2** Pivotal studies supporting approval of therapeutic cancer vaccines

Vaccine platform type	Product name	Antigen(s)	Identifier (phase, name)	Patient population	Regimens	Findings	Reference
Cell-based	Sipuleucel-T	PA2024	NCT00065442 (phase 3, IMPACT)	Metastatic castration-resistant prostate cancer	Sipuleucel-T vs. placebo	Primary endpoint (overall survival): Median of 25.8 months (sipuleucel-T) vs. 21.7 months (placebo); HR 0.78; 95% CI 0.61–0.98; $p = 0.03$	[66]
Cell-based (oncolytic virus)	Talimogene laherparepvec (T-VEC)	N/A	NCT00769704 (phase 3, OPTiM)	Unresected stage IIIB to IV melanoma	Intralesional T-VEC vs. subcutaneous recombinant GM-CSF	Primary endpoint (durable response rate): 16.3% (T-VEC) vs. 2.1% (GM-CSF); odds ratio 8.9; $p < 0.001$	[67]

CI confidence interval, *GM-CSF* granulocyte-macrophage colony-stimulating factor, *HR* hazard ratio, *N/A* not applicable

effects lead to the immune-mediated regression of distant tumor lesions, presumably either through amplification of previously activated host-immunity and/or the priming of new anti-tumor immune responses [75–77]. Evidence for tumor-specific T cell induction after T-VEC treatment was observed in a clinical study of patients with stage IIIc and stage IV melanoma. In this study, a patient with a complete response after T-VEC injection had an increase in MART-1-specific effector T cells, both in the injected target lesion and in a nontarget lesion [77].

### 3 The Evolution of Cancer Vaccine Development: Current Strategies Based on Historical Experience and Scientific Advances

The limited success of therapeutic cancer vaccines despite decades of development by academia and industry raises the questions of why expectations have not been fulfilled and how barriers to successful development can be overcome. Advances in our understanding of antigen immunogenicity, the importance of antigen presentation, and the dynamics of how cancer cells evade and suppress the host immune system suggest that previous studies may have used suboptimal antigen targets, vaccine designs, and/or trial designs (including patient populations). In 2015, Melief et al. formulated a list of attributes that cancer vaccines would need to have to be successful [6]. In brief, these attributes stress the importance of broad stimulation of CTLs and T helper cells through two mechanisms: (1) selection of appropriate

antigens that induce both T cell populations and (2) rational vaccine designs that achieve concentrated delivery of tumor antigens to activated DCs, where epitopes derived from exogenous tumor antigens can be loaded onto both MHC class I (through the cross-presentation pathway) and MHC class II molecules to stimulate CTLs and T helper cells, respectively (Fig. 2) [6, 78]. Over the last decade, therapeutic cancer vaccine strategies have improved, incorporating better immunogenicity, antigen selection, and structural design to meet these criteria.

#### 3.1 Selecting the Appropriate Antigen

Choosing optimal antigens has been described as the most important consideration in the design of therapeutic vaccines [11]. Antigen selection affects critical vaccine properties, including the ability to generate a strong and broad immune response, target cancer stem cells to prevent relapse, and avoid off-target effects on normal cells. Optimal antigen discovery is hindered by the limited number of suitable immunogenic antigens fitting these criteria within the context of an immense number of potential antigens [14]. Current strategies aim to efficiently identify appropriate antigens for cancer vaccines either in the form of “ideal” shared tumor antigens or more personalized neoantigens.

Shared TAAs are self-proteins with preferential or abnormal expression in cancer cells versus normal cells [11], and these have been the primary type of antigen tested in clinical trials [11]. An important advance in the field occurred in 2009 when a National Cancer Institute (NCI) Pilot Project developed a list of nine “ideal” cancer antigen criteria, which

comprised therapeutic function, immunogenicity, stem cell expression, specificity, oncogenicity, expression level and percentage of antigen-positive cells, number of patients with antigen-positive cancers, number of epitopes, and cellular location of expression [14]. The NCI project then used a mathematical model to quantitatively rank 75 antigens according to how well they met these criteria [14]. As shown in Table 1, antigens used in past pivotal trials of cancer vaccines were often not ranked highly on the NCI list, indicating that early vaccines may have lacked efficacy because of sub-optimal antigen selection. Tumor antigens employed in these vaccines may also have been insufficiently immunogenic. Several peptide-based vaccines with negative phase 3 results targeted tumor antigens that would have been retrospectively deemed not a high priority target by consensus NCI criteria [14]. For example, the GV1001 peptide vaccine targeted a TAA (human telomerase reverse transcriptase [hTERT]) ranked 23 out of 75 possible antigens by the NCI list [14]. This vaccine failed to generate sufficient immune responses and did not improve OS compared with chemotherapy alone in patients with pancreatic cancer [49].

Development of the NCI's prioritized list provides the impetus to investigate highly ranked TAAs. Within this list, Wilms' tumor 1 (WT1) protein, a zinc finger transcription factor, was considered the most encouraging antigen among the 75 assessed [14]. Early-stage trials of cancer vaccines targeting WT1 or other highly ranked TAAs, such as mucin 1 (MUC1) and human epidermal growth factor receptor 2 (HER2)/neu, are ongoing and have demonstrated tumor-specific immune responses [79–93]. Notably, randomized phase 2 studies of WT1 vaccines have shown trends toward survival benefit [84, 94], and a phase 3 study of a WT1-based vaccine is being planned (Table 3) [95]. Additionally, the phase 3 ATALANTE-1 trial (NCT02654587) comparing a cancer vaccine targeting five TAAs (angiotensin-converting enzyme [ACE], HER2, melanoma-associated antigen (MAGE) 2, MAGE3, and p53) with chemotherapy in patients with advanced non-small-cell lung cancer who relapsed after checkpoint inhibitor therapy demonstrated an encouraging 1-year OS rate of 46% in the vaccine group ( $n = 61$ ) compared with 36% in the chemotherapy group ( $n = 31$ ) [96]. Taken together, these late-stage studies underscore the potential and maturity of targeting highly ranked TAAs as a treatment strategy for cancer.

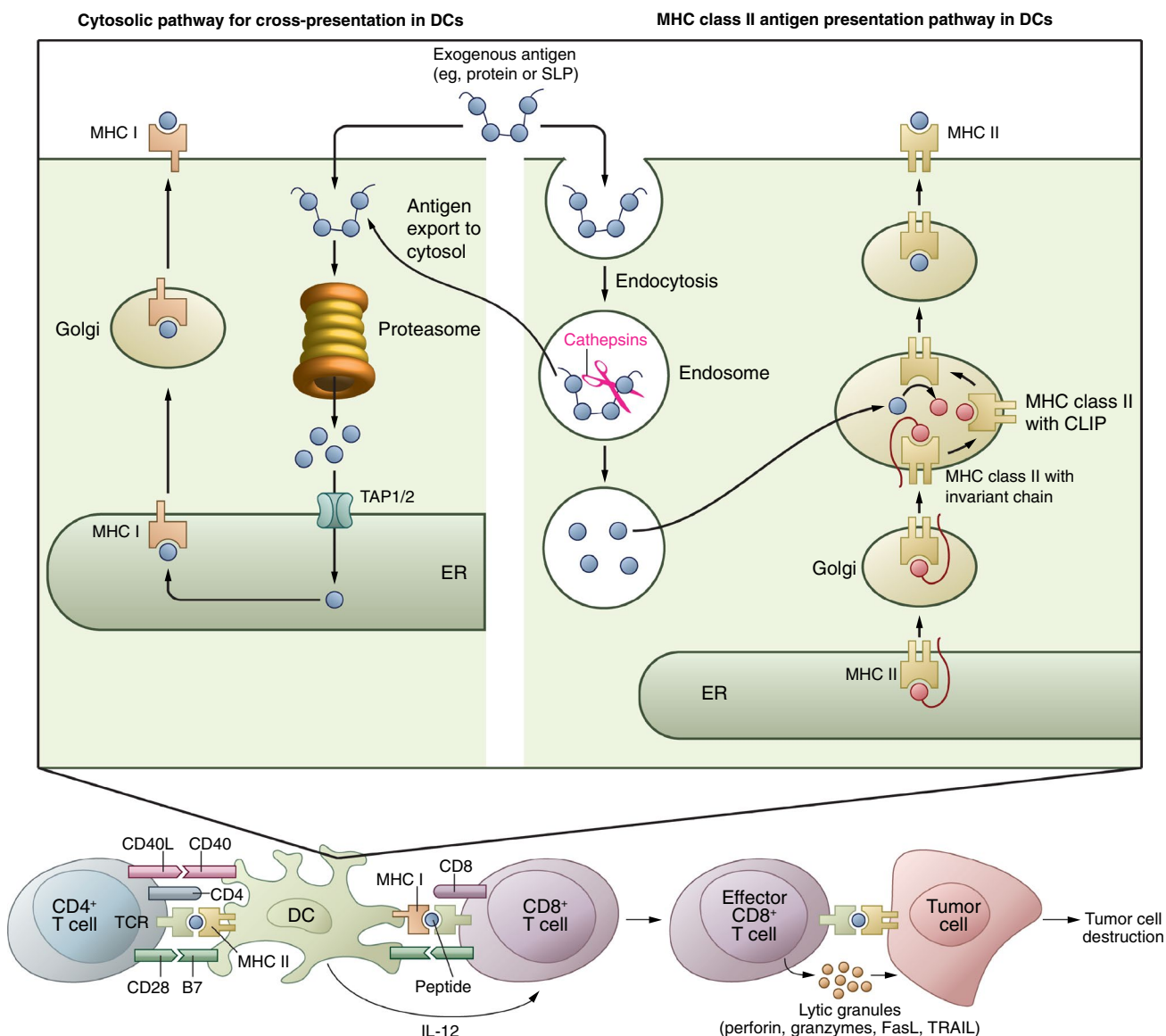
A limitation of shared TAAs is that they are autologous antigens, and immune self-tolerance mechanisms may deplete or eliminate TAA-specific T cells with high functional avidity, resulting in a tolerized T cell repertoire with relatively low reactivity toward the TAA [6]. Although increased immunogenicity has enabled TAA-based vaccines to break this tolerance, parallel advances in genomics have now allowed the efficient identification of another class of

cancer antigen, termed neoantigen, which is not subject to immune self-tolerance mechanisms [6]. Neoantigens are aberrant peptides that arise from genetic and epigenetic alterations (point mutations, insertions/deletions, gene fusions/translocations, splice variants, and post-translational modifications) in cancer cells [97]. Because these alterations are not part of the normal exome or transcriptome, the encoded neoantigens are tumor specific. Mutated peptides that are dissimilar to the self-proteome are more likely to be seen as novel by the immune system, and therefore be more immunogenic, compared with those that are similar to the self-proteome [98].

Broadly, candidate neoantigens are identified through a two-step process. First, whole exome and transcriptome sequencing of normal and cancerous cells allows identification of tumor-specific mutations (i.e., the mutanome) [100]. Next, neoepitopes are prioritized by *in silico* prediction of the binding affinity of each mutant peptide to MHC molecules. Advances in massive parallel sequencing have drastically accelerated this process, enabling feasible and high-throughput identification of tumor neoantigens for cancer vaccines [99, 100]. Because of these technological advancements and parallel innovations in cancer immunotherapy, the Human Vaccines Project (a public–private partnership with a goal of accelerating the development of cancer vaccines) ranked neoantigens as a high priority target for clinical translation [100].

Several early-stage clinical studies in patients with solid tumors showed that personalized neoantigen vaccines are safe, feasible, and able to augment neoantigen-specific T cell responses [101–108]. Notably, although most neoantigen vaccine clinical studies have been restricted to cancer types with high mutation burdens (e.g., melanoma) and thus more neoantigen potential, recent data from patients with glioblastoma also show neoantigen immune responses for cold tumors with low mutation burdens [106, 109]. Whether the generation of immune responses to neoantigens is therapeutically relevant remains uncertain, as mutations in expressed genes rarely result in the presentation of T cell targetable neoantigens on the cell surface [110] and few neoantigen vaccine studies have attempted to verify that the mutated peptide is present on the surface of the tumor cell. One of the few studies to assess the presence of mutated peptides on the tumor cell surface found 643 genomic mutations among 15 patients with glioblastoma but did not identify any of these mutations in the HLA peptidome by mass spectrometry [111]. Although this study found that neoepitopes induced T cell responses [111], the failure to identify surface mutated peptides calls into question the role of neoantigens for tumors with low tumor mutation burden. Overall, these initial studies of neoantigen cancer vaccines have provided proof of concept, have provided rationale for larger studies





**Fig. 2** Optimal antigen processing and presentation by DCs is important for effective immune-mediated tumor cell destruction [6]. Antigens enter DCs through multiple mechanisms, including endocytosis, phagocytosis, pinocytosis, and receptor-mediated uptake. These antigens are processed by DCs into peptide fragments (epitopes) before being loaded onto MHC class I molecules through cross priming or MHC class II molecules through the classical exogenous presentation pathway. T cell recognition of these epitopes occurs via binding between the TCR and the peptide-MHC complex on the DC. Following epitope recognition, CD40L expressed by CD4+ T cells activates DC-expressed CD40 to promote DC maturation and IL-12 secretion. This subsequently stimulates CD28 signaling and activation of CD8+ T cells. When the TCR of an effector CD8+ T cell binds to a tumor cell, an immunological synapse forms and lytic granules are secreted

by the effector CD8+ T cell, resulting in tumor cell destruction. Note: Cytosolic and vacuolar pathways for cross-presentation have been described. The figure presents the cytosolic pathway as it has been suggested this is the predominant pathway for cross-presentation [78]. [Figure adapted with permission of the *Journal of Clinical Investigation*, from “Therapeutic Cancer Vaccines,” Cornelis J.M. Melief et al, Volume 125, Issue 9, 2015; permission conveyed through Copyright Clearance Center, Inc.] CD cluster of differentiation, CLIP class II-associated invariant chain peptide, DC dendritic cell, ER endoplasmic reticulum, FasL Fas ligand, IL interleukin, MHC major histocompatibility complex, SLP synthetic long peptide, TAP transporter of antigen processing, TCR T cell receptor, TNF tumor necrosis factor, TRAIL TNF-related apoptosis-inducing ligand

and continued research and development [11], and have highlighted a need for neoantigen vaccine studies to verify the presence of targetable, mutated peptides on the tumor cell surface.

There are several ongoing efforts to optimize neoantigen vaccines. Cost-effective and efficient workflows and algorithms for more accurate prediction of which mutated peptides will stimulate the most potent anti-tumor T cell

Table 3 Select ongoing phase 3 studies evaluating cancer vaccines

Vaccine platform type	Product name	Antigen(s)	Identifier (phase, name)	Patient population	Enrollment	Regimens	Primary outcome measures
Cell-based (trivalent DC)	-	Autologous tumor stem cells, survivin, and hTERT	NCT03548571 (phase 2/3, DEN-STEM)	Glioblastoma IDH wild-type, with unmethylated MGMT-gene promoter	60	Trivalent DC immunization vs. radiotherapy with concomitant and adjuvant temozolomide	PFS
Peptide	GP96 heat shock protein-peptide complex	-	NCT04206254 (phase 2/3)	Liver cancer	80	GP96 vaccination after surgery vs. no treatment after surgery	2-year recurrence-free survival rate
Adenoviral vector containing the herpes simplex virus thymidine kinase gene	ProstAtak® (AdV-tk) + valacyclovir	-	NCT01436968 (phase 3)	Localized prostate cancer (intermediate risk or one NCCN high-risk feature) due to undergo standard prostate-only EBRT	711	ProstAtak® (AdV-tk) + valacyclovir + radiation therapy ± androgen deprivation therapy vs. placebo + valacyclovir + radiation therapy ± androgen deprivation therapy	DFS
Cell-based (bacterial)	BCG Tokyo-172 strain solution	-	NCT03091660 (phase 3)	Stage 0/Is/1 urothelial carcinoma	969	Tokyo-172 strain BCG (arm 2) vs. Tokyo-172 strain BCG solution with priming (arm 3) vs. TICE® BCG (arm 1)	Time to high-grade recurrence for arm 1 vs. arm 2, and arm 2 vs. arm 3
Cell-based (DCs)	DCs plus autologous tumor RNA	-	NCT01983748 (phase 3)	Stage T2, T3, or T4 melanoma of the uvea	200	Autologous DCs loaded with autologous tumor RNA vs. SOC	Prolongation of OS
Cell-based (tumor cell)	OncovAX®	-	NCT02448173 (phase 3)	Stage II colon cancer	550	OncovAX® and surgery vs. surgery	DFS
Oral vaccine (tablet) derived from pooled blood	Hepcortespensimut-L (Hepko-V5)	-	NCT02232490 (phase 3, Hepko-V5)	Advanced hepatocellular carcinoma	120	Hepcortespensimut-L vs. placebo	Changes in plasma AFP
Cell-based (bacterial)	BCG	-	NCT04165317 (phase 3)	High-risk non-muscle-invasive transitional cell carcinoma of the urothelium and complete resection of all Ta/T1 papillary disease	999	PF-06801591 + BCG induction and maintenance (arm A) vs. PF-06801591 + BCG induction only (arm B) vs. BCG induction and maintenance (arm C)	EFS (arm A vs. arm C and arm B vs. arm C)



Table 3 (continued)

Vaccine platform type	Product name	Antigen(s)	Identifier (phase, name)	Patient population	Enrollment	Regimens	Primary outcome measures
Cell-based (bacterial)	OncoTICE®/ ImmuCYST®/ TheraCys®	–	NCT02948543 (phase 3, ANZUP 1301)	Confirmed high-grade pT <sub>a</sub> or stage pT <sub>1</sub> (any grade) non-muscle-invasive bladder cancer and transurethral resection	500	Intravesical BCG (arm A) vs. intravesical BCG + mitomycin C	DFS
Cell-based (bacterial)	BCG-Medac®	–	NCT03799835 (phase 3, ALBAN)	High-risk non-muscle-invasive urothelial carcinoma	614	BCG vs. BCG + atezolizumab	RFS
Cell-based (DCs)	nDC	–	NCT02993315 (phase 3, MIND-DC)	Stage III cutaneous melanoma, classified as IIIB or IIIC disease	210	nDC vaccination vs. placebo	RFS
Cell-based (tumor cell)	Vigil	–	NCT03495921 (phase 3, VITA)	Patients aged ≥ 2 years with histologically confirmed ESFT and relapsed/refractory to 1 line of systemic chemotherapy	114	Vigil + irinotecan and temozolomide vs. irinotecan and temozolomide	PFS
Cell-based (bacterial)	BCG	–	NCT03528694 (phase 3, POTOMAC)	High-risk transitional cell carcinoma of the urothelium of the urinary bladder confined to the mucosa/submucosa; for patients who have received complete resection of all T <sub>a</sub> /T <sub>1</sub> papillary disease prior to randomization	973	Durvalumab plus BCG (induction + maintenance) vs. durvalumab plus BCG (induction only) vs. BCG	DFS for durvalumab plus BCG (induction + maintenance) vs. BCG maintenance) vs. BCG
Cell-based (bacterial)	TICE® BCG OncoTICE®	–	NCT03711032 (phase 3, KEYNOTE-676)	Histologically confirmed non-muscle-invasive (T <sub>1</sub> , high grade T <sub>a</sub> and/or CIS) transitional cell carcinoma of the bladder previously treated with BCG induction therapy followed by persistent/recurrent disease and has received cystoscopy/TURBT to remove all resectable disease	1525	BCG (induction and maintenance) plus pembrolizumab vs. BCG monotherapy (induction and maintenance)	CRR by blinded independent central review, EFS

Table 3 (continued)

Vaccine platform type	Product name	Antigen(s)	Identifier (phase, name)	Patient population	Enrollment	Regimens	Primary outcome measures
TAA vaccine	OSE2101/ Tedopi/ EP-2101 EP2101 IDM-2101	HLA A2 restricted "optimized epitopes" from ACE, HER2, MAGE2, MAGE3, and P53	NCT02654587 (phase 3, ATALANTE-1)	Locally advanced NSCLC (stage III) unsuitable for radiotherapy, or metastatic (stage IV) with disease recurrence/progression after an immune checkpoint inhibitor and platinum-based chemotherapy	363	OSE2101 vs. docetaxel or pemetrexed	OS
Analog peptide vaccine	Galimpepimut-S (SL-S-001)	WT1	NCT04229979 (phase 3)	AML in second complete remission or in second complete remission with incomplete platelet recovery	116	Galimpepimut-S vs. best available therapy (observation, HMA monotherapy, venetoclax monotherapy, or low-dose cytarabine)	OS
Cell-based (bacterial)	BCG	–	NCT03664869 (phase 3, Finnbladder-10)	High-risk non-muscle-invasive bladder cancer confined to the bladder (high-grade Ta/any T1 following 2nd resection)	300	BCG instillation and maintenance therapy vs. sequential BCG and EMDA mitomycin C	Bladder cancer recurrence rate

Includes studies that are planned or are recruiting patients

ACE angiotensin-converting enzyme, AFP  $\alpha$ -fetoprotein, AML acute myelogenous leukemia, BCG bacillus Calmette-Guérin, CIS carcinoma in situ, CRR complete response rate, DC dendritic cell, DFS disease-free survival, EBRT external beam radiation therapy, EFS event-free survival, EMDA electromotive drug administration, ESFT Ewing sarcoma family of tumors, HER2 human epidermal growth factor receptor 2, HLA human leukocyte antigen, HMA hypomethylating agent, hTERT human telomerase reverse transcriptase, IDH isocitrate dehydrogenase, MAGE melanoma-associated antigen, MGMT O<sup>6</sup>-methylguanine-DNA methyltransferase, NCCN National Comprehensive Cancer Network, nDC natural dendritic cell, NSCLC non-small-cell lung cancer, OS overall survival, PFS progression-free survival, RFS recurrence-free survival, SOC standard of care, TAA tumor-associated antigen, TURBT transurethral resection of the bladder tumor, WT1 Wilms' tumor 1

response are being investigated [112, 113]. Innovative neoantigen vaccine designs are also being explored, with pre-clinical data showing that DNA-based neoantigen vaccines can generate robust CTL-driven anti-tumor responses and delay tumor progression [114]. Furthermore, results from a combined approach using both TAAs and neoantigens in patients with newly diagnosed glioblastoma suggest that mixtures of antigenic targets may provide sustained anti-tumor responses by central memory CTLs and T helper cells [111].

## 3.2 Evolution of Therapeutic Cancer Vaccine Designs

Over the past decade, vaccine designs have evolved to elicit effective immune responses characterized by potent and broad stimulation of CTLs and T helper cells as well as enhanced antigen presentation by activated DCs. These optimizations, summarized below for the most encouraging platforms, are currently being used in the next generation of therapeutic cancer vaccines with the hope they will lead to improved immune and clinical responses compared with historical experience.

### 3.2.1 Peptide Vaccines

In response to the observation that short peptides and long protein sequences resulted in inadequate clinical activity (Table 1), extension of the amino acid sequence beyond the minimal-length CTL epitope or other short sequences has been shown to achieve more concentrated and selective delivery of antigens to DCs with sustained presentation [63]. Vaccination with synthetic long peptides has induced more robust and durable T cell responses compared with the minimal-length epitopes in preclinical models [6, 115, 116]. Because of these advantages, modern synthetic peptide vaccine designs commonly use at least one long peptide [6, 117], representing an important advance over the use of minimal-length peptide constructs.

Amino acid substitutions on the native TAA sequence (epitope enhancement) can be rationally implemented to improve binding stability to APCs and thus increase the likelihood of successful antigen presentation to T cells [60, 61, 87]. Modification of the peptide structure to increase amphiphilicity may also increase peptide immunogenicity, as demonstrated in the development of the BiVax peptide/polyinosinic-polycytidylic acid (poly I:C) subunit vaccine [118].

Recent evidence demonstrates that T helper cells play critical roles in induction of strong and long-lasting immune-mediated anti-tumor responses [119, 120]. In particular, targeting T helper type 1 cells is thought to be optimal, as they have been shown to have potent effects in inducing and

maintaining anti-tumor immunity, whereas T helper type 2 cells may actually promote neoplastic transformation in certain contexts [121]. While early peptide vaccine designs only targeted CTLs, modern synthetic peptide vaccines now typically include CTL and helper peptides in an effort to increase immunogenicity and improve clinical efficacy [85, 87].

### 3.2.2 DC Vaccines

The evolution of DC cancer vaccines was sparked by increased insight into DC biology and technology advances, recently reviewed in 2017 by Garg et al., who classified the development of this DC vaccine platform according to first-generation, second-generation, and next-generation designs [122]. Notably, sipuleucel-T, the only approved DC-based vaccine (licensed in 2010), was considered by Garg et al. to be on the borderline of first- and second-generation designs.

In brief, first-generation designs were characterized by the use of immature monocyte-derived DCs; the development of maturation cocktails enabled the consistent use of mature monocyte-derived DCs in second-generation constructs [122]. This advancement was important because compared with immature DCs, mature DCs express higher levels of MHC and costimulatory molecules, produce more cytokines, and traffic more efficiently to lymph nodes [123]. All these effects make mature DCs more potent activators of the immune system, which in turn has been found to translate to improved efficacy in clinical trials. Indeed, Garg et al. reported that in many trials, second-generation DC vaccines produced higher response rates and increased median OS compared with first-generation designs [122].

The transition from second-generation to next-generation DC vaccines was characterized by the use of patient-derived specific DC subsets (e.g., myeloid and plasmacytoid DCs) with specialized functionalities (antigen presentation, interferon responses, migratory capacity) superior to those of monocyte-derived DCs [122, 124]. This transition was enabled by incorporation of antibody-coated magnetic bead technology for more rapid and pure isolation of native DCs compared with older techniques such as density centrifugation [122, 124]. Next-generation DC vaccines are currently being tested in clinical trials [124]; results presented so far from phase 1 or 2 studies have demonstrated their feasibility and safety, with some encouraging OS durations observed [124–127]. A phase 3 study (NCT02993315) comparing next-generation DC vaccination with placebo as adjuvant therapy for patients with stage III melanoma will provide more robust survival data and clarify the clinical efficacy of this vaccination approach [124].

More recently, a strategy employing intratumoral DCs as part of an *in situ* vaccine has been described. This *in situ* vaccine approach uses a triplet consisting of injection of FMS-like tyrosine kinase 3 ligand at the target lesion to

generate accumulation of intratumoral DCs, local tumor irradiation to load the intratumoral DCs with TAAs released from dying tumor cells, and injections of a Toll-like receptor (TLR) agonist at the target lesion to drive intratumoral DC activation [128]. In essence, this triplet creates a DC-based vaccine at the site of the tumor [128]. This *in situ* vaccine was recently evaluated in a phase 1 clinical trial (NCT01976585) for patients with advanced stage indolent non-Hodgkin lymphoma, where it was reported to be well tolerated and capable of producing durable regressions at distant tumor sites via an abscopal effect [128]. Of 11 patients who received the *in situ* vaccine, eight had partial or complete remissions of the treated tumor with regard to the non-treated tumors, six patients had stable disease or minor regressions lasting 3–18 months, and three achieved remission [128].

### 3.2.3 Genetic Vaccines

The main limitations of early nucleic acid-based vaccine designs have been limited uptake of the nucleic acid by DCs and other cells, either because of low transfection efficiency or degradation [11], and the resultant low immunogenicity observed in clinical trials. For DNA- or RNA-based vaccines, several upgrades have allowed the prospect of improved transfection rates and immunogenicity as described by Hollingsworth and Jansen [11] including use of electroporation, sonoporation, nanoparticles [129], gene guns, microneedle arrays, needle-free injection [130], and liposomal encapsulation.

Based on encouraging phase 2b data showing significantly higher regression rates of cervical intraepithelial neoplasia compared with placebo [11, 131], a DNA-based vaccine using electroporation is currently being evaluated in two phase 3 studies (REVEAL 1, NCT03185013; REVEAL 2, NCT03721978) to treat patients with precancerous lesions of the cervix (high-grade squamous intraepithelial lesions associated with human papillomavirus). Early-stage studies are evaluating DNA-based vaccines using electroporation for a variety of solid tumors (NCT03199040, NCT03122106, NCT03532217, NCT03439085, NCT02204098, and NCT04397003). In phase 1 studies, RNA-based vaccines using either electroporation [108] or RNA-lipoplexes [132] to improve systemic DC targeting have demonstrated encouraging immune-mediated anti-tumor activity for patients with melanoma.

Other methods to improve the immunogenicity of nucleic acid-based vaccines have been recently reviewed by Lopes et al. [133]. To break immune tolerance and target multiple TAAs, DNA vaccines encoding xenoantigens or chimeric

proteins have been studied [133]. In animal models, chimeric DNA vaccines have induced potential anti-tumor effects [134, 135], and one such vaccine is approved for the treatment of canine melanoma. Numerous human clinical trials are currently evaluating polyepitope DNA vaccines, which aim to induce a broad T cell response through the simultaneous delivery of multiple antigens [133].

### 3.3 The Role of Adjuvants

Effective therapeutic cancer vaccines rely on antigen presentation and activation of the immune system by DCs; however, suppression of DC maturation and function is a hallmark of cancer immune evasion [136]. Even worse, many subsets of DCs are in an immature state and produce “self”-tolerizing messages to the immune system. Therefore, cancer vaccines without DC activators may actually convey a tolerizing signal to the immune system and diminish endogenous immune response [137]. Thus, activation of DCs with immunostimulatory adjuvants is a critical component of many cancer vaccine strategies [136].

An advance in the design of cancer vaccines has been the inclusion of adjuvants that can trigger pattern recognition receptors, such as TLRs, Nod-like receptors (NLRs), retinoic acid-inducible gene (RIG)-I-like receptors (RLRs), stimulator of interferon genes (STING), and CD40 agonists, to signal the immune system that the vaccine antigen is both foreign and dangerous [2, 6, 11, 138]. This is especially important for peptide vaccine platforms, because unlike microbe- or nucleic acid-based platforms, peptide antigens do not inherently present danger signals to the immune system [11]. Without this signal, the immune system cannot mount a strong anti-tumor response due to a lack of costimulation and efficient antigen presentation by DCs.

Novel adjuvants, such as TLR agonists, have been tested in preclinical and clinical studies [11], with evidence of potent DC activation and generation of strong T cell responses [139, 140]. An important finding is that co-delivery of the peptide antigen and TLR agonist, either through peptide-agonist conjugation or nanocarriers, results in improved DC targeting, DC activation, and trafficking to draining lymph nodes [141, 142]. Recent peptide vaccine formulations have primarily employed Montanide ISA-51, TLR agonists such as poly I:C or CpG, or immunostimulatory cytokines such as GM-CSF as adjuvants [117]; however, there is currently no consensus about what the optimal adjuvant is for a given peptide vaccine [117], representing a potentially fruitful avenue of research to further optimize vaccine design.

## 4 Future Perspective

The historical experience with therapeutic cancer vaccines coupled with fundamental advances in understanding of the immunobiology of cancer have provided a road map for future vaccine development. The key challenges that must be overcome are identifying antigens and vaccine vectors that will lead to strong and broad T cell responses, tailoring vaccine designs to achieve optimal antigen presentation by professional APCs, and finding combination partners employing complementary mechanisms of action to overcome the diverse methods that cancer cells use to evade and suppress the immune system [11]. In recent years, the field has risen to meet this challenge, with many encouraging upgrades to antigen selection and vaccine designs. Combination strategies with a variety of other agents, including immunotherapies, chemotherapies, and radiotherapy, have also been investigated in preclinical and clinical studies [11]. These refinements will need to be validated in appropriately designed, randomized, phase 3 studies. Consequently, despite decades of lackluster progress, therapeutic cancer vaccines are now primed to emerge as central components of cancer therapy due to these advancements in biology and technology.

Therapeutic cancer vaccines may fill a niche not currently met by conventional therapies or other immunotherapies. Clinical experience suggests that vaccines are safe and can elicit long-term immune memory responses important for durable disease control [11]. This experience coupled with the existence of multiple mechanisms of immunosuppression in advanced disease suggest that vaccines may be particularly well-suited early in the course of the disease or in the minimal residual disease setting. Indeed, when there has been apparent benefit from cancer vaccines, it has been in the minimal residual disease setting [143].

Another potential role for vaccination is to augment lost immunity to oncogenic proteins where immunity is lost through oncogenesis [144, 145]. Cancer vaccines could also potentially be used to prevent disease through the targeting of antigens whose upregulation is associated with resistance to therapy [146].

Therapeutic cancer vaccines may also help actualize the full potential of immunotherapies that have already had an impact in the clinic [12]. For example, it is known that although checkpoint inhibitors are effective for “hot” tumors characterized by infiltration of primed and active T cells, they lack potency for “cold” tumors that do not have these immune cells. By priming tumor-specific T cells and mobilizing them to the tumor, essentially turning “cold” tumors “hot,” vaccine therapies restore the ability of checkpoint inhibitors to unleash T cell-mediated tumor

destruction [12, 128]. Similarly, whereas chimeric antigen receptor (CAR)-T cell therapy is now an established therapy for certain hematologic malignancies, demonstrating efficacy in solid tumors has been difficult. A recent preclinical study showed that injection of a vaccine consisting of amphiphile CAR-T ligands primed CAR-T cells and enhanced their efficacy in solid tumor models [147].

There are several encouraging avenues to further improve the efficacy of therapeutic cancer vaccines. One currently being investigated is the use of heterologous prime boosting whereby a TAA is first delivered by a specific vector during a priming vaccination and then subsequently delivered by a different vector during later boosting vaccinations [11]. This strategy overcomes immune-mediated inactivation of the initial viral vector, allowing repeated vaccination against the TAA target and potentially enhanced immunogenicity [11]. Another approach to improve the efficacy of cancer vaccines is the development of strategies to induce antibody responses with anti-tumor activity. For example, one study found that multi-site injections enhanced the number of TAA-specific antibodies compared with single bolus injections [148].

Finally, it is anticipated that cancer vaccines will benefit from advances in the speed, cost, and efficiency of molecular sequencing, artificial intelligence, and cellular engineering. These techniques may enable the quick and complete interrogation of the immune response (changes in immune milieu, tumor immune escape mechanisms) to a cancer vaccine, allowing subsequent vaccines to be tailored based on this response [149].

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revised it critically for important intellectual content; approved the final version; and agree to be accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved.

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