

# A combination trial of vaccine plus ipilimumab in metastatic castration-resistant prostate cancer patients: immune correlates

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**Abstract** We recently reported the clinical results of a Phase I trial combining ipilimumab with a vaccine containing transgenes for prostate-specific antigen (PSA) and for a triad of costimulatory molecules (PROSTVAC) in patients with metastatic castration-resistant prostate cancer. Thirty patients were treated with escalating ipilimumab and a fixed dose of vaccine. Of 24 chemotherapy-naïve patients, 58 % had a PSA decline. Combination therapy did not exacerbate the immune-related adverse events associated with ipilimumab. Here, we present updated survival data and an evaluation of 36 immune cell subsets pre- and post-therapy. Peripheral blood mononuclear cells were collected before therapy, at 13 days and at 70 days post-initiation of therapy, and phenotyped by flow cytometry for the subsets of T cells, regulatory T cells, natural killer cells, and myeloid-derived suppressor cells. Associations between overall survival (OS) and immune cell subsets prior to treatment, and the change in a

given immune cell subset 70 days post-initiation of therapy, were evaluated. The median OS was 2.63 years (1.77–3.45). There were trends toward associations for longer OS and certain immune cell subsets before immunotherapy: lower PD-1<sup>+</sup>Tim-3<sup>NEG</sup>CD4<sub>EM</sub> ( $P = 0.005$ , adjusted  $P = 0.010$ ), higher PD-1<sup>NEG</sup>Tim-3<sup>+</sup>CD8 ( $P = 0.002$ , adjusted  $P = 0.004$ ), and a higher number of CTLA-4<sup>NEG</sup> Tregs ( $P = 0.005$ , adjusted  $P = 0.010$ ). We also found that an increase in Tim-3<sup>+</sup> natural killer cells post- versus pre-vaccination associated with longer OS ( $P = 0.0074$ , adjusted  $P = 0.015$ ). These results should be considered as hypothesis generating and should be further evaluated in larger immunotherapy trials.

**Keywords** Ipilimumab · Vaccine · PROSTVAC · T cells · NK cells · Immunotherapy

## Abbreviations

ALC	Absolute lymphocyte count
CTLA-4	Cytotoxic T-lymphocyte-associated antigen-4
DT	Doubling time
EM	Effector memory
GM-CSF	Granulocyte–macrophage colony-stimulating factor
ICOS	Inducible costimulator
IFN	Interferon
IL	Interleukin
mCRPC	Metastatic castration-resistant prostate cancer
MDSC	Myeloid-derived suppressor cell
NK	Natural killer
OS	Overall survival
PAP	Prostatic acid phosphatase
PBMC	Peripheral blood mononuclear cell
PD-1	Programmed death 1 receptor
PSA	Prostate-specific antigen
PSMA	Prostate-specific membrane antigen

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TIM-3	T-cell immunoglobulin and mucin domain-containing molecule-3
Tregs	Regulatory T cells
TRICOM	Triad of costimulatory molecules (ICAM-1, B7.1, and LFA-3)

## Introduction

Two immunotherapeutic agents for cancer have recently been approved by the Food and Drug Administration (FDA): sipuleucel-T for prostate cancer and ipilimumab for metastatic melanoma. Sipuleucel-T (PROVENGE<sup>®</sup>, Dendreon Corp.) is a therapeutic vaccine generated by the isolation of the patient's peripheral blood mononuclear cells (PBMCs) and culturing them in vitro with a fusion protein of prostatic acid phosphatase (PAP) and granulocyte-macrophage colony-stimulating factor (GM-CSF). The product is then reinfused into the patient. The Phase III IMPACT trial showed a 4.1-month improvement in overall survival (OS) and a 22 % relative reduction in risk of death, and it was approved for use in prostate cancer in 2010 [1].

Another vaccine, designated as PROSTVAC, has shown evidence of clinical activity in metastatic prostate cancer in two Phase II trials [2, 3], and a Phase III trial is currently ongoing (NCT01322490 [4]). PROSTVAC (PSA-TRICOM, Bavarian Nordic, Inc.) consists of a prime-boost regimen with recombinant vaccinia (prime) and fowlpox (boost) vectors, containing transgenes for prostate-specific antigen (PSA) and three costimulatory molecules for cytotoxic T lymphocytes (B7.1, ICAM-1, and LFA-3, designated TRICOM) [5]. A multicenter, randomized, placebo-controlled Phase II study showed an 8.5-month improvement in overall survival and a 44 % reduction in death rate compared to placebo in patients with asymptomatic or minimally symptomatic metastatic castration-resistant prostate cancer (mCRPC) [2]. The median OS was 25.1 months for vaccinated patients ( $n = 82$ ) versus 16.6 months for controls ( $n = 40$ ). In a second Phase II single-arm trial in mCRPC at the National Cancer Institute (NCI), the median survival was 26.6 months ( $n = 32$ ) [3]. A retrospective analysis of this trial evaluated patients based on the Halabi nomogram [6] and found that patients with a more indolent disease (predicted survival >18 months) displayed greater improvements in survival than patients with more aggressive disease. PROSTVAC vaccination was also shown to generate an antigen-specific immune response [3]. In addition, it was shown that patients who had a decrease in the cytotoxic T-lymphocyte-associated antigen-4 (CTLA-4)<sup>+</sup> regulatory T-cell (Treg) population post-vaccination displayed longer overall survival [7].

Ipilimumab (Yervoy<sup>®</sup>, Bristol-Myers Squibb) is a fully human monoclonal antibody that targets CTLA-4. It is the

first in a new class of agents called immune checkpoint inhibitors. It has been extensively studied in metastatic melanoma and has shown an improvement of overall survival of 2–4 months compared to active control groups, which led to FDA approval [8, 9]. Ipilimumab has also previously been investigated for the treatment for prostate cancer in a pilot trial of patients with hormone-refractory prostate cancer [10]. They found a PSA decline of  $\geq 50$  % in 2/14 patients and concluded that further investigations were warranted. Ipilimumab alone or in combination with radiotherapy was also investigated in a recently reported Phase I/II trial of 75 patients with mCRPC [11]. Both PSA decline and tumor response were observed, and 8/34 patients in the 10 mg/kg  $\pm$  radiotherapy group had a confirmed PSA decline of  $\geq 50$  %. Of these, six had received prior chemotherapy and two were chemotherapy-naïve. One of the tumor-evaluable patients in the 10 mg/kg  $\pm$  radiotherapy group achieved a confirmed complete response, and 2 patients achieved an unconfirmed partial response. Six patients had stable disease. The median OS was 17.4 months [11].

In the PSA-TRICOM trials, no adverse events above grade 1 or 2 toxicity and no evidence of autoimmunity were observed. In the ipilimumab trials, there were some severe adverse events involving autoimmunity, including colitis, panhypophysitis, adrenal insufficiency, raised aminotransferases, and neutropenia. Since PSA-TRICOM has three costimulatory molecules designed to enhance T-cell immunity, and the ipilimumab checkpoint inhibitor is designed to reduce the immune suppressive CTLA-4 entity, it was important to determine whether the combination of PSA-TRICOM and ipilimumab would exacerbate the autoimmunity seen with ipilimumab alone.

We have recently reported [12] the clinical results of a Phase I study combining ipilimumab with PSA-TRICOM vaccine in patients with mCRPC. Thirty patients were treated with an escalating dose of ipilimumab and a fixed dose of vaccine. Of the 24 chemotherapy-naïve patients, 14 patients (58 %) had a PSA decline from baseline, with six of these  $\geq 50$  %. Combination therapy did not seem to exacerbate the immune-related adverse events associated with ipilimumab, and there was no apparent association between immune-related adverse events and clinical outcome. In the present study, we report on updated survival data, which was evaluated in terms of several patient characteristics such as Gleason score and Halabi nomogram. Here, we have also investigated whether any of 36 specific immune cell subsets of patients prior to therapy correlate with clinical outcome and whether changes in any of these subsets during therapy correlate with survival. For each of these immune cell subsets, we have analyzed phenotypes based on known immunologic markers, many of which have previously been shown to correlate with biologic activity [7, 13–19].

## Materials and methods

### Patients

Thirty patients with mCRPC were enrolled on a Phase I trial of combination therapy with ipilimumab and PROST-VAC, a poxviral vaccine targeting PSA and containing transgenes for three T-cell costimulatory molecules (NCT00113984) [12, 20]. Recombinant vaccinia PROST-VAC was given as a prime with recombinant fowlpox PROSTVAC given as monthly boosts starting on day 15. GM-CSF was given on 4 consecutive days with each vaccination. Ipilimumab was given at the dose levels of 1, 3, 5, and 10 mg/kg. Ipilimumab treatment was started after 2 weeks, at the time of the first boost vaccination, and given monthly on the same day as vaccine. Initially, our protocol allowed for only six courses with ipilimumab; however, a protocol amendment gave patients with stable disease the option of additional ipilimumab every 3 months for a maximum of four additional doses. The maintenance dose of monthly vaccine could continue until there was evidence of disease progression on imaging studies, or toxic effects that required discontinuation. All injections were given at the NIH Clinical Center (Bethesda, MD, USA). All patients reviewed and signed an informed consent form approved by the NCI's Institutional Review Board.

### Collection of peripheral blood mononuclear cells

Peripheral blood mononuclear cells were collected at baseline, after 13 days and after approximately 70 days of treatment. Briefly, 60 ml of blood was collected, and the mononuclear fraction was separated by Ficoll–Hypaque density gradient separation, washed three times, and preserved in 90 % heat-inactivated human AB serum (Gemini Bio-Products, W Sacramento, CA, USA) and 10 % DMSO in liquid nitrogen at a concentration of  $1 \times 10^7$  cells/ml until assayed.

### Flow cytometry

Multi-color flow cytometry analysis was performed on PBMCs from all time points by staining for 30 min at 4 °C with CD3-V450, CD8-FITC or APC, ICOS-PE, HLA-DR-PerCP-Cy5.5, CD25-PE-Cy7, CD45RA-PerCP-Cy5.5, CD62L-FITC, CD127-V450, PD-1-PE, Tim-3-AF700, CD4-APC-Cy7 (BD Biosciences, San Jose, CA, USA), CCR7-PE-Cy7 (R&D Systems, Minneapolis, MN, USA), CTLA-4-FITC (LSBio, Seattle, WA, USA), and FoxP3-APC (eBioscience, San Diego, CA, USA) for T cells. For natural killer (NK) cells, CD3-V450, CD16-APC-Cy7, CD56-PE-Cy7, and Tim-3-AF700 (BD) were used. For myeloid-derived suppressor cells (MDSCs), CD33-PE,

CD11b-APC-Cy7, HLA-DR-PerCP-Cy5.5, CD14-V450, and CD15-APC (BD) were used.  $1 \times 10^5$  cells were acquired on an LSRII (BD), and data were analyzed using FlowJo software (Tree Star Inc., Ashland, OR, USA). The appropriate isotype controls were used, and dead cells were excluded from the analysis.

### Induction and analysis of T<sub>H</sub>17 cells

T<sub>H</sub>17 cells were analyzed using the Human T<sub>H</sub>1/T<sub>H</sub>17 Phenotyping kit (BD). Briefly, PBMCs were thawed and incubated overnight at 37 °C.  $1 \times 10^6$  cells/ml were stimulated for 5 h with PMA/Ionomycin in the presence of Golgi-Stop (Leukocyte Activation Cocktail with BD GolgiPlug, BD). The cells were then fixed, permeabilized, and stained according to the manufacturer's instructions. CD4-PerCP-Cy5.5, interleukin (IL)-17A-PE, and interferon (IFN) $\gamma$ -FITC (BD) were used.  $1 \times 10^5$  cells were acquired on an LSRII (BD), and data were analyzed using FACSDiva software (BD). The appropriate isotype controls were used, and dead cells were excluded from the analysis.

### Statistical analysis

In an exploratory manner, an actuarial analysis was performed on overall survival using the Kaplan–Meier method. OS was calculated as the period between the on-study date and date of death, or last follow-up. The log-rank test was used to compare strata or test for a trend (where appropriate). For both immune cell parameters and clinical parameters, baseline values were used to create strata for use in the actuarial analysis. For immune cell parameters, the percent difference from baseline (day 70–day 0) data was also used to create strata. The cutoffs were selected post hoc. Based upon the number of subjects available, the data were divided in tertiles to perform an exploratory evaluation of the association between the parameters and OS. For those parameters in which the log-rank  $P < 0.10$ , adjacent strata were combined and the two new strata with the smallest  $P$  value were used (in which case the log-rank test  $P$  value was adjusted for the implicit number of tests performed). Subsequently, a Cox proportional hazards regression analysis was performed on the data. The initial regression model included parameters from the actuarial analysis such that the log-rank  $P < 0.05$ . Both stepwise and backward selection processes were performed on the data.

Either a parametric or nonparametric analysis was performed on the immunological data, as appropriate. A repeated measures analysis of variance (ANOVA) was performed on the data if the ANOVA assumptions were satisfied. A Box–Cox transformation was performed on the data prior to ANOVA, and the data were transformed as appropriate. We also tested for linear and curvilinear

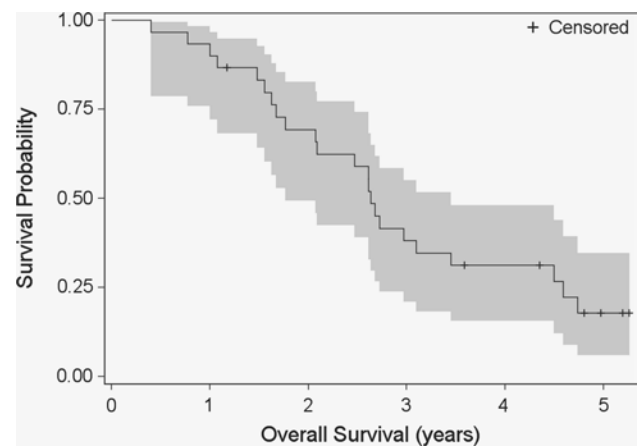
trends over time using orthogonal polynomial contrasts. Residuals were examined for normality to verify ANOVA assumptions. If ANOVA was not appropriate for the data, we first used Friedman's test and then used the Wilcoxon signed rank test to make pairwise comparisons between distributions of time periods. For both methods, all three pairwise comparisons were made and the  $P$  values were adjusted using Holm's method (step down Bonferroni). In view of the very large number of tests performed on the survival and immunological data, we consider  $P < 0.005$  as being statistically significant, while  $0.005 < P < 0.05$  would be considered trends. All reported  $P$  values are two-tailed.

## Results

Overall survival of all 30 patients has been updated from the previous publication [12] and was calculated as the difference between the on-study date and the date of death ( $n = 23$ ), or the date of last follow-up ( $n = 7$ ). Figure 1 shows the Kaplan–Meier plot for overall survival for all patients. The median survival time was 2.63 years (95 % confidence limits 1.77–3.45). Probability of survival (95 % confidence limits) at 1, 2, and 3 years was 0.93 (0.76–0.98), 0.69 (0.49–0.83), and 0.38 (0.21–0.55), respectively.

We performed an actuarial analysis of overall survival on the clinical characteristics data. As can be seen in Table 1, there were trends favoring a low Halabi score  $<117$ , which corresponds to a Halabi predicted survival of greater than approximately 18 months, a longer PSA-doubling time (DT) at baseline  $>2.42$  months, and a baseline hemoglobin  $>12.4$  g/dl. These results have previously been reported to be prognostic favorable factors [6, 21]. No other clinical variables were found to associate significantly with overall survival.

Using seven-color flow cytometry, we have now evaluated the subsets of CD4, CD8, NK, Tregs,  $T_H17$  cells, and MDSC at three time points: pre-treatment, day 13 (post-first vaccine and pre-ipilimumab), and day 70 (during vaccine/ipilimumab treatment). For each of these immune cell subsets, we have analyzed phenotypes based on known immunologic markers, some of which have previously been shown to correlate with a specific biologic activity [7, 13–19]. The description of each of these 36 subsets is given in Table 2. Figure 2 shows the three immune cell subsets that increased during therapy. For the three parameters shown, the differences between baseline (BL) and day 70, and day 13 and day 70, were generally significantly larger than zero, that is, the day 70 values were significantly larger than the baseline and day 13 values. Linear trends tests confirmed these findings for absolute lymphocyte count (ALC) ( $P < 0.0001$ ) (Fig. 2A), ICOS<sup>+</sup> CD4<sup>+</sup> T cells ( $P < 0.0080$ )



**Fig. 1** Overall survival. Kaplan–Meier curve for overall survival in years for all patients ( $n = 30$ ), calculated as the difference between the on-study date and the date of death ( $n = 23$ ), or the date of last follow-up ( $n = 7$ ). The median survival time was 2.63 years (95 % confidence limits 1.77–3.45). Probability of survival (95 % confidence limits) at 1, 2, and 3 years was 0.93 (0.76–0.98), 0.69 (0.49–0.83), and 0.38 (0.21–0.55), respectively

**Table 1** Risk analysis for clinical parameters versus overall survival

	Log rank	Trend	Favored
Halabi score	0.070	0.063	$<117$
Gleason score	0.024	0.42	None
Baseline PSA	0.28	0.69	None
Off study PSA	0.46	0.61	None
PSA–DT (months)	0.023	0.015	$>2.42$
Baseline hemoglobin	0.061	0.029	$>12.4$

Actuarial analysis results after dichotomizing the data for clinical characteristics showing the log-rank and trend test  $P$  values, as well as the favored group

PSA–DT prostate-specific antigen-doubling time

(Fig. 2B), and IFN $\gamma$ <sup>+</sup> CD4<sup>+</sup> T cells ( $P < 0.0006$ ) (Fig. 2C). The significance of these immune cell subsets will be further discussed. All other studied immune cell subsets shown in Table 2 did not change significantly from baseline to day 13 or day 70 of treatment, or from day 13 to 70.

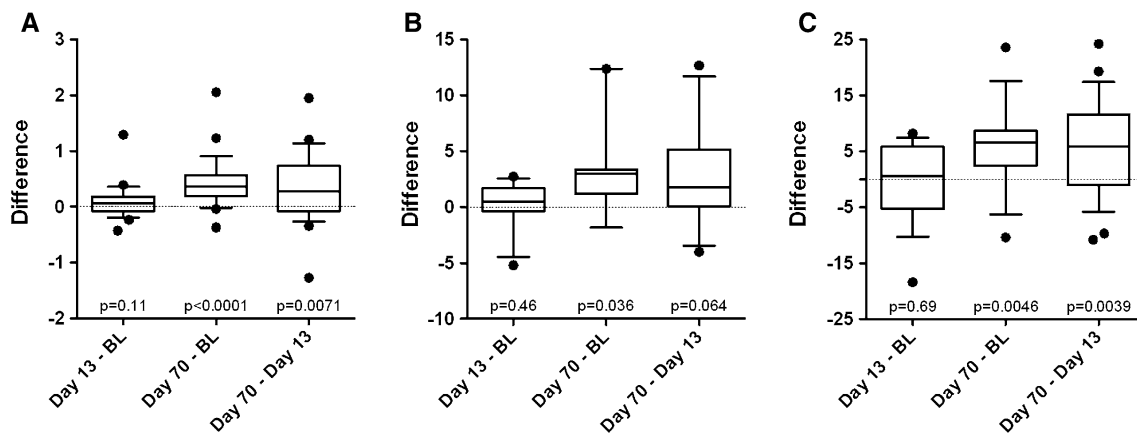
Analyses of clinical and immune cell subset baseline values were performed, as well as differences from baseline of the immune cell subsets, to evaluate whether any association existed with subsequent overall survival (Table 3). PBMCs were not available for flow cytometry analysis for 2 out of the 30 patients, so they were excluded from these comparisons.

Actuarial analyses were performed to identify immune cell subsets that were associated with longer OS. Subsequently, Cox regression analyses were performed on the immune cell subsets showing evidence for being associated

**Table 2** Immune cell subsets

Immune cell	Meaning
ALC	Absolute lymphocyte count
CD4	T helper cells
CD4 <sub>CM</sub>	Central memory T helper cells (CD4 <sup>+</sup> CD45RA <sup>NEG</sup> CD62L <sup>+</sup> CCR7 <sup>+</sup> )
PD-1 <sup>+</sup> TIM-3 <sup>NEG</sup> CD4 <sub>CM</sub>	Activated central memory CD4 T cells, negative immune regulator
PD-1 <sup>NEG</sup> TIM-3 <sup>+</sup> CD4 <sub>CM</sub>	Activated central memory CD4 T cells, negative immune regulator
PD-1 <sup>+</sup> TIM-3 <sup>+</sup> CD4 <sub>CM</sub>	Severe exhaustion of central memory CD4 T cells
CD4 <sub>EM</sub>	Effector memory T helper cells (CD4 <sup>+</sup> CD45RA <sup>NEG</sup> CD62L <sup>NEG</sup> CCR7 <sup>NEG</sup> )
<b>PD-1<sup>+</sup> TIM-3<sup>NEG</sup> CD4<sub>EM</sub></b>	<b>Activated effector memory CD4 T cells, negative immune regulator</b>
PD-1 <sup>NEG</sup> TIM-3 <sup>+</sup> CD4 <sub>EM</sub>	Exhausted effector memory CD4 T cells, negative immune regulator
PD-1 <sup>+</sup> TIM-3 <sup>+</sup> CD4 <sub>EM</sub>	Severe exhaustion of effector memory CD4 T cells
ICOS <sup>+</sup> CD4	Activated T helper cells
IFN $\gamma$ <sup>+</sup> CD4	Activated T helper cells
T <sub>H</sub> 17	T helper cell type 17, unclear role in cancer
IFN $\gamma$ <sup>+</sup> T <sub>H</sub> 17	More activated T helper cell type 17
CD8	Cytotoxic T cells
<b>PD-1<sup>NEG</sup> TIM-3<sup>+</sup> CD8</b>	<b>Activated CD8 T cells, negative immune regulator</b>
CD8 <sub>CM</sub>	Central memory cytotoxic T cells (CD8 <sup>+</sup> CD45RA <sup>NEG</sup> CD62L <sup>+</sup> CCR7 <sup>+</sup> )
PD-1 <sup>+</sup> TIM-3 <sup>NEG</sup> CD8 <sub>CM</sub>	Activated central memory CD8 T cells, negative immune regulator
PD-1 <sup>NEG</sup> TIM-3 <sup>+</sup> CD8 <sub>CM</sub>	Activated central memory CD8 T cells, negative immune regulator
PD-1 <sup>+</sup> TIM-3 <sup>+</sup> CD8 <sub>CM</sub>	Severe exhaustion of central memory cytotoxic T cells
CD8 <sub>EM</sub>	Effector memory CD8 T cells (CD8 <sup>+</sup> CD45RA <sup>NEG</sup> CD62L <sup>NEG</sup> CCR7 <sup>NEG</sup> )
PD-1 <sup>+</sup> TIM-3 <sup>NEG</sup> CD8 <sub>EM</sub>	Activated effector memory CD8 T cells, negative immune regulator
PD-1 <sup>NEG</sup> TIM-3 <sup>+</sup> CD8 <sub>EM</sub>	Activated effector memory CD8 T cells, negative immune regulator
PD-1 <sup>+</sup> TIM-3 <sup>+</sup> CD8 <sub>EM</sub>	Severe exhaustion of effector memory cytotoxic T cells
ICOS <sup>+</sup> CD8	Activated cytotoxic T cells
<b>TREGS</b>	<b>Regulatory T cells (CD4<sup>+</sup>CD25<sup>HI</sup>FoxP3<sup>+</sup>CD127<sup>NEG</sup>)</b>
CD4 : TREG RATIO	Ratio of effector T cells to regulatory T cells
CD8 : TREG RATIO	Ratio of effector T cells to regulatory T cells
NK CELLS	Natural killer cells (CD3 <sup>NEG</sup> CD56 <sup>+</sup> )
CD16 <sup>+</sup> CD56 <sup>BR</sup>	Functional intermediate, lytic, and cytokine production
TIM-3 <sup>+</sup> CD16 <sup>+</sup> CD56 <sup>BR</sup>	Fully functional intermediate
CD16 <sup>+</sup> CD56 <sup>DIM</sup>	Mature, more cytokine production
TIM-3 <sup>+</sup> CD16 <sup>+</sup> CD56 <sup>DIM</sup>	Fully functional mature
CD16 <sup>NEG</sup> CD56 <sup>BR</sup>	Immature, more lytic
<b>TIM-3<sup>+</sup> CD16<sup>NEG</sup>CD56<sup>BR</sup></b>	<b>Immature NK cells transitioning into more mature (CD16<sup>+</sup>CD56<sup>DIM</sup>) phenotype</b>
MDSC	Myeloid-derived suppressor cells (HLA-DR <sup>NEG</sup> CD33 <sup>+</sup> CD11b <sup>+</sup> )
MDSC <sub>MO</sub>	Monocytic MDSC (CD14 <sup>+</sup> CD15 <sup>NEG</sup> )
MDSC <sub>GR</sub>	Granulocytic MDSC (CD14 <sup>NEG</sup> CD15 <sup>+</sup> )
MDSC <sub>LIN-</sub>	Non-lineage MDSC (CD14 <sup>NEG</sup> CD15 <sup>NEG</sup> )
MARKERS	
CTLA-4	Cytotoxic T-lymphocyte-associated antigen-4
ICOS	Inducible costimulator Costimulatory for the activation of T cells
PD-1	Programmed death 1 receptor On activated T cells and B cells, and on mature dendritic cells. Negative immune regulator, engagement with PD-L1, can downregulate T-cell activation.
TIM-3	T-cell immunoglobulin and mucin domain-containing molecule-3 Activation and maturation marker, and negative regulator of NK cells. Negative immune regulator expressed on T cells.

These 36 different immune cell subsets were analyzed by flow cytometry at baseline, day 13 and day 70, as described in “[Materials and methods](#).” Subsets shown in **bold** were associated with overall survival in subsequent analyses. One parameter, PD-1<sup>+</sup> TIM-3<sup>NEG</sup> CD8<sub>CM</sub>, was not analyzed because many values were zero



**Fig. 2** Analysis of immune cell subsets pre-therapy and during therapy. Thirty patients with metastatic castration-resistant prostate cancer were treated with an increasing dose (1, 3, 5, or 10 mg/kg) of ipilimumab in combination with PROSTVAC vaccine and GM-CSF. Peripheral blood mononuclear cells (PBMCs) were collected at baseline, day 13, and day 70 and analyzed by flow cytometry. **a** There was a trend for increase in the absolute lymphocyte count (ALC) during therapy ( $n = 27\text{--}29$ ). **b** There was a trend for increase in the frequency of ICOS<sup>+</sup> CD4<sup>+</sup> T cells during therapy (baseline:  $n = 12$ ,

day 13:  $n = 16$ , day 70:  $n = 14$ ). **c** There was a trend for increase in the frequency of IFN $\gamma$ <sup>+</sup> CD4<sup>+</sup> T cells during therapy ( $n = 19\text{--}29$ ). For the three parameters shown, the differences between baseline (BL) and day 70, and day 13 and day 70, were generally significantly larger than zero, that is, the day 70 values were significantly larger than the baseline and day 13 values. Linear trends tests confirmed these findings. The box plots display the differences for each pairwise comparison between the time points, from baseline to day 13 or day 70 of treatment, or from day 13 to day 70. BL baseline

**Table 3** Cox model results for specific immune cell subsets versus overall survival

Parameter <sup>a</sup>	Parameter levels <sup>b</sup>	<i>P</i> value	Hazard ratio	95 % confidence limits
Baseline % PD1 <sup>+</sup> of CD4 <sub>EM</sub>	<4.77 versus >4.77	0.026	3.17	1.15–8.75
Baseline number of Tregs	<98 versus >98	0.0013	0.153	0.049–0.481
Baseline % Tim-3 <sup>+</sup> of CD8	<40.9 versus >40.9	0.002	0.155	0.047–0.505
% PD-1 <sup>+</sup> of CD8 <sub>EM</sub>	<50 versus >50	0.027	3.03	1.33–8.07
% Tim-3 <sup>+</sup> of CD16 <sup>NEG</sup> CD56 <sup>BR</sup>	<57.8 versus >57.8	0.055	0.283	0.078–1.029
Baseline hemoglobin	<12.4 versus >12.4	0.011	0.271	0.100–0.739

<sup>a</sup> The first three laboratory parameters are baseline values and were included in an analysis of clinical and immune cell subset baseline values, as well as differences from baseline of the immune cell subsets. The other laboratory parameters are percent differences from baseline, and this analysis was based solely on the differences between baseline and the clinical baseline variables

<sup>b</sup> The first level indicated is the reference level

with longer OS. Three immune cell subsets at baseline were found to be jointly predictive of OS, but are presented in a univariate manner. As seen in Fig. 3a, patients with a lower percentage of PD-1<sup>+</sup>Tim-3<sup>NEG</sup> CD4 effector memory (CD4<sub>EM</sub>) cells at baseline (<4.77 %) displayed longer survival ( $P = 0.005$ , adjusted  $P = 0.010$ ). As shown in Fig. 3b, patients with a higher percentage (>40.9 %) of activated Tim-3 single positive (Tim-3<sup>+</sup>PD-1<sup>NEG</sup>) CD8<sup>+</sup> T lymphocytes at baseline displayed longer overall survival ( $P = 0.002$ , adjusted  $P = 0.004$ ). Surprisingly, an increased number of Tregs at baseline also correlated with longer OS ( $P = 0.005$ , adjusted  $P = 0.010$ ) (Fig. 3c); however, it should be noted that these were the CTLA-4<sup>NEG</sup> Tregs (CD4<sup>+</sup> CD25<sup>HI</sup> FoxP3<sup>+</sup> CD127<sup>NEG</sup> CTLA-4<sup>NEG</sup>), and not the CTLA-4<sup>+</sup> Tregs, which have previously been found to

be the most highly suppressive subset [7]. These immune cell subsets will be further discussed. The Cox model did not identify any other immune cell subsets as being predictive of OS.

We then evaluated whether any association existed between overall survival and the change in a specific immune cell subset at 70 days of therapy, compared to the baseline level. This analysis was based solely on the change in immune cell subsets and the clinical baseline variables. Adjusting for the baseline level of hemoglobin, two other immune cell subsets were associated with OS (Table 3). There was a trend toward an increased OS in patients who had a larger increase in the population of immature NK cells transitioning into a more mature NK phenotype (TIM-3<sup>+</sup> CD3<sup>NEG</sup> CD16<sup>NEG</sup> CD56<sup>BR</sup>) versus

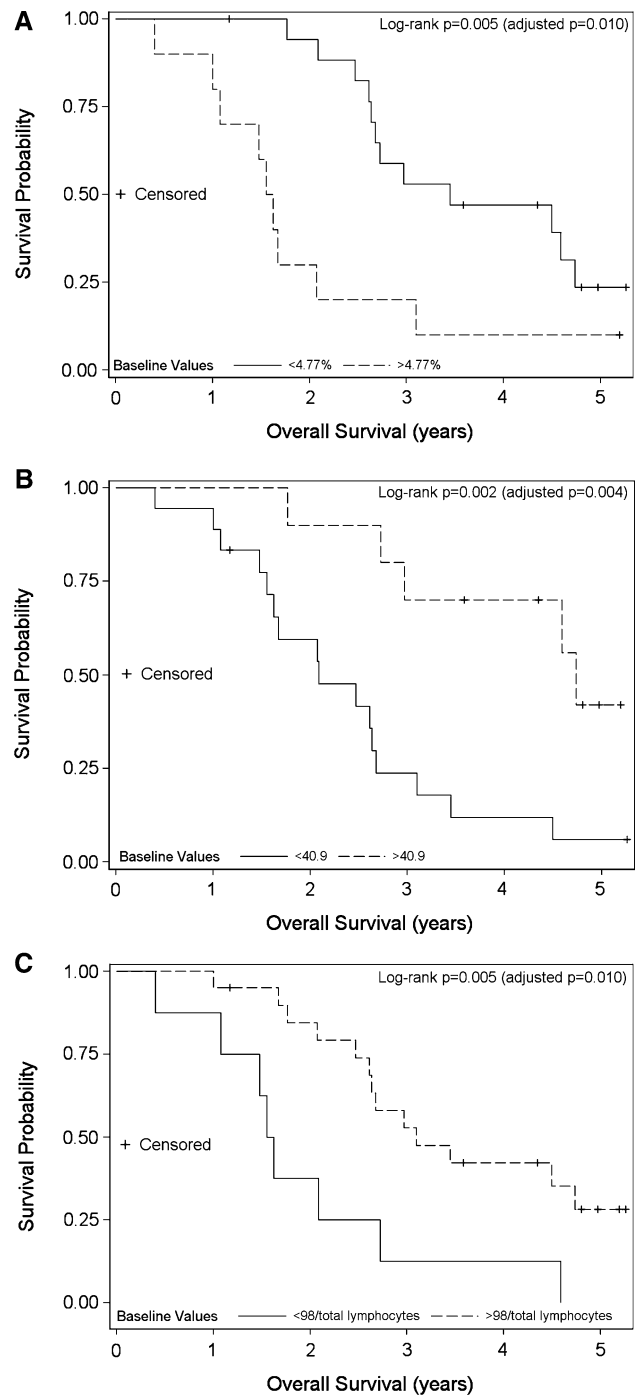
**Fig. 3** Analysis of immune cell subsets at baseline versus overall survival. Thirty patients with metastatic castration-resistant prostate cancer were treated with an increasing dose (1, 3, 5, or 10 mg/kg) of ipilimumab in combination with PROSTVAC vaccine and GM-CSF. Peripheral blood mononuclear cells from baseline and day 70 were available for 28 patients, and were analyzed by flow cytometry. Kaplan–Meier curves representing overall survival versus immune cell subsets after dichotomizing the data at one of the tertiles are shown. **a** Baseline % PD-1<sup>+</sup>Tim-3<sup>NEG</sup> activated effector memory CD4<sup>+</sup> T cells. *Dashed lines* denote overall survival of patients with immune cell subset values greater than the upper tertile. *Solid lines* denote overall survival of patients with immune cell subset values below the upper tertile. **b** Baseline % Tim-3<sup>+</sup>PD-1<sup>NEG</sup> activated CD8<sup>+</sup> T cells. *Dashed lines* denote overall survival of patients with immune cell subset values greater than the upper tertile. *Solid lines* denote overall survival of patients with immune cell subset values below the upper tertile. **c** Baseline number of CTLA-4<sup>NEG</sup> Tregs (CD4<sup>+</sup> CD25<sup>HI</sup> FoxP3<sup>+</sup> CD127<sup>NEG</sup> CTLA-4<sup>NEG</sup>). *Dashed lines* denote overall survival of patients with immune cell subset values greater than the lower tertile. *Solid lines* denote overall survival of patients with immune cell subset values below the lower tertile

those patients who had a smaller increase or a decrease in this NK-cell subset ( $P = 0.0074$ , adjusted  $P = 0.015$ ) (Fig. 4a), as well as a trend toward an increased OS in patients who had a decrease or less than 50 % increase in the percentage of PD-1<sup>+</sup> Tim-3<sup>NEG</sup> CD8<sub>EM</sub> T cells versus those patients who had an increase >50 % in this subset on day 70 ( $P = 0.0062$ , adjusted  $P = 0.012$ ) (Fig. 4b). No additional associations with OS were observed for the other immune cell subsets studied.

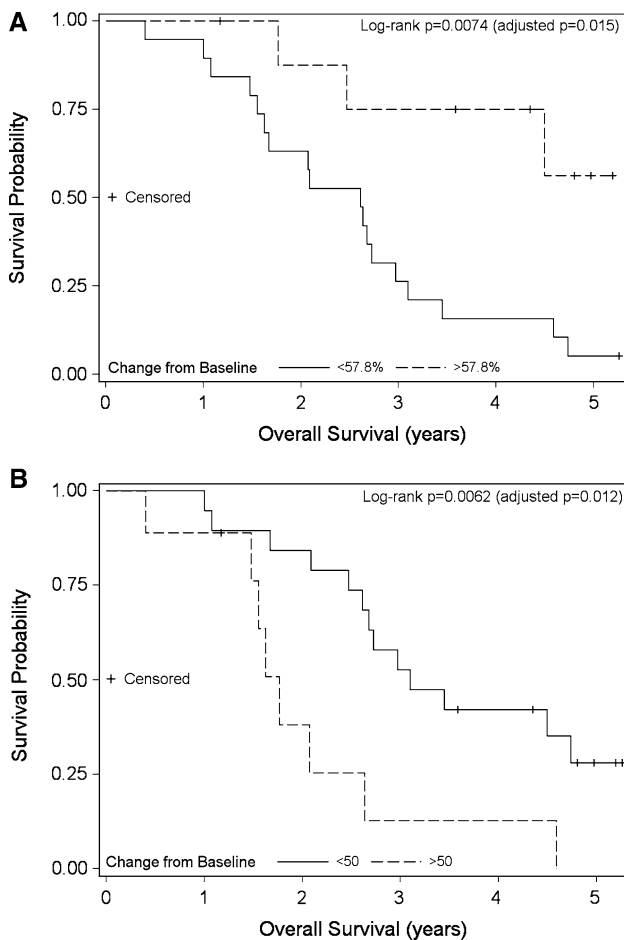
## Discussion

The study reported here provides an update on survival data from a Phase I trial combining therapy with a viral vector vaccine, PROSTVAC, and ipilimumab, an immune checkpoint inhibitor [12], as well as an evaluation of 36 discrete immune cell subsets in peripheral blood before and during this therapy.

The rationale for employing immunotherapy in prostate cancer is that previous studies have shown the presence of tumor-infiltrating lymphocytes in prostate cancer tissue, even when no therapy has been given previously [22, 23], and increased infiltration after androgen deprivation therapy [24]. In one study, the magnitude and quality of the infiltrate was shown to be a prognostic factor for survival [22]. This suggests that an immune reaction can be mounted, but it is not strong enough to inhibit tumor growth. A recent phenotypic study has shown that skewing of the intraprostatic immune cell infiltrate toward the T<sub>H</sub>17 and Treg phenotypes may be involved in the development and progression of prostate cancer [25]. Immunotherapies provide several different strategies to increase the immune response by increasing the immune cell infiltrate,



by making the effector cells more proficient at killing the tumor cells, by decreasing immune suppressive entities such as Tregs and MDSC, or by changing the composition of the immune cell infiltrate, and thereby decreasing immunosuppression and overcoming immune tolerance. In addition, the presence of specific immune cell subsets prior to immunotherapy in some patients may render them more amenable to immunotherapy and vice versa. Prostate cancer provides a good model for immunotherapy since



**Fig. 4** Analysis of change in immune cell subsets versus overall survival. Thirty patients with metastatic castration-resistant prostate cancer were treated with an increasing dose (1, 3, 5, or 10 mg/kg) of ipilimumab in combination with PROSTVAC vaccine and GM-CSF. Peripheral blood mononuclear cells from baseline and day 70 were available for 28 patients and were analyzed by flow cytometry. Kaplan–Meier curves representing overall survival versus immune cell subsets after dichotomizing the data at one of the tertiles. **a** The change from baseline to day 70 in the percentage of immature natural killer cells transitioning into a more mature phenotype (TIM-3<sup>+</sup> CD16<sup>NEG</sup> CD56<sup>BR</sup>). *Dashed lines* denote overall survival of patients with immune cell subset changes greater than the upper tertile. *Solid lines* denote overall survival of patients with immune cell subset changes below the upper tertile. **b** The change from baseline to day 70 in the percentage of PD-1<sup>+</sup>Tim-3<sup>NEG</sup> CD8<sup>EM</sup> T cells (CD8<sup>+</sup> CD45RA<sup>NEG</sup> CD62L<sup>NEG</sup> CCR7<sup>NEG</sup>). *Dashed lines* denote overall survival of patients with an increase >50 % in this immune cell subset. *Solid lines* denote overall survival of patients with a decrease, or an increase <50 %, in this immune cell subset

there are several known tumor-associated antigens [PSA, prostate-specific membrane antigen (PSMA), and PAP, for example] that are minimally expressed in other organs, decreasing the risk of immune-related side effects.

It has previously been shown that incorporation of transgenes for a tumor antigen and a triad of costimulatory molecules into the vaccine enhances the quantity and the

quality of the CD8<sup>+</sup> T cells generated [26, 27]. In addition, another preclinical study showed that simultaneously providing positive costimulation and inhibiting negative costimulatory signals using anti-CTLA-4 monoclonal antibodies resulted in greatly enhanced (10-fold) avidity of the T cells [28]. Since PSA–TRICOM has three costimulatory molecules designed to enhance T-cell immunity, and the ipilimumab checkpoint inhibitor is designed to reduce the immune suppressive CTLA-4 entity, it was important to determine whether the combination of PSA–TRICOM and ipilimumab would exacerbate the autoimmunity seen with ipilimumab alone. However, no increase in the frequency or severity of immune-related adverse events above that observed with ipilimumab alone was seen. The most common toxic effect seen in this study was grade 1–2 vaccination site reactions (3 patients had grade 1 and 26 patients had grade 2). Twenty-one patients had immune-related adverse events of grade 2 or higher. These included grade 3–4 diarrhea or colitis (4 patients), grade 3 rash (2 patients), grade 3 raised aminotransferases (2 patients), grade 3 endocrine events (2 patients), and grade 4 neutropenia (1 patient) [12].

The updated median OS in the trial reported here was 2.63 years (31.6 months). The patient population was similar to that in the previous Phase II trial of PSA–TRICOM alone, where the median survival in the vaccine arm was 25.1 months versus 16.6 months in the control arm. The results compare quite favorably with the results of a Phase II study employing PROSTVAC alone in a similar population. There also appeared to be a greater serum PSA response in the chemotherapy-naïve patients in the combination study [12, 29]. A Phase III trial of ipilimumab with radiation in advanced metastatic prostate cancer did not show a statistical survival benefit, i.e., only a 1.2-month difference in OS versus the placebo arm [30]. A 3-arm randomized trial will need to be conducted comparing the efficacy of vaccine versus ipilimumab versus vaccine + ipilimumab.

In the current trial, there were trends for longer overall survival favoring a lower Halabi score (i.e., a longer predicted survival), a longer PSA–DT, and a higher hemoglobin level at baseline, but no other clinical variables (Table 1). Interestingly, after adjusting for baseline hemoglobin levels, the immune subset variables still seem to significantly associate with longer survival (Table 3). It should be pointed out that all four doses of ipilimumab were included in the comparisons with OS, and one must be well aware of the risk that numerous comparisons could lead to false positives, which could lead to false conclusions. Therefore, the data generated in this study should strictly be considered as hypothesis generating data, and larger randomized studies are necessary to draw more definitive conclusions.



The rate of increase in ALC during the treatment for melanoma patients with ipilimumab has been shown to associate with clinical benefit in some previous trials, but many trials have also refuted this hypothesis [31, 32]. The current study could not show an association of increased ALC with clinical benefit, which may be due to any number of factors (e.g., melanoma vs. prostate cancer patients), although there was a slight trend for increased OS in patients with higher ALC at baseline ( $P = 0.057$ , adjusted  $P = 0.11$ ).

We evaluated the frequencies of 36 different immune cell subsets at three time points—one prior to and two post-therapy—and performed correlative studies with overall survival. We found an increased frequency of ICOS<sup>+</sup> CD4<sup>+</sup> T cells after therapy (Fig. 2b), in accordance with the previous studies [33, 34], although this increase did not correlate with survival. Expression of ICOS on CD4<sup>+</sup> T cells is necessary for effector memory development, reactivation, and survival [35, 36]. The frequency of ICOS<sup>+</sup> was analyzed from the entire CD4<sup>+</sup> population excluding Tregs. The ICOS/ICOSL pathway has been reported to be required for maximal anti-tumor effects following treatment with anti-CTLA-4 monoclonal antibodies [37], and a persistent increase in ICOS<sup>+</sup> CD4<sup>+</sup> T cells over 12 weeks correlated with OS in a retrospective analysis of melanoma patients treated with ipilimumab [38]. As seen in Fig. 2C, there was an increase in the frequency of IFN $\gamma$ <sup>+</sup> CD4<sup>+</sup> T cells during therapy in the study reported here. This may be beneficial by activating CD8<sup>+</sup> T cells and macrophages in the tumor microenvironment [39], thereby enhancing anti-tumor immunity.

Regulatory T cells are a major immunosuppressive entity, which increases tolerance and counteracts successful immunotherapy. We defined Tregs as CD4<sup>+</sup> CD25<sup>HI</sup> FoxP3<sup>+</sup> CD127<sup>NEG</sup> and further evaluated the expression of CTLA-4 on the surface. The CTLA-4<sup>+</sup> Treg population was previously shown to be more suppressive than the CTLA-4<sup>NEG</sup> population in prostate cancer patients treated with PSA-TRICOM [7]. In the present study, there was an association ( $P = 0.005$ , adjusted  $P = 0.01$ ) between the absolute number of CTLA-4<sup>NEG</sup> Tregs at baseline and survival (Fig. 3C). However, there was no association of survival with the more suppressive subset of Tregs (CTLA-4<sup>+</sup>) at baseline ( $P = 0.82$ ). Regimens that decrease the numbers and/or activity of the Treg population have shown promising results, and one previous study showed that the efficacy of anti-CTLA-4 treatment against melanoma was mediated by Fc-dependent depletion of tumor-infiltrating Tregs [40]. It has also been shown in melanoma that CTLA-4 blockade of T effectors and Tregs concomitantly gives the greatest treatment efficacy [41].

In this trial, lower levels of PD-1<sup>+</sup> Tim-3<sup>NEG</sup> CD4<sub>EM</sub> cells at baseline associated with longer survival (Fig. 3a,

$P = 0.005$ , adjusted  $P = 0.01$ ). This could suggest that therapy blocking PD-1 could be beneficial for the patients who display high levels of PD-1<sup>+</sup> T cells. PD-1 is expressed by activated lymphocytes [42] and inhibits the effector functions and proliferation after binding to its ligand, PD-L1 (B7-H1). Interruption of the PD-1/PD-L1 pathway is currently being investigated and has shown promising results in melanoma and several carcinomas [43, 44]. In T-cell exhaustion, PD-1 and Tim-3 are coexpressed on the cell surface [45], and these cells produce fewer cytokines and show less proliferation. Some reports suggest that when PD-1 is expressed without Tim-3, this may be more indicative of T-cell activation than of T-cell exhaustion.

We also found a trend that higher levels of Tim-3<sup>+</sup> PD-1<sup>NEG</sup> CD8<sup>+</sup> T cells at baseline associated with longer overall survival (Fig. 3b,  $P = 0.002$ , adjusted  $P = 0.004$ ). It has previously been reported that Tim-3-expressing human CD8<sup>+</sup> T cells exhibit an effector memory phenotype, and strong effector functions in tuberculosis [46], which would support our finding. In contrast, T cells expressing both Tim-3 and PD-1 may exhibit an exhausted phenotype. We did not find any associations between the central memory subsets of CD8<sup>+</sup> T cells and clinical outcome in this trial.

We found that an increase during therapy in the NK-cell immature subset that expresses Tim-3 was associated with increased survival (Fig. 4a,  $P = 0.0074$ , adjusted  $P = 0.015$ ). Tim-3 is a maturation marker on NK cells and acts as a coreceptor to enhance IFN $\gamma$  production [47]. Tim-3<sup>+</sup> NK cells are fully responsive with respect to cytokine production and cytotoxicity, but may be negatively regulated when encountering target cells expressing ligands of Tim-3 [16]. This may thus be an important immune cell subset to monitor in future clinical immunotherapy trials.

In addition to the comparisons with all patients, we also evaluated whether there were any differences in OS between the cohort of patients that received 10 mg/kg of ipilimumab ( $n = 15$ ), compared to the combined cohorts that received <10 mg/kg ( $n = 15$ ) for 6 of the parameters; there were no statistical differences in the baseline Tim-3<sup>+</sup> PD-1<sup>NEG</sup> CD8 T cells, baseline PSA-DT, baseline hemoglobin, or the change in Tim-3<sup>+</sup> NK cells. There were greater differences in OS at baseline in PD-1<sup>+</sup> Tim-3<sup>NEG</sup> effector memory CD4<sup>+</sup> T cells and CTLA-4<sup>NEG</sup> Tregs in patients receiving lower doses of ipilimumab versus 10 mg/kg dose ( $P = 0.0001$  vs.  $P = 0.36$ , and  $P = 0.021$  vs.  $P = 0.22$ , respectively). This may suggest that the higher dose level of ipilimumab can overcome some of the underlying immune deficiencies present in patients, whereas in the lower dose groups, a deficiency in either of these cell subsets has an impact on the clinical outcome.

We have previously published the results of IFN $\gamma$  ELISPOT analysis for PSA-peptide responses in this trial [12]. Unfortunately, only 9 of the 30 patients enrolled on this

trial had the MHC class I allele HLA-A2, which to date is the only allele for which we have a functional ELISPOT assay to measure PSA-specific responses. Of the 9 patients who were HLA-A2<sup>+</sup>, 6 were in the <10 mg/kg cohort and 3 were in the 10 mg/kg cohort. The only 3 patients that had any post-vaccine level of elevation of PSA-specific T cells above the pre-vaccine levels were the 3 patients in the 10 mg/kg cohort. Therefore, it was not feasible for the current trial to relate the findings in immune cell subsets with the PSA-specific responses.

It should also be pointed out that both preclinical and clinical studies have indicated the potential importance of “antigen cascade,” also termed “epitope spreading,” in anti-tumor responses. For example, in one study [48, 49], CEA transgenic mice bearing CEA-expressing tumors were vaccinated with a recombinant vaccine directed against CEA. While control studies showed CEA was needed to be present in both the vaccine and tumor for the induction of anti-tumor responses, it was determined that the more significant immune response primarily responsible for the anti-tumor effect was that directed against an endogenous antigen (gp70) present in the tumor. Clinical studies in breast cancer also showed that there was a correlation with clinical benefit for those patients who demonstrated the phenomenon of antigen cascade in PBMCs post-vaccination [50–53]. Future vaccine-based clinical studies are being designed to measure the breadth of antigen-specific responses post-vaccination if sufficient PBMCs are available.

The data reported here are intended to be hypothesis generating, and larger randomized, controlled immunotherapy trials need to be evaluated in a similar manner to determine whether the analysis of specific immune cell subsets pre-treatment or early in the treatment cycle are predictive of clinical outcome. Results of such analysis may of course depend on the immunotherapeutic agent being evaluated, prior therapies of patients, and the disease and disease stage being investigated. It is interesting to note that in the study reported here, strong associations with OS were seen in specific immune cell subsets (Figs. 3, 4) as compared to conventional clinical parameters such as Gleason score, baseline PSA, and PSA-doubling time (Table 1). Adjusting for the independent effect of baseline hemoglobin levels, the changes during therapy in the other variables still seem to be associated with longer survival.

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