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### Phase II trial of a GM-CSF-producing and CD40L-expressing bystander cell line combined with an allogeneic tumor cellbased vaccine for refractory lung adenocarcinoma

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#### Abstract

We created a vaccine in which irradiated allogeneic lung adenocarcinoma cells are combined with a bystander K562 cell line transfected with hCD40L and hGM-CSF. By recruiting and activating dendritic cells, we hypothesized that the vaccine would induce tumor regression in metastatic lung adenocarcinoma. Intradermal vaccine was given q14 days x3, followed by monthly x3. Cyclophosphamide (300 mg/m<sup>2</sup> IV) was administered before the 1st and 4th vaccines to deplete regulatory T-cells. All-trans retinoic acid was given (150/mg/m<sup>2</sup>/day) after the 1st and 4th vaccines to enhance dendritic cell differentiation. Twenty-four participants were accrued at a single institution from 10/2006 to 6/2008, with a median age 64 and median of 4 previous lines of systemic therapy. A total of 101 vaccines were administered. Common toxicities were headache (54%) and site reaction (38%). No radiologic responses were observed. Median overall survival (OS) was 7.9 months (mo) and median progression-free survival (PFS) was 1.7 mo. Of 14 patients evaluable for immunological study, 5 had peptide-induced CD8+ T-cell activation after vaccination. Overall, vaccine administration was feasible in an extensively pretreated population of metastatic lung cancer. Despite a suggestion of clinical activity in the subset with immune response, the trial did not meet the primary endpoint of inducing radiologic tumor regression.

#### INTRODUCTION

Due to high annual incidence and poor long-term survival, lung cancer remains an ideal target for novel agents such as immunotherapy. In particular, the treatment of patients with

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The University of South Florida holds a patent for GM.CD40L and CCL21 with one author (S.A.), application: 12/933,568, publication: US 2011/0059137, Filing date: Mar 19, 2009.

The authors have no other conflicts of interest to disclose.

#### HUMAN SUBJECTS STATEMENT:

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The procedures within were approved by the Institution Review Board (IRB) approval from the University of South Florida (FWA00001669), in accordance with the Helsinki Declaration of 1975.

advanced non-small cell lung cancer (NSCLC) is frequently complicated by co-morbid conditions and older age.<sup>1</sup> Thus, tumor vaccines may be ideal in this population due to their favorable toxicity profile.<sup>2</sup> Unfortunately, tumor-associated antigen (TAA) vaccination alone is usually insufficient to induce innate immunity, likely due to host immune incompetence and tumor-related immune suppression.<sup>3</sup> Therefore, strategies to induce or deregulate co-stimulatory protein interactions have been investigated. In particular, dendritic cells (DC) are the most potent antigen presenting cells (APC) that express co-stimulatory molecules.<sup>4</sup> DCs are activated by the cytokine granulocyte-macrophage colony stimulating factor (GM-CSF).<sup>5</sup> Furthermore, the maturation of DCs from immunosuppressive myeloid-derived suppressor cells (MDSCs) is induced by the combination of GM-CSF with IL-4<sup>6</sup> or IL-10.<sup>7</sup>

Several previous vaccine trials in NSCLC have tested methods of recruiting dendritic cells with GM-CSF. An adenoviral vector for delivery of hGM-CSF gene was safe in NSCLC<sup>89</sup> and a larger trial in NSCLC suggested a correlation of cell dose to survival.<sup>10</sup> Unfortunately, this approach was hampered by feasibility, since genetic transduction of individual tumors required a median of 50 days from harvest to treatment. A scientific advance was the creation of a "bystander" cell line derived from K562, which is universally major histocompatability complex (MHC) negative.<sup>11</sup> This line was stably transfected with plasmid vector to secrete GM-CSF, eliminating the burden of genetic modification of autologous cells. However, when this bystander was combined with autologous TAAs in NSCLC, no tumor regression was observed.<sup>12</sup> Subsequently, it was discovered that GM-CSF-expressing bystander vaccine at high doses may actually impair immunity by recruitment of induced Gr1<sup>+</sup>/CD11b<sup>+</sup> myeloid suppressor cells.<sup>1314</sup> Similarly, anti-tumor vaccine activity is often attenuated by induction of regulatory T-cells.<sup>151615</sup> Due to the antigenic heterogeneity of NSCLC, many trials have relied upon autologous tumor for vaccine TAAs. However, autologous collection suffers from several potential drawbacks: failure to harvest, unsuccessful processing or contamination, and patient progression while awaiting vaccine preparation.<sup>17</sup>

To address these problems, we created a bystander K562 cell line which was transfected with GM-CSF and CD40L plasmids admixed with two lung adenocarcinoma cell lines as the source of shared tumor antigens.<sup>18</sup> CD40L expression is believed to augment DC activation at the local vaccine site.<sup>19</sup> Our bystander cell line (GM.CD40L) was more effective in inducing responses in cultures of tumor-draining lymph nodes compared to autologous vaccine alone.<sup>20</sup> In a Phase I trial of GM.CD40L with an autologous tumor vaccine, antitumor immune responses as well as some durable radiologic stable disease was observed.<sup>21</sup> Next, we designed a preparation of two lethally irradiated lung adenocarcinoma cell lines as a shared tumor antigen. This contains together HER-2/neu, CEA, GD-2, WT-1, MAGE-A1 and -A3.<sup>22</sup> In this approach, thousands of potential lung tumor epitopes within the lysate may be taken up and cross-presented by both MHC class I and II molecules on DCs.<sup>23, 24</sup> Thus, screening for specific TAAs or matching HLA type in individual patients is not required. All-trans-retinoic acid (ATRA) was added to induce differentiation of immature DCs at the local vaccine site.<sup>25</sup> Cyclophosphamide pretreatment was included to reduce the number and function of CD25+ T-regulatory cells.<sup>26</sup> We hypothesized that this vaccine preparation would induce objective responses in a clinical trial of patients with advanced lung adenocarcinoma.

#### **METHODS**

#### Study Design

This was an open-label, single institution, single-arm, phase II trial designed to evaluate the safety, efficacy, and immunogenicity of a GM.CD40L plus tumor cell-based vaccine.

Intravenous cyclophosphamide at 300 mg/m<sup>2</sup> was given on day 1 and 57. The vaccine was then administered intradermally in the axillary and inguinal lymph node basins; every 2 weeks for the first 4 weeks (days 4, 18, and 32), then monthly for the next 3 months (days 60, 88, and 116). All-trans retinoic acid (ATRA,) at 150 mg/m<sup>2</sup>/day, was given orally thrice daily after the first and fourth vaccine (days 5–7 and days 61–63). CT of chest, abdomen, and pelvis was done at baseline, with reassessment at week 8, week 17, and every three months thereafter using Response Evaluation Criteria in Solid Tumors (RECIST) v 1.0. This schedule was followed uniformly and consistently. Hematologic, liver, and renal function was assessed, and toxicity was graded according to National Cancer Institute Common Toxicity Criteria v 3.0. Biologic materials were registered with the Food and Drug Administration (BB-IND-13088) and Office of Biologic Activities (NIH-OBA-0608-801). The trial was nationally registered (NCT00601796). All patients signed the protocol-specific informed consent approved by local institutional review board (FWA00001669).

#### **Study Population**

Eligible patients for this study were 18 years or older, with histologically-confirmed American Joint Committee on Cancer (AJCC) clinical stage IIIB or IV lung adenocarcinoma that was refractory or relapsed after at least one previous systemic therapy. Participants had to have an Eastern Cooperative Oncology Group performance status of 0 or 1, an absolute granulocyte count > 1,500/ $\mu$ L, platelet count > 100,000 / $\mu$ L, hematocrit >25%, total bilirubin < 2 mg/dL, and serum creatinine < 2 mg/dL. Participants were required to have progressive, measurable disease by RECIST criteria. Patients with brain metastases were eligible if the disease was asymptomatic. Patients were excluded if they had received prior systemic corticosteroids, radiation, or chemotherapy within 4 weeks of first vaccination. Patients withpregnancy or a history of HIV were not eligible to participate in the study.

#### Study Drug

The GM.CD40L bystander cell line was prepared as previously described.<sup>21</sup> Two human cell lines, NCI-H1944 and NCI-H2122, were used as the TAA source. These cell lines were originally isolated from two smokers with stage IIIB lung adenocarcinoma.<sup>27</sup> H2122 and H1944 were tested for correct identification and karyology, and products were free of viral, retroviral, and bacterial contaminants (WuXi AppTec®, St Paul MN). These cell lines do not constitutively express HLA-A2.<sup>22</sup>

For the therapeutic vaccine, one vial containing  $7.5 \times 10^6$  radiated H1944 adenocarcinoma cells, 1 vial containing  $7.5 \times 10^6$  radiated H2122 adenocarcinoma cells, and 1 vial containing  $15 \times 10^6$  radiated GM.CD40L bystander cells were thawed rapidly to  $37^\circ$ C, combined, diluted in 10 mL sterile saline for 15–30 min at  $37^\circ$ C, centrifuged, and resuspended in a final volume of 1.1 mL sterile saline. A 0.1 mL aliquot was sent for microbiological testing (including Gram stain and culturing for aerobes, anaerobes, and fungi) and immunofluorescence analysis after staining with DAPI, to eliminate the possibility of mycoplasma contamination. If the testing ruled out microbial contamination, the vaccine was transported on ice and injected intradermally. Each injection consisted of 0.25 mL of cell suspension, containing  $15 \times 10^6$  irradiated tumor cells and  $15 \times 10^6$  irradiated GM.CD40L bystander cells. Patients were monitored for acute toxicity for 1 hour after injection.

#### **Immune Function Testing**

At baseline and after each vaccination, 10 mL of blood was collected by phlebotomy. Peripheral blood mononuclear cells (PBMCs) were isolated using a density gradient protocol (Ficoll-PaqueTM PLUS, GE Healthcare Biosciences, PA), and frozen in liquid nitrogen. All samples from each patient were analyzed simultaneously to reduce inter-assay variability.

MNC were thawed, incubated overnight in complete medium (RPMI-1640 supplemented with antibiotics and 10 % FBS), and then used in experiments. For ELISPOT assay,  $2 \times 10^5$  PBMCs /well of AIM-V medium containing 10% human AB serum and 2ng/ml rhIL-2 were added to 96-well plates. HLA-A2-presented peptides derived from WT1, RMFPNAPYL<sup>28</sup>; CEA, YLSGANLNL<sup>29</sup>; and hTERT, ILAKFLHWL<sup>30</sup> were added to the appropriate wells. If HLA-A2 negative, irradiated H1944 cells were used for peptide. After a 36 h restimulation culture, ELISPOT assays were performed with anti- $\gamma$ -IFN (Serial # 552138 - BD bioscience, San Jose CA). Controls included PBMCs stimulated with an irrelevant peptide as a negative control and PBMCs stimulated with anti-CD3 as a positive control. The plates were cultured for 18 hours, washed, and the detection antibodies were added. Resultant spots were counted with Immunospot Series I Analyzer (Cellular Technology Ltd, Shaker Heights OH). A response was defined as positive if it met these criteria: 1) the mean number of spots present in the peptide wells was greater than 20 spots per 200,000 PBMC added, and 2) a greater than 3-fold increase over the standard deviation of the pre-treatment PBMC ELISPOT was observed.

#### Analysis of Cell Phenotype

Cell phenotype was evaluated by multicolor flow cytometry using a LSR II flow cytometer and monoclonal antibodies obtained from Becton–Dickinson. The following antibodies were used: CD4-Alexa 700; CD3-PE, CD14-PE, CD19-PE, CD56-PE (as lineage-PE); CD25-PE; CD127-Alexa 647; HLA-DR-APC; CD33-PE-Cy7. The following combinations of antibodies were used to identify cell populations: Lineage<sup>–</sup> (Lin) (CD3, 14, 19, 56) HL-DR<sup>+</sup>; Mature DC: Lin<sup>–</sup> HLA-DR<sup>+</sup>; MDSC: Lin<sup>–</sup> HLA-DR<sup>–</sup> CD33<sup>+</sup>; T-reg: CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>–</sup>. Dead cells were removed from the analysis by using DAPI staining.

For PD-L1 expression, NCI-H1944 and NCI-2122 cells were incubated in complete media with or without 50 U/mL interferon- $\gamma$  (R&D Systems, Minneapolis) for 18 hours. The cells were analyzed by staining monoclonal anti-human CD274-FITC, clone MIH1 (BD Bioscience, New Jersey) in combination with corresponding isotype control, then imaged using FACScan (Becton Dickinson, Mountain View, California) and CellQuest software (Becton Dickinson).

#### **Statistical Plan**

This trial used a Simon minimax design<sup>31</sup> based upon the primary endpoint of objective response rate, in which  $H_0 = 15\%$  and  $H_1 = 30\%$  with one-sided  $\alpha = 0.05$  and  $\beta = 0.20$ . The trial had 54% probability of stopping early at 23 patients if  $H_0$  was true, and 5% chance of stopping early if  $H_1$  was true. All patients exposed to study drug were included in the safety and survival analyses. All statistical tests were two-tailed unless otherwise specified. Mean ELISPOT responses were compared by Wilcoxon signed rank, or if two groups were present, Friedman with Dunn's post-test to control for multiple comparisons.<sup>32</sup> Best overall response was defined as best response recorded from first vaccine to progression or death. Progression free survival (OS) defined from date of treatment until date of progression or death, and overall survival (OS) defined from date of treatment until death. Survival curves were calculated according to the Kaplan-Meier method. Since no patients were censored, the effect of immune response on PFS and OS was evaluated using the Wilcoxon rank-sum test. Figures and survival plots were generated using GraphPad Prism® (version 5.04, San Diego CA).

#### RESULTS

Twenty-four participants were accrued at a single institution from October 2006 to June 2008, with data cut-off in January 2012. Participant demographics are presented in Table 1.

Patients had a median age of 64 years (range 48 - 84) and had received a median of 4 previous lines of systemic therapy prior to entry. All were Caucasian. Twenty were former or active smokers, and 4 had brain metastases. Eleven patients were HLA-A2 positive.

#### Safety

A total of 101 vaccines were administered. One grade 3 diarrhea, one grade 3 nausea, one grade 3 abdomen pain, and three grade 3 headache were observed related to study medication. Adverse events are presented in Table 2. The most common treatment-related adverse events included injection site reaction (38%), fatigue (21%), and headache (54%). All adverse effects were manageable and without significant clinical sequela. No auto-immune adverse events such as endocrine dysfunction or vitiligo were observed. No participants stopped therapy due to adverse events or elective discontinuation. Ten patients (42%) completed the initial four month course of vaccine therapy and continued vaccination, with three remaining on treatment at one year.

#### Survival

After at least three vaccines, ten (42%) of patients had stable disease by RECIST criteria. No confirmed radiologic responses were observed. Therefore, the trial met predefined stopping criteria and closed to accrual. Median progression-free survival was 1.7 months (mo) [95%CI NE – 4.6 mo], and median overall survival was 7.9 mo [95%CI 4.1 – 11.6 mo] (Figure 1). One-year survival was 33% and 3-year survival was 8%. Four patients expired from cardiac disease, four from obstructive lung disease, and the remainder died from progressive lung cancer. For unclear reasons, the eleven HLA-A2 positive participants appeared to have improved PFS compared to the thirteen HLA-A2 negative (median PFS 4.4 vs. 1.5 mo, p = 0.02) and trend towards improved OS (median OS 14.0 vs 5.0 mo, p = 0.06).

#### Immune Response

Samples for evaluation of vaccine-induced immunity were available from 14 patients. Patients who were HLA-A2 positive were tested for a response to three peptides (CEA, TERT2, and WT-1) as a sentinel measure of vaccine activity. Similarly, patients who were HLA-A2 negative were tested for response to H1944 cell lysate. Responses ranged between 40 and 700 spots/ $2 \times 10^5$  PBMCs, presented in Figure 2. Overall, 36% of evaluable patients met the pre-specified definition for immune response. All immune responses were statistically significant compared to baseline with p < 0.05. An association of immune response with risk of progression was not statistically significant (HR = 0.33 and p = 0.10, one-tailed  $\chi^2$ ), as presented in Figure 3B. Compared to non-responders, immune responders appeared to have longer median OS (23.0 vs 7.1 mo, p = 0.04) and a similar but nonstatistically significant median PFS (5.6 vs. 1.7 mo, p = 0.18), as presented in Figure 4. However, immune responders also appeared to have more favorable baseline prognostic features, such as less grade 3 cancer (p = 0.02) and trend toward more metastatic disease limited to the lungs alone (p=0.08), as presented in Table 3. No statistically significant association between HLA A2 and immune response was observed, although this was limited by small sample size (RR 1.8, 95% CI 0.79 – 4.2, *p* = 0.30).

#### Peripheral Mononuclear Cells

Of the patients evaluable, peripheral blood lymphocytes decreased by Day 45 (p = 0.04) and Day 88 (p=0.02) compared to baseline, as presented in Table 4. However, no significant difference in absolute blood lymphocytes between immune responders and non-responders was observed. An increase in Lin<sup>-</sup> HLA-DR<sup>-</sup> CD33<sup>+</sup> MDSCs was observed at Day 45 compared to Day 0 (p = 0.04), which appeared to be more pronounced in non-responders (p

= 0.03), as seen in Figure 5A. No change in Lin<sup>-</sup>HLA-DR<sup>+</sup> mature dendritic cells was observed at Day 45 compared to Day 0, with mean % total live PBMCs of 6.7 vs. 6.2; p = 0.82. These findings were limited by small sample size due to only nine evaluable paired samples, with the remainder of patients non-evaluable due to off-study, deceased, or poor sample viability. Despite a bolus of cyclophosphamide at Day 0, no statistically significant change in peripheral blood CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>-</sup> T-regulatory cells was observed at Day 45, as shown in Figure 5B, although this finding was similarly limited by small sample size.

#### **Tumor-associated Vaccine Antigen**

To help identify why the allogeneic vaccine may have been unsuccessful, we tested the cell lines used as TAA for cell surface PD-L1 expression. As shown in Figure 6, no constitutive PD-L1 expression was observed in either H1944 or H2122. However, expression was inducible after incubation with IFN- $\gamma$ , in accordance with reports in other lung cancer cell lines.<sup>3334</sup>

#### DISCUSSION

This GM-CSF-producing and CD40L-expressing bystander cell line combined with an allogeneic (adenocarcinoma) tumor cell-based vaccine appeared to be safe, despite an extensively pretreated study population. However, the trial failed to meet its primary endpoint. Even in participants with an immune response to vaccination, no radiologic responses were observed. These findings are not dissimilar to previous peptide or whole-cell lysate vaccine trials, in which successful immunogenicity only rarely translated into tumor regression.<sup>3536</sup> Several mechanisms may have contributed to the historical failure of this vaccine strategy: inappropriate clinical response endpoints<sup>37</sup>, presence of tumor-induced immune suppression<sup>38</sup>, and application to populations suffering from extensive disease or impaired host immunity.<sup>39</sup> In this regard, the current success of checkpoint inhibitors in overcoming immune suppression is of particular interest because it presents an attractive companion for future immunotherapy strategies in general, and for tumor cell-based vaccines specifically.<sup>40</sup> In particular, combination of GM-CSF-secreting vaccines with CTLA-4 inhibitors has shown some promise.<sup>4142</sup> Furthermore, optimal timing and context of vaccination may also be particularly relevant in contributing to the failure of this trial. For example, a GM-CSF bystander vaccine, GVAX ®, appeared effective only if given after chemotherapy in an autologous mice model of lung cancer.<sup>43</sup>

Our findings are characterized by important limitations. First, this study had a small sample size, and only a proportion were evaluable for immune correlates. Next, we used IFN- $\gamma$ release as an indirect surrogate for host immune response to sentinel epitopes within the whole cell lysate. This method has established limitations in reproducibility and specificity,<sup>44</sup>and also does not confirm that effective cross-priming occurred.<sup>45</sup> Next, the likelihood of confounding due to nutritional status, organ function, or subsequent medications upon immune response was not addressed.<sup>46</sup> This is because the sample size was too small to permit the use of a multivariable regression model.<sup>47</sup> These co-morbid factors may confound any association between host immunity and survival, and would be expected to result in bias away from the null. As shown in Table 3, the immune responders did have more favorable tumor grade and extent of metastatic disease, so it is plausible that this explains the difference in survival observed. Therefore, conclusions about the relationship between immune response and survival should be regarded as exploratory. Moreover, the relationship between IFN-y immune response and other important correlates of T-cell immunity, such as delayed hypersensitivity response (DTH)<sup>48</sup> and presence of TGF-β-producing FOXP3+ regulatory T-cells<sup>49</sup> was not examined. In addition, measurement of T-cells in the peripheral blood may not directly correlate with the activity of

T-cells at the site of the tumor.<sup>50</sup> Thus, testing of pre- and post-treatment tumor biopsies for the density of either CD 8 + or memory T-cell infiltration would have provided corroboration for the serologic findings in this trial.

The significance of improved PFS in HLA-A2 patients is unclear. Neither the K562 bystander nor allogeneic tumor cell lines express this haplotype.<sup>22</sup> Since the allogeneic cells were lysed and irradiated, it is unlikely their MHC Class I components would have impacted the presentation of their derived proteins., and the bystander K562 cell line does not express any MHC Class I components.<sup>51</sup> It is conceivable that HLA-A2 performed better due to random chance, or unknown confounders unrelated to immunity.

Based upon a limited subset of patients, no durable change in the number of T-regs was observed, which suggests that single-bolus cyclophosphamide may be suboptimal. Despite a report that cyclophosphamide pretreatment improves G-CSF vaccine-induced immunity<sup>52</sup>, bolus cyclophosphamide has since been shown to be inefficient in reducing T-regs.<sup>26</sup> In contrast, metronomic dosing appears to be more effective<sup>53</sup>, perhaps mediated through ATP.<sup>54</sup> Since the peripheral blood MDSCs did not decrease during treatment, the utility of the ATRA treatment in this trial also remains uncertain. However, we have previously demonstrated that ATRA reduces immature myeloid derived cells in patients receiving IL-2 for metastatic renal cell<sup>55</sup>, as well as small cell lung cancer<sup>56</sup> and the latter trial is ongoing (NCT00617409).<sup>57</sup>

Some lung cancer cell lines can be induced to express cell surface PD-L1, while others have constitutive PD-L1 expression.<sup>33</sup> Since the cell lines used as TAA in this trial were lysed and irradiated prior to injection, it is unclear if PD-L1 was present in sufficient quantity at the local vaccination site to interfere with effector T-cell activity. Despite the negative clinical result, the immune responses observed in this trial suggest that this vaccine strategy may still hold potential with further improvement. Along these lines, the expression of the chemokine C-C motif ligand (CCL21) has been shown to augment anti-tumor immune responses and has been explored as an adjuvant in several vaccines.<sup>5859</sup> This evidence has led us to design another clinical trial, incorporating GM.CD40L, in combination with a cell line engineered to express exogenous CCL21, in metastatic NSCLC (NCT01433172).<sup>60</sup> This randomized phase II trial is currently underway and is expected to be completed in late 2013.<sup>61</sup>

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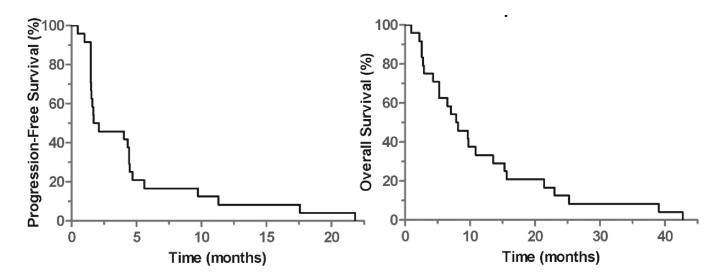
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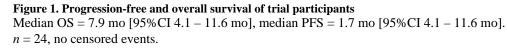
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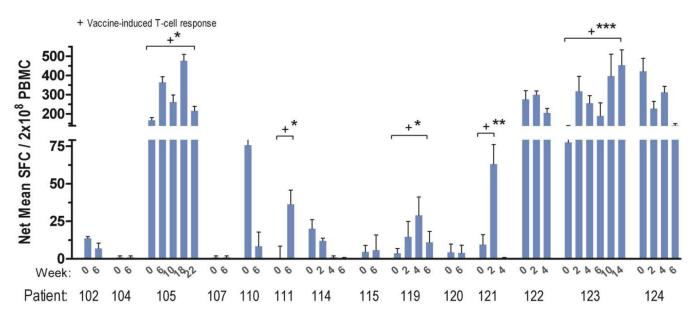


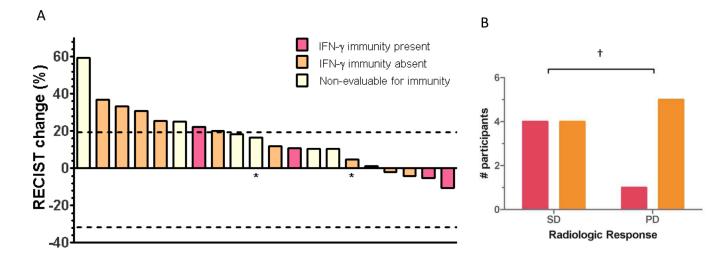
Figure 2. Patient IFN- $\gamma$  immune responses. Mean CEA / hTERT / WT-1, or H1944 lysate if A2-(n = 14)

Shown are enzyme-linked immunosorbent spot assay responses from a direct assay, comparing pre-vaccination and weekly time points after vaccination. Bars = mean  $\pm$  SD. + indicates responses that met pre-defined immune response criteria.

\* p < 0.05, \*\* p < 0.001, \*\*\* p < 0.0001 (Wilcoxon signed rank, or Friedman with Dunn's post-test if more than two comparators). SFC: spot-forming cells. PBMC: Peripheral blood mononuclear cells.

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#### Figure 3.

A. Waterfall plot of best overall radiologic response to vaccine. Immunity assessed by IFN- $\gamma$  ELISPOT. Of 24 total, 3 were non-evaluable for RECIST due to lung cancer death (n = 2) or missing evaluation (n = 1). \* PD due to unequivocal progression of malignant pleural effusion.

**B.** Participants who met pre-specified immune response criteria had non-statistically significant trend towards more favorable radiologic response. SD: Stable disease, PD: Progressive disease. RECIST v 1.0. † p = 0.10 (1-tailed  $\chi^2$ ).

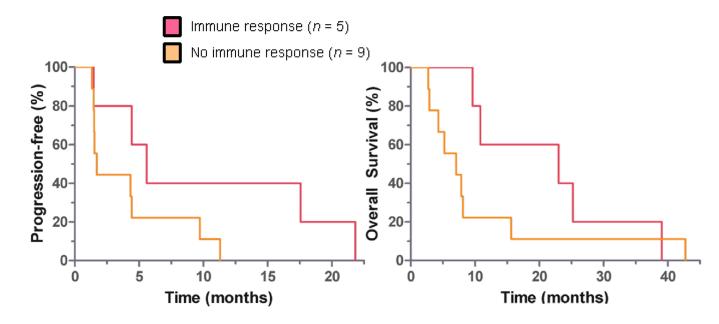
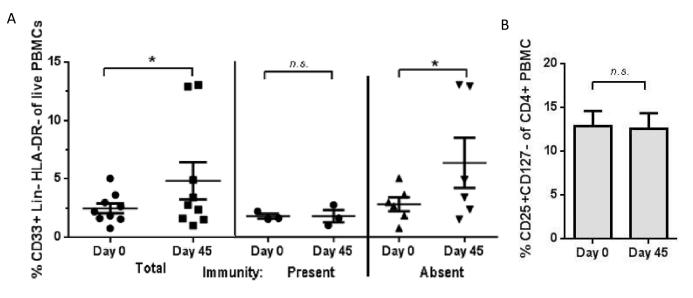


Figure 4. Participants who met pre-specified IFN- $\gamma$  immune response criteria appeared to have improved overall survival

Median OS 23.0 vs. 7.1 months  $\circ p = 0.04$ . Median PFS 5.6 vs. 1.7 months,  $\circ p = 0.18$ . Immune response determined by IFN- $\gamma$  ELISPOT assay.  $\circ$  Wilcoxon rank-sum test.

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#### Figure 5.

A. MDSCs increased from baseline (day 0) to next consecutive evaluation (day 45). MDSCs may have contributed to failure to achieve an IFN- $\gamma$  response to vaccine antigen. Fraction CD33+ of Lin-DR- were analyzed by flow cytometry. Remaining patients were not evaluable due to poor cell viability, sample not collected, or death.

Bars: mean  $\pm$  standard error. \* p < 0.05, Wilcoxon signed rank test. *n.s.* = not significant. For pairing, Spearman  $r^2 = 0.94$ .

**B.** Peripheral T-regulatory cells did not appreciably decrease from baseline (day 0) to next consecutive evaluation (day 45), despite single bolus of intravenous

**cyclophosphamide**. n = 8. Fraction CD25+CD127- of CD4+ were analyzed by flow cytometry. Remaining patients were not evaluable due to poor cell viability, sample not collected, or death.

Bars: mean  $\pm$  standard error. *n.s.* = not significant, Wilcoxon signed rank test. For pairing, Spearman  $r^2 = 1.0$ .

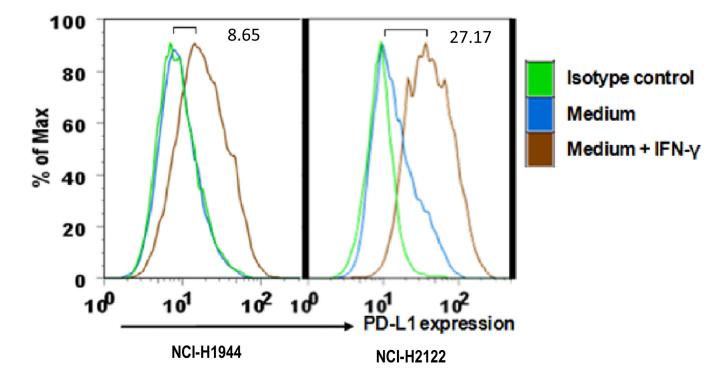


Figure 6. FACS analysis of cell-surface PD-L1 expression in the two human adenocarcinoma cell lines used as tumor antigen

Cell lines were incubated for 18 hours with or without 50 U/mL IFN- $\gamma$  and stained with monoclonal antibody against human B7-H1. Numbers on the bar indicate the mean fluorescent intensity of IFN- $\gamma$  incubated cells as compared with media-incubated cells alone. Positive staining is indicated by the blue or brown lines and control antibody (mouse IgG1) staining is shown as green lines.

#### Table 1

#### Demographics

Participant Demographics and Baseline Measurements.

Characteristics	Patients n (%)
Total	24
Age in years (Median)	64.3
Range	48 - 84
Race	
White	24 (100)
Gender	
Female	12 (50)
Male	12 (50)
ECOG <sup>*</sup> performance status	
PS 0	8 (33)
PS 1	15 (63)
PS 2	1 (4)
Smoking status	
Smoker	3 (12)
Former smoker	17 (71)
Non smoker	4 (17)
AJCC <sup>‡</sup> clinical stage	
IIIB	1 (4)
IV	23 (96)
Histotype of NSCLC	
Adenocarcinoma	24 (100)
Previous treatments	
Surgical resection	10 (42)
Radiation	13 (54)
Chemotherapy	22 (92)
Tyrosine kinase inhibitor	9 (38)
Metastatic sites	
Visceral	17 (71)
Bone	7 (29)
Brain	4 (17)

\* Eastern Cooperative Oncology Group.

 $^{\ddagger}$ American Joint Committee on Cancer

Table 2

## Therapy-related adverse events

Therapy-related adverse events

	21%				.9	.9	38%	9	~	17%	29%	~	54%		~	9
Total %	5 21	4%	4%	4%	4%	4%	9 38	2 8%	4%	4 17	7 29	4%	13 54	4%	4%	4%
Grade 4 7	5			-	1	1	5	2	1	7						
Grade 3										1	1		3		1	
Grade 2	1				1						2		3	1		
Grade 1	4	1	1	1		1	6	2	1	3	4	1	7			1
Toxicity	Fatigue	Dizzyness	Anorexia	Chills	Leukopenia	Low Platelet	Site Reaction	Dry skin	Edema: limb	Diarrhea	Nausea	ALT elevation	Headache	Joint pain	Abdomen pain	Injection pain
Category	Constitutional				Hematologic		Dermatologic			GI			Pain			

### Table 3Baseline factors related to immune response

Baseline factors related to immune response

Characteristic				p value
Immune category	Total	+	-	+ vs all others
n (%)	24 (100)	5 (26)	9 (38)	5 vs. 19
Weight, in kg (mean $\pm$ SD)	69.7 ± 14.0	$61.6 \pm 12.5$	$74.6 \pm 13.8$	0.14 <sup>§</sup>
BMI (mean ± SD)	$25.6\pm4.6$	25.1 ± 4.0	$26.8\pm5.8$	0.85 <sup>§</sup>
Albumin, g/dL (mean $\pm$ SD)	$3.91\pm0.43$	$3.90\pm0.47$	$4.08\pm0.41$	0.76 <sup>§</sup>
Poorly differentiated (n,%)	7 (30)	0 (0)	2 (22)	0.03 <sup>†</sup>
Prior bevacimab (n, %)	6 (25)	1 (20)	1 (11)	$0.99^{\dagger}$
Lung-only mets (n,%)	11 (46)	4 (80)	5 (56)	$0.08^{\dagger}$

'Lung-only mets' denotes stage IV disease with only contralateral lung metastases. "+" = IFN- $\gamma$  immunity observed, "-" = not observed. 13 patients were not evaluable for immunity.

§= Wilcoxon rank-sum

 $^{\dagger} = \chi^2$ test.

## Table 4

# Effect upon peripheral blood count

Effect upon peripheral blood count.

Time		Day 0			Day 45			Day 88	
Immune category	Total	+	-	Total	+	I	Total	+	Ι
u (%)	24 (100) 5 (26)		9 (38)	22 (100) 5 (23)		9 (41)	15 (100) 5 (33)		6 (40)
WBC ( $\times 10^{6}/ml$ )	$7.1 \pm 1.8$	$5.6\pm0.9$	$7.1 \pm 1.5$	$7.1 \pm 1.8  5.6 \pm 0.9  7.1 \pm 1.5  7.4 \pm 2.0  6.3 \pm 1.8  6.5 \pm 1.2  9.4 \pm 4.5  5.8 \pm 1.0  10.6 \pm 5.0  10.6 \pm 5.0 $	$6.3\pm1.8$	$6.5\pm1.2$	$9.4\pm4.5$	$5.8 \pm 1.0$	$10.6\pm5.0$
Neutrophils (×10 <sup>6</sup> /ml) 4.7 ± 1.4 3.6 ± 1.0 4.8 ± 0.9 5.3 ± 1.6 <sup>*</sup> 4.2 ± 1.4 4.9 ± 1.3 7.1 \pm 4.3 <sup>*</sup> 4.1 \pm 0.6 8.1 \pm 4.6	$4.7 \pm 1.4$	$3.6 \pm 1.0$	$4.8\pm0.9$	$5.3\pm1.6^*$	$4.2 \pm 1.4$	$4.9\pm1.3$	$7.1\pm4.3^*$	$4.1\pm0.6$	$8.1 \pm 4.6$
$ \label{eq:Lymphocytes} Lymphocytes (\times 10^6/ml)  1.5 \pm 0.8  1.4 \pm 0.7  1.5 \pm 0.6  1.4 \pm 0.7^*  1.4 \pm 0.6  1.2 \pm 0.7  1.3 \pm 0.6^*  1.1 \pm 0.4  1.4  1.4 \pm 0.4  1.4 $	$1.5 \pm 0.8$	$1.4 \pm 0.7$	$1.5\pm0.6$	$1.4\pm0.7^*$	$1.4 \pm 0.6$	$1.2 \pm 0.7$	$1.3\pm0.6^*$	$1.1 \pm 0.4$	$1.4 \pm 0.4$
$Monocytes  (\times 10^6/ml) \qquad 0.6 \pm 0.2 \qquad 0.5 \pm 0.2 \qquad 0.5 \pm 0.2 \qquad 0.5 \pm 0.3 \qquad 0.5 \pm 0.1 \qquad 0.5 \pm 0.2 \qquad 0.6 \pm 0.3 \qquad 0.5 \pm 0.1 \qquad 0.5 \pm 0.5 = 0.5 = 0.5 \pm 0.$	$0.6 \pm 0.2$	$0.5\pm0.2$	$0.5\pm0.2$	$0.6\pm0.3$	$0.5\pm0.1$	$0.5\pm0.2$	$0.6\pm0.3$	$0.5\pm0.1$	$0.7\pm0.3$

Mean absolute cell count  $\pm$  standard deviation are shown. "+" = IFN- $\gamma$  immune response observed by Day 88, "-" = immune response not observed. 13 patients were not evaluable for immunity. 'Day 0' is day of cytoxan pretreatment. Day 45 and 88 are  $\pm 3$  days.

Asterisks indicate p < 0.05, with each timepoint compared to pretreatment level by Wilcoxon signed rank.

No significant differences were observed between "+' and "-" groups by Wilcoxon rank-sum