**ORIGINAL ARTICLE**

**In vitro activation of hTERT-specific T cell responses in lung cancer patients following chemotherapy**

Benedikt Gahn¹, Matthias Staudinger², Katrin Woester¹, Dominique Wellnitz¹, Sebastian Boettcher¹, Martin Gramatzki², Michael Kneba¹

¹Second Department of Medicine, University Medical Center Schleswig-Holstein, 24116 Kiel, Germany; ²Division of Stem Cell Transplantation and Immunotherapy, Second Department of Medicine, University Medical Center Schleswig-Holstein, Kiel, Germany

**ABSTRACT**

**Objective:** The aim of this study was to examine chemotherapy concomitant *in vitro* activation of human telomerase reverse transcriptase (hTERT)-specific T cell responses in peripheral blood mononuclear cell (PBMC) samples of patients with advanced non-small cell lung cancer (NSCLC).

**Methods:** PBMCs depleted of regulatory T cells were stimulated by peptide loaded dendritic cells (DC) matured either by application of cytokines (cDC) or a Toll-like receptor 7/8-agonist combined with a soluble CD40-ligand (ligDC). The hTERT peptide-specific T cell responses were assessed using flow cytometry for intracellular interferon-γ (IFN-γ).

**Results:** After cDC activation, T cells producing IFN-γ in response to hTERT were found in PBMC samples of 4 patients. In 2 of these patients the hTERT-specific T cell responses were further increased after ligDC application. However, PBMC of 3 other patients showed little or no induction of hTERT-specific T cell responses as a result of the methods applied during this study.

**Conclusions:** These results indicate, that concomitant to chemotherapy hTERT-specific T cell responses can be activated in PBMC of NSCLC patients *in vitro*. This activation can be further increased by ligDC though the number of responding patients is still limited.

**KEY WORDS**

Lung cancer; dendritic cells; telomerase; T cells

**Corresponding to:** Benedikt Gahn, MD, Assistant Professor. Second Department of Medicine- Hematology/Oncology, University Medical Center Schleswig-Holstein, Chemnitzstraße 33, 24116 Kiel, Germany. Email: B.Gahn@med2.uni-kiel.de.

Submitted Apr 04, 2013. Accepted for publication May 14, 2013. Available at www.jthoracdis.com

ISSN: 2072-1439 © Pioneer Bioscience Publishing Company. All rights reserved.

---

**Introduction**

Patients with advanced non-small cell lung cancer (NSCLC) have an unfavourable prognosis. Although recent therapeutic advances using tyrosine kinase inhibitors enabled improved prognosis—particularly for patients with epidermal growth factor receptor (EGFR) mutations—most patients still show an unsatisfactory median survival of only 8 months (1).

In addition to current radio- and chemotherapy protocols, tumor antigen-specific T cell mediated immunotherapy is widely accepted as an attractive therapeutic strategy. In particular, CD8⁺ cytotoxic T lymphocytes (CTL) are considered to be major mediators of tumor immuno-surveillance (2,3) recognising tumor-associated antigens (TAA) as cognate peptides bound to major histocompatibility molecules expressed on the surface of tumor cells. A specific T cell receptor (TCR) binds to these peptide-MHC (pMHC) complexes causing CTL to proliferate, produce cytokines, detect and mark target cells presenting the same antigen for lysis. Both active and passive immunotherapies using TAA-specific T cells have already demonstrated their efficiency in a number of clinical studies (4-6).

Human telomerase reverse transcriptase, hTERT, has been identified as the catalytic enzyme required for telomere elongation (7) and is expressed in 93-100% of NSCLC (8,9). Former studies have already demonstrated that peptides derived from hTERT are (I) naturally processed by tumors; (II) are presented on MHC molecules and (III) trigger effector functions of specific CTL (10-12).

Clinical trials of multiple vaccine formulations have illustrated that hTERT-specific immune responses can be safely induced in cancer patients and have a noticeable impact on clinical results. In HLA-A2⁺ women with metastatic breast cancer who were subcutaneously vaccinated with the hTERT 1540 peptide, an
The induction of NY-ESO-1 resulted in the induction of an intensive maturation signal.

**Materials and methods**

**Patients**

All patients were treated with standard 1st line platinum containing chemotherapy regimes at the University Medical Center Schleswig-Holstein. After informed consent, peripheral blood samples from 7 patients (A-G) with advanced NSCLC (Table 1) were obtained between chemotherapy cycles. Corresponding HLA-typing of the peripheral blood mononuclear cells (PBMC) was performed by PCR.

**Isolation of PBMC and generation of DC**

PBMCs were isolated by centrifugation on a Ficoll-Paque plus (GE Healthcare Bio-Sciences AB, Munich, Germany) density gradient from peripheral blood samples. DCs were obtained as previously described (22) with minor modifications. Briefly, PBMCs were resuspended at 2×10^7 cells/mL in 10 mL CellGroDC (CellGenix, Freiburg, Germany) and incubated for 2 h in a 250 mL plastic flask. The non-adherent lymphocytes (PBL) were removed and frozen in 50% RPMI 1640 (Biochrom, Berlin, Germany) supplemented with 40% heat-inactivated pooled human AB serum (Baxter, Unterschleissheim, Germany) and 10% DMSO according to standard protocols. The adherent fraction was cultivated in presence of 800 U/mL GM-CSF and 500 U/mL IL-4 (R&D, Wiesbaden, Germany). Fresh medium supplemented with cytokines was added on days 3 and 5. On day 6, floating cells were harvested and used for antigen loading.

**DC pulsing with hTERT peptides and maturation protocols**

For activation of HLA-A and HLA-DR-restricted T cell responses hTERT peptides (Table 2) were used to load stimulator cells. 30 μg of each hTERT peptide were added to the
DCs (3×10⁶) in 1mL CellGro supplemented with 500 U/mL IL-4 and 800 U/mL GM-CSF. DCs were incubated at 37 °C for 2 h. For DC maturation a terminal differentiation-inducing standard cytokine cocktail consisting of 2,000 IE/mL IL-1β, 1,000 IE/mL IL-6, 1,000 IE/mL TNF-α (all: R&D Systems, Wiesbaden, Germany) and 1 μg/mL prostaglandin E2 (Sigma Aldrich, St. Louis, USA) was added and the culture was incubated overnight (26). For comparison, T cells were treated with 1 μg/mL CL097 (InvivoGen, San Diego, USA) for 2 h and matured with 1 μg/mL sCD40L (Miltenyi, Bergisch-Gladbach, Germany) for an additional 20 h. Subsequently, DCs were washed and used for stimulation of autologous T cells.

Prior to T cell culture non-adherent PBMCs from 7 patients (A-G, Table 1) were depleted of CD4⁺CD25⁺ T regs using the CD4⁺CD25⁺ regulatory T cell isolation kit (Miltenyi Biotec, Bergisch-Gladbach, Germany) according to the manufacturer’s instructions. CD4⁺ cells were separated with a biotinylated CD4 cell antibody cocktail. Following positive selection of CD4⁺CD25⁺ cells with CD25 MicroBeads, the CD4⁺ and CD4⁺CD25⁺ population were co-cultured in vitro with hTERT peptide-pulsed DCs using a protocol adapted from a previous study (22). Briefly, 1×10⁶ non adherent PBMC depleted of T regs were cultured in RPMI 1640 and 2 mM L-Glutamine (both Biochrom, Berlin, Germany), 5% AB-Serum (Baxter, Unterschleissheim, Germany), 50 μg/mL Gentamicin (GIBCO, Invitrogen, Darmstadt, Germany) with hTERT-pulsed DC (1×10⁵). After 7 days in culture, a second stimulation was performed followed by addition of 20 IU/mL IL-2 (R&D Systems, Wiesbaden, Germany) on day 10. After 14 days in culture, a third stimulation was performed. After a total of 15 days in culture, the peptide-stimulated T cells were tested for specificity by intracellular IFN-γ FACS.

**Intracellular cytokine staining**

To assess the capacity of T cells to produce IFN-γ upon recognition of a specific target, intracellular IFN-γ staining was performed. Target cells used in both the experimental and the standard groups included hTERT-peptide-loaded DCs. 2 h after start of stimulation 1 μL/mL Brefeldin A (Sigma-Aldrich, St. Louis, USA) was added to the cultures. After 10 h cells were fixed and permeabilized with Inside Fix and Inside Perm (BD Biosciences, San Jose, USA) and stained with the following intracellular and cell surface markers: PE-Cy5-CD3, PE-CD8, PerCP-Cy5.5-CD4 and ECD-CD45 (Beckman Coulter, Krefeld, Germany) and IFN-γ-FITC (Miltenyi Biotec, Bergisch-Gladbach, Germany). Samples were analyzed on a FACS-Calibur flow cytometer. Gating for cytokine was based on a positive T cell control sample which had been stimulated with staphylococcal enterotoxin B (SEB) or staphylococcal enterotoxin A (SEA) (Sigma-Aldrich, St. Louis, USA). T cells incubated with non-peptide loaded DC were used as negative controls.

### Table 1. Patients’ characteristics.

<table>
<thead>
<tr>
<th>Patient</th>
<th>Age (years)</th>
<th>Tumor histology</th>
<th>Tumor stage</th>
<th>Therapy</th>
</tr>
</thead>
<tbody>
<tr>
<td>A</td>
<td>74</td>
<td>Squamous cell</td>
<td>IV</td>
<td>Carboplatin/Paclitaxel</td>
</tr>
<tr>
<td>B</td>
<td>68</td>
<td>Adenocarcinoma</td>
<td>IV</td>
<td>Carboplatin/Docetaxel</td>
</tr>
<tr>
<td>C</td>
<td>55</td>
<td>Adenocarcinoma</td>
<td>IIB</td>
<td>Carboplatin/Vinorelbine</td>
</tr>
<tr>
<td>D</td>
<td>60</td>
<td>Squamous cell</td>
<td>IIB</td>
<td>Carboplatin/Paclitaxel</td>
</tr>
<tr>
<td>E</td>
<td>72</td>
<td>Large cell</td>
<td>IV</td>
<td>Carboplatin/Docetaxel</td>
</tr>
<tr>
<td>F</td>
<td>69</td>
<td>Squamous cell</td>
<td>IIB</td>
<td>Carboplatin/Vinorelbine</td>
</tr>
<tr>
<td>G</td>
<td>73</td>
<td>Squamous cell</td>
<td>IIIA</td>
<td>Carboplatin/Docetaxel</td>
</tr>
</tbody>
</table>

### Table 2. hTERT-derived T cell epitopes.

<table>
<thead>
<tr>
<th>Epitope</th>
<th>Sequence</th>
<th>Restricting MHC allele</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>I540</td>
<td>ILAKFLHWL</td>
<td>HLA-A2</td>
<td>(23)</td>
</tr>
<tr>
<td>V461</td>
<td>VYGFVRACL</td>
<td>HLA-A24</td>
<td>(12)</td>
</tr>
<tr>
<td>Y325</td>
<td>YAETKHFLY</td>
<td>HLA-A1</td>
<td>(24)</td>
</tr>
<tr>
<td>R672</td>
<td>RPGLLGASVGLLDDI</td>
<td>DR1, DR7, DR15</td>
<td>(25)</td>
</tr>
</tbody>
</table>
Cytotoxicity assay

T cells were tested for specific cytotoxicity against T2 cells loaded with hTERT peptides. Target cells (1 × 10⁵) were labeled with 0.1 mCi ⁵¹Cr and mixed with various numbers of effector cells to give E:T ratios of 40:1, 20:1, 10:1 and 5:1. Target cells incubated in complete medium or 5% Triton X-100 (Sigma) were used to determine spontaneous and maximal ⁵¹Cr release, respectively. After 4 h, supernatants were collected and radioactivity was measured on a gamma counter. The mean percentage of specific lysis of triplicate wells was calculated as 100× (experimental release-spontaneous release)/(maximal release-spontaneous release).

Results

Expression of surface maturation markers on dendritic cells

Maturation and costimulatory molecules like CD80, CD83 and CD86 are essential for DCs to induce an effective cytotoxic T-lymphocyte (CTL) response. Because alteration in surface protein expression is an important step in the maturation of DCs, we investigated whether the TLR7/8-agonist in combination with sCD40L had an additive effect on the expression of DC costimulatory molecules.

After culture in GM-CSF and IL-4 adherent PBMCs showed typical DC morphology including dendrites extending in several directions. Based on forward-side-scatter dot plots these large granular cells were gated as DC (Figure 1A). The immature DCs were either activated by the standard cytokine cocktail (cDC) or first primed for 2 h with the CD7/8-agonist and then matured for an additional 20 h with the sCD40L (ligDC). The PBMC-derived large granular cells showed upregulation of CD80, CD86 and CD83 indicating maturation of DCs. However DCs activated by the TLR7/8-agonist and sCD40L displayed lower levels of costimulatory molecules than DCs matured by the standard cytokine cocktail. Almost all DCs matured by the cytokine cocktail upregulated CD80, CD83 and CD86 (Figure 1B), but only 50% of DCs that had been matured by the TLR7/8-agonist and sCD40L upregulated CD83 and CD86 (Figure 1C). Though CD80 was upregulated in all DCs, this surface marker was less intensely expressed in the DCs matured by the TLR7/8-agonist and sCD40L. These results are representative for 5 experiments performed in patients A, B, C, D and G. The findings show that the expression of surface markers analyzed in this study is more efficiently induced by the cytokine cocktail than by TLR7/8-agonists combined with sCD40L.

Depletion of regulatory T cells

As regulatory T cells prevent in vitro expansion of tumor-reactive T cells, we first depleted CD4⁺CD25⁺ regulatory T cells in non-adherent PBMC. As exemplified in HLA-A‘02 patient G, depletion of regulatory T cells resulted in 0% CD4⁺CD25⁺ regulatory T cells (Figure 2). In 3 analyzed patients this approach resulted in a mean of 0.1% regulatory T cells in the non-adherent PBMC being subsequently stimulated with hTERT-peptide loaded DC.

Activation of hTERT-specific T cell responses

Before analysis of hTERT-specific T cell reactivity a positive control for the T cells’ ability to produce cytokines was performed. The T cell responses to the strong immunogenic stimuli SEA and SEB, respectively, were analyzed on non-adherent, T<sub>reg</sub> depleted PBMC of all NSCLC patients (Figure 3A). The specific IFN-γ production of CD8⁺ T cells was detected within a range of 4.1% to 14.7% and a median of 7.8% (Figure 3B).

Detection of hTERT-specific cellular immune responses was performed on non-adherent, T<sub>reg</sub> depleted PBMC of NSCLC patients after in vitro stimulation with autologous hTERT-peptide loaded DC. The peptides used for DC pulsation were selected following published data after MHC class I and II complex analysis (Table 3). Recognition of the HLA-matched hTERT peptides by non-adherent, T<sub>reg</sub> depleted PBMC was analyzed using functional flow cytometry of the peptide-induced intracellular IFN-γ production. Target cells used in both the experimental ligDC and the standard cDC groups included hTERT-peptide-loaded DCs. Non-peptide-loaded DCs were used as negative controls. The percentage of hTERT-specific T cells was calculated by subtraction of IFN-γ⁺ T cells in absence of antigen (negative control) from the number of IFN-γ⁺ T cells in presence of hTERT-peptide loaded DC.

With DC that had been matured with the standard cytokine cocktail hTERT-specific IFN-γ production of CD8⁺ T cells could be induced in PBMC samples of 4 (A-D) out of 7 patients. Using this standard approach, 0.5-1.5% IFN-γ⁺ CD8⁺ T cells could be activated (Figure 4).

Interestingly, the most intense T cell response was observed in PBMC samples of patient A. Due to the patient’s HLA-type the DCs were loaded with both the HLA-A2 restricted I540 and the HLA-A24 restricted V461 hTERT-peptide.

In 1 additional patient (F) only a very small population of T cells could be activated comprising of 0.1% T cells with hTERT-reactivity.

Since there is strong correlation between the intracellular IFN-γ cytokine-staining assay and the ⁵¹Cr-release assay we used the intracellular cytokine staining in the present study as a surrogate marker for cytolytic activity (23). To confirm the function of the activated IFN-γ⁺ hTERT-specific T cells a cytotoxicity assay was done in patient F. However, in this patient T lymphocytes that had been stimulated with hTERT-peptide pulsed cDC did not lyse hTERT-peptide loaded T2 target cells (data not shown).
In the remaining 2 patients (E, G) immunophenotypic analysis of PBMC samples could not reveal any activation of hTERT-specific IFN-γ+ T cell responses after stimulation with cytokine matured DC.

In 2 patients (A, B) in whom the hTERT-specific T cell response had already been activated by the standard cytokine approach, the reaction to hTERT could be further intensified by TLR7/8-agonist and sCD40L matured DCs resulting in 0.9% and 2.0% IFN-γ+CD8+ T cells, respectively (Figure 5A,B).

In PBMC samples of patient C and patient E a small population of 0.1% IFN-γ+CD8+ T cells could be activated by hTERT-peptide loaded ligDC (Figure 6). This population could
Figure 2. Depletion of CD4^+CD25^+ regulatory T cells in patient G: before depletion of CD4^+CD25^+ T cells 3.5% regulatory T cells were present in non-adherent PBMC of patient G (A). After depletion of CD4^+CD25^+ T cells no T_{reg} were detectable (B).

Figure 3. Direct ex vivo FACS analysis of intracellular IFN-γ release by peripheral blood T-cells in responses to SEA/SEB. A, Representative data from patient B; B, Detailed results from all 7 patients after SEA (patient A-E) and SEB (patient F-G) stimulation in NSCLC patients.

<table>
<thead>
<tr>
<th>Patient</th>
<th>HLA-A</th>
<th>HLA-DR</th>
<th>MHC class I peptide</th>
<th>MHC class II peptide</th>
</tr>
</thead>
<tbody>
<tr>
<td>A</td>
<td>2,24</td>
<td></td>
<td>I540, V461</td>
<td>R672</td>
</tr>
<tr>
<td>B</td>
<td>24</td>
<td>7,15</td>
<td>V461</td>
<td>R672</td>
</tr>
<tr>
<td>C</td>
<td>2</td>
<td>7</td>
<td>I540</td>
<td>R672</td>
</tr>
<tr>
<td>D</td>
<td>2</td>
<td>7</td>
<td>I540</td>
<td>R672</td>
</tr>
<tr>
<td>E</td>
<td>1</td>
<td>7</td>
<td>Y325, V461</td>
<td>R672</td>
</tr>
<tr>
<td>F</td>
<td>1,24</td>
<td>1,15</td>
<td>I540</td>
<td>R672</td>
</tr>
<tr>
<td>G</td>
<td>2</td>
<td>1,15</td>
<td>I540</td>
<td>R672</td>
</tr>
</tbody>
</table>
Figure 4. Activation of T cell responses to hTERT in T<sup>reg</sup> depleted PBMC of NSCLC patients after weekly stimulations with peptide loaded cDC. Figure shows the CD8/IFN-γ profile of CD3 gated lymphocytes after incubation of T cells with hTERT-peptide loaded autologous target cells/DC (dahed bars). Responses against non-peptide loaded targets/controls/DC serve as negative controls (black bars).

Figure 5. Increased activation of hTERT-specific T cells after stimulation of T<sup>reg</sup> depleted PBMC with peptide-loaded ligDC. Flow cytometric analysis of peripheral blood CD8<sup>+</sup> T cells specifically producing IFN-γ in response to hTERT-peptides loaded targets. Target cells with no peptide were used as controls. The T cells were either stimulated by cytokine (cDC) or by sCD40-and TLR7/8-ligand (ligDC) matured, hTERT-peptide loaded DC. IFN-γ, interferon gamma. A, Representative data from patient A. Frequencies of CD8<sup>+</sup> T cells that produced IFN-γ are shown as percentages of total numbers of CD8<sup>+</sup> T cells; B, In patient A and B the frequency of T cells that produced IFN-γ in response to hTERT-peptides was higher in T cells that had been stimulated with hTERT-peptide loaded ligDC than in T cells that had been stimulated with cDC.

Figure 6. Very low (patient C and E) and no hTERT-specific activation signal, respectively (patient D, F, G). The T cells were stimulated weekly by hTERT-peptide loaded ligDC. Figure shows the flow cytometric analysis of peripheral blood CD8<sup>+</sup> T cells specifically producing IFN-γ in response to hTERT-peptides loaded targets. Target cells with no peptide were used as controls. IFN-γ, interferon gamma.
not be expanded and functionally tested. Lacking functional data we cannot say if this CD8+ T cell population is recognizing hTERT expressing target cells.

In PBMCs of patients D, F and G there was no hTERT-specific activation signal after stimulation of T cells with hTERT peptide loaded ligDC (Figure 6).

Taken together, hTERT-specific T cells could be activated in PBMC samples of 4 patients by usage of the standard cytokine cocktail. In 2 of these patients the percentage of hTERT-specific T cells could be further increased by DC that had been matured with TLR7/8-agonist and sCD40L.

Additional functional assays were prevented by the insufficient T cell numbers available. Although in the present study hTERT-specific T cells were activated, the total number of T cells decreased. Starting with a median of 6×10^6 T cells, there was a loss of CD8+ cells resulting in a median of 1.3×10^4 T cells after 15 days of culture.

**Discussion**

In present vaccination and adoptive T cell transfer studies activation of cancer-specific T cell responses is associated with a favourable prognosis. In HLA-A2+ patients with advanced NSCLC the estimated overall survival after vaccination with hTERT peptides was 30 months for immunological responders vs. 4 months for non-responders (15) demonstrating a survival advantage in patients with a hTERT-specific immune response. Since the induction of tumor-specific CD8+ T cell response correlates with anti-tumor efficiency, improved protocols for the activation of tumor-specific CD8+ T cell responses are urgently needed.

Regulatory T cells suppress tumor-specific T cell responses and thereby play a key role in cancer immune escape. In patients with metastatic urothelial carcinoma the NYESO-1 directed T cell responses were augmented by Treg depletion (27). Also in cancer patients WT1-specific CD8+ T cell responses could be activated in vitro only after depletion of Treg (20). In accordance with these data in the present study we have stimulated PBMC with peptide loaded DC after depletion of Treg. Following the depletion of regulatory T cells hTERT-specific T cell reactivity could be generated. However, CD25 is also expressed on activated CD4+ T cell subsets. Because of that the depletion of CD4+CD25+ T cells potentially impairs the CD4+ helper T cell response leading to a decreased activation of the hTERT-specific CD8+ T cells. This might especially be important in those patients in whom the non-adherent PBMC were stimulated with MHC class II restricted peptides.

Boosting pre-existing, naturally occurring CD8+ T cell responses directed against class-I MHC-restricted peptides of tumor antigens, represents a primary goal of both active and passive immunotherapy. In a previous study the frequency of circulating hTERT-specific T cells detected in lung cancer patient upon diagnosis was 1×10^-7-4×10^-7 peripheral blood CD8+ cells (28). In the present study, hTERT-specific IFN-γ production of CD8+ T cells could be induced in vitro by different DC maturation cocktails in PBMC samples of 4 out of 7 patients with advanced NSCLC. In these patients, hTERT-specific T cell responses between 0.5% and 2.1% could be activated potentially indicating expansion of base line hTERT-specific T cells. This expansion process could be monitored by hTERT-tetramers.

These data are in accordance with other groups aiming at the activation of hTERT-specific T cells. Filaci et al. (24) demonstrated I540-specific activation of CD8+ T cell responses in patients with different types of cancer after in vitro stimulation with peptide-loaded antigen presenting cells (APC). Generation of p540 peptide-specific CTL lines was tested in 16 cancer patients, but CTL lines could be generated in 11 patients (68%) only, in particular from 8 of 12 prostate cancer patients, 1 of 2 lung cancer patients and 2 patients affected with gastric and liver cancer, respectively (24).

In patients with multiple myeloma a single ex vivo peptide stimulation led to activation of 0.3-0.5% of hTERT-specific cells, but this effect occurred to be limited to only 2 of 27 patients (25).

In 6 out of 8 lymphoma and solid tumor patients 0.5% I540-specific CD8+ cells were identified after three in vitro stimulations of PBMC with peptide loaded APC. Four ex vivo stimulations were necessary to identify 1-3% I540-specific CD8+ T cells (29).

Though we and others have reproducibly activated TAA-specific T cell using dendritic cells as APCs (22) in all other studies mentioned above different types of peptide loaded APC were used. All approaches were finally activating similar quantities of hTERT-specific T cells. Since the generation of DC is both time consuming and laborious the use of peptide loaded PBMCs as stimulators might be advantageous.

To further analyze the functionality of the activated hTERT-specific T cells we performed a cytotoxicity assay in patient F. In this assay we assessed whether the hTERT-reactive T cells detected by measuring specific, intracellular IFN-production mediated cytotoxicity to hTERT+ target cells. However, the in vitro PBMC derived hTERT-specific T cells of this patient did not lyse the hTERT expressing target cells. This data might be related to the very low percentage of hTERT reactive T cells since the number of IFN-γ secreting T cells is closely correlated to their cytolytic activity (30). To detect very low numbers of cytolytic T cells analysis by Flow cytometry of cytolytic makers like granzyme and perforin could be performed in future studies.

Further functional analyses were prevented by the limited absolute number of T cells. After 15 days of in vitro stimulations the number of T cells dropped from 6×10^6 T cells to a median of 1.3×10^4 T cells. This drop was to some extent expected within
the initial activation period. Most likely apoptosis of T cells not recognizing their specific antigen is the leading cause. However, also hTERT-specific T cells might have died within these 15 days due to exhaustion or culture conditions not facilitating survival of TAA-specific T cells. Allogeneic feeder cells like lymphoblastoid cell lines might be of better use for T cell expansion (22). These data indicate that for both clinical adoptive transfer studies as well as for further functional analyses the initial activation of hTERT-specific T cells need to be combined with efficient T cell expansion protocols (31).

These studies show, that by application of the standard cytokine approach hTERT-specific T cells can be generated in vitro in a subset of patients only. Although DCs in our study were phenotypically less matured by a TLR7/8-agonist in combination with a sCD40L, in PBMC samples of 2 patients the number of hTERT-specific T cells was improved. Other authors have previously shown the in vitro effectivity of this combination of a TLR7/8-agonist and the maturation signal sCD40L activating MART-1-specific CD8+ T cells with high functional avidity and enhanced tumor cell recognition (17). These findings were induced by high-levels of IL-12 expression. IL-12 produced by the TLR7/8-primed DC polarized CD4+ T cells for Th1 cytokine production leading to MART-1-specific CD8+ T cells (17). IL-12 being expressed by the TLR7/8-agonist and sCD40L primed DC greatly enhances T cell functional avidity (32). The analysis of the DC in our study was restricted to the expression of a maturation marker and costimulatory molecules. Only 50% of DCs that had been matured by the TLR7/8-agonist and sCD40L upregulated CD83 and CD86. Although CD80 was also upregulated in all DCs, this surface marker was expressed less intensely. However, the secretion of IL-12 is crucial for the induction of specific T-cell responses and may characterize the generated DC in more detail rather than expression of surface maturation molecules. In addition the adherent cells that we have designated DC after culture in IL-4 and GM-CSF could be further characterized by flow cytometry for specific DC markers like CD11c.

Further improvement of DC T cell stimulatory capacity might be achieved by the combination of cytokines with CD40- and TLR7/8 ligands. Paustian et al. (33) have shown that DC activated with IFN-gamma in combination with TLR4-, TLR7/8-agonists and CD40-ligands were capable of responding with strong IL-12 production. This quadruple activating cocktail was not tested for its capacity to sensitize T cells but further studies are needed to determine the best formulation for DC maturation. In a clinical phase I-study in patients with advanced malignant melanoma a subcutaneous vaccine consisting of melan-A-peptide, in combination with the TLR9L CPG 7909 and montanide ISA-51 stronger activated CD8+ T cell responses than any other vaccine induced T cell reaction in men (34).

Summarizing these data, even with the DC maturation protocols used in this study it was not possible to increase the number of immune responders. Only in PBMCs of a subgroup of lung cancer patients receiving standard platinum containing chemotherapy hTERT-specific T cells could be activated in vitro. Potentially these T cells can be isolated and expanded facilitating adoptive T cell transfer for treatment of tumor cells remaining after standard chemotherapy. In addition, hTERT-specific vaccinations might be simultaneously applied to first line chemotherapy protocols. The results of our study indicate, that following chemotherapy hTERT-specific T cell responses can be activated in vitro in a subgroup of NSCLC patients only. This activation by peptide pulsed CDC can be further increased by ligation. We have used a set of hTERT peptides covering the MHC-class I molecules HLA A1, A2, A24 and the MHC-class II molecules HLA DR1/7/15. However, in our study the number of patients with these HLA-types was limited. Experimental settings using overlapping peptides covering the whole hTERT protein for T cell activation would potentially be advantageous. In such a study patients of may be all HLA-types could be included leading to high numbers of patients analyzed.

Naturally occurring T cells against multiple TAA are a common phenomenon in tumor patients. In lung cancer patients the number of TAA-reactive T cells was greatly different between the MAGE-A1, MAGE-A3 and hTERT peptides (27). Following these findings one could expect T cell reactivity to other TAA in those patients who in the present study did not activate hTERT-specific T cell reactivity. Based on this assumption recent immunotherapy studies are investigating multiepitope peptides for T cell activation.

We have not analyzed the frequency of base line T cell reactivity to hTERT. In a previous study the frequency of circulating hTERT-specific T cells was found significantly higher in newly diagnosed lung cancer patients as compared with healthy individuals (27) but varied widely. Upon diagnosis T cell responses to hTERT-peptides were detected in up to 40% of lung cancer patients only. Potentially in our study patients not activating hTERT-specific T cell in vitro did not have any base line hTERT-specific T cell reactivity.

The identification of appropriate immune surrogates of tumor-specific T-cell activation is still pending. A candidate predictive biomarker that identifies patients being able to activate hTERT-specific immune responses in vitro might be the presence of preexisting hTERT-specific T-cells. These tumor-specific T cells should have long term memory, lytic function, proliferative capacity, the ability to migrate to the tumor sites and could potentially be detected by advanced tetramer staining technologies.

Acknowledgements

Disclosure: The authors declare no conflict of interest.
References

27. Horn T, Grab J, Schusdziarra J, et al. Antitumor T cell responses in bladder cancer are directed against a limited set of antigens and are modulated by regulatory T cells and routine treatment approaches. Int J Cancer 2013. [Epub ahead of print].