Polyfunctional tumor-reactive T cells are effectively expanded from non-small cell lung cancers, and correlate with an immune-engaged T cell profile in situ

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Abstract

Non-small cell lung cancer (NSCLC) is the second most prevalent type of cancer. With the current treatment regimens, the mortality rate remains high. Therefore, better therapeutic approaches are necessary. NSCLCs generally possess many genetic mutations and are well infiltrated by T cells (TIL), making TIL therapy an attractive option. Here we show that T cells from treatment naive, stage I-IVa NSCLC tumors can effectively be isolated and expanded, with similar efficiency as from normal lung tissue. Importantly, 76% (13/17) of tested TIL products isolates from NSCLC lesions exhibited clear reactivity against primary tumor digests, with 0.5%-30% of T cells producing the inflammatory cytokine Interferon (IFN)-γ. Both CD4⁺ and CD8⁺ T cells displayed tumor reactivity. The cytokine production correlated well with CD137 and CD40L expression. Furthermore, almost half (7/17) of the TIL products contained polyfunctional T cells that produced Tumor Necrosis Factor (TNF)-α and/or IL-2 in addition to IFN-γ, a hallmark of effective immune responses. Tumor-reactivity in the TIL products correlated with high percentages of CD103⁺CD69⁺CD8⁺ T cell infiltrates in the tumor lesions, with PD-1hiCD4⁺ T cells, and with FoxP3⁺CD25⁺CD4⁺ regulatory T cell infiltrates, suggesting that the composition of T cell infiltrates may predict the level of tumor reactivity. In conclusion, the effective generation of tumor-reactive and polyfunctional TIL products implies that TIL therapy will be a successful treatment regimen for NSCLC patients.
INTRODUCTION

Non-small cell lung cancer (NSCLC) is one of the most prevalent types of cancer worldwide. With the current treatment regimens, the 5-year survival rate of patients suffering from NSCLC is still limited. Because of the striking effects of immunotherapy in melanoma (1–4), recent efforts are directed towards treatment with immunotherapy for NSCLC patients. NSCLC are considered immunogenic, because they often contain high numbers of somatic mutations (often induced by smoking tobacco) (5), and are well infiltrated by T cells (6–8). The functionality of these tumor infiltrated T lymphocytes (TILs) has, however, not been well documented. Treatment with anti-PD-1 is currently standard of care for stage IV NSCLC patients with distant metastasis. This therapy resulted in tumor regression in 17-30% of the patients (9–13). The majority of patients with NSCLC thus does not benefit from immune checkpoint blockade. Tumors in these patients utilize different immunosuppressive pathways (14) that need to be overcome in order for immunotherapy to work. Therefore, additional immunotherapy interventions should be explored for the treatment of NSCLC patients, possibly in combination with the existing anti-PD-1 therapy.

One therapeutic opportunity for intervention is the infusion of TILs. Adoptive transfer of *ex vivo* expanded TILs has proven highly effective for stage IV melanoma patients (15), with impressive 50% overall response rates in pretreated patients (1, 2). Of these melanoma patients, 10-20% experience durable complete remission (1, 2). Tumor-reactive T cells were already detected in the mid 1990s in NSCLC lesions (16–18), and a clinical effect of TIL therapy for stage IV NSCLC patients has been
reported, albeit with very minor improvements in survival (19). The protocols to
culture and expand TILs from tumor lesions, and the read-outs to define tumor-
reactive T cells substantially improved since that time. Therefore, the efficacy to grow
tumor-reactive TIL products from NSCLC lesions should be re-assessed, both in
terms of cell expansion and the presence of cytokine-producing TILs in response to
tumors. Furthermore, it is yet to be determined whether a specific T cell profile in
tumor lesions correlates with the level of tumor reactivity of expanded TIL products.

Here, we show that most TIL products contain tumor-reactive T cells. In particular
TIL products with high tumor reactivity are polyfunctional. Furthermore, tumor
reactivity of the expanded TIL product correlated with the composition of the T cell
compartment in the tumor lesions. We conclude that the generation of NSCLC-
specific TIL products for therapeutic purposes is feasible and should be reconsidered
for clinical application.
Materials and methods

Patient cohort and study design

Between June 2015 and June 2017, 25 treatment-naive NSCLC patients were included in this study. Samples from 2 patients were excluded because of logistic issues. Table 1 depicts the patient characteristics of the remaining 23 donors. The cohort consisted of 10 male and 13 female donors between the age of 38 and 79 years (average 66.1 years) with clinical stage Ia-IVa according to the TNM7 staging system for NSCLC, based on tumor size, nodal involvement and level of metastasis. All but two patients had a history of smoking.

The study protocol was designed to define the phenotype of TILs isolated from treatment naive NSCLC compared to non-tumorous lung tissue (N=16), to explore the capacity to expand TILs (N=17) and to measure the presence of tumor-reactive T cells within TIL cultures (N=17). The study was performed according to the Declaration of Helsinki (seventh revision, 2013), and executed with consent of the Institutional Review Board of the Netherlands Cancer Institute-Antoni van Leeuwenhoek Hospital (NKI-AvL), Amsterdam, the Netherlands. Donor tissues were obtained directly after surgery, transported at room temperature in RPMI medium 1640 (Gibco) containing 50ug/ml gentamycin (Sigma-Aldrich), 12.5ug/ml Fungizone (Amphotericin B, Gibco) and 20% fetal bovine serum (FBS) (Bodego), weighed and processed within four hours. Distal healthy lung tissue as defined by the pathologist was taken as far away as possible from the tumor lesion.
Table 1: Patient characteristics

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*American Joint committee on Cancer: Lung Cancer Staging, 7th edition, 2009

Tumor stage and differentiation, and weight of the obtained tissues for this study are shown in Table 2. Tumor size ranged from T1a-T4 according to the TNM7 system, and specified as adenocarcinoma (n=14), squamous carcinoma (n=5) and as NSCLC not otherwise specified (NOS, n=4). On average 1420 mg tumor tissue was obtained, ranging from 23-9590 mg (Table 2).
Table 2: Tumor characteristics

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Tissue digestion

The first five donors were used to optimize the digestion protocol for the highest yield of live CD45+ cells in single cell digests. Briefly, freshly isolated tumorous and non-tumorous lung tissue were finely chopped, and incubated for 45 min shaking at 37°C in IMDM (Gibco) containing 30IU/ml collagenase IV (Worthington), 12.5μg/ml DNAse (Roche), and 1% FBS. The digest was pelleted at 360g for 10 min, and
resuspended in FACS buffer (PBS containing 2% FCS and 2mM EDTA). Digest was filtered first through a tea mesh and then over a 100um filter. After red blood cell lysis for 15 min at 4°C with 155 mM NH₄Cl, 10 mM KHCO₃, 0.1 mM EDTA (pH 7.4), live and dead cells were manually counted with trypan blue solution (Sigma) on the hemocytometer. From six donors, the tumor tissue was divided in three regions. A sample from each region was separately processed for TIL culture. The remaining tissue was combined for the total tumor digest. After digestion, 1-2x10⁶ live cells were used for flow cytometry analysis, and 1-3x10⁶ live cells were used for TIL cultures. The remaining digest was cryo-preserved until further use.

*T cell expansion*

TIL cultures were performed as previously described for melanoma-derived TILs (1, 2). Briefly, 2-3 wells containing 0.5-1x10⁶ live cells from the tissue digest were cultured for 10-13 days in 24 wells plates in 20/80 T-cell mixed media (Miltenyi) containing 5% human serum (HS) (Sanquin), 5% FBS, 50μg/ml gentamycin, 1,25 μg/ml fungizone, and 6000 IU human recombinant (hr) IL-2 (Proleukin, Novartis) (pre-Rapid Expansion Phase; pre-REP). Medium was refreshed on days 7, 9 and 11. Wells were split when a monolayer of cells was observed in the entire well. When more than 30% of the cells stopped dividing (determined as rounding up of cells), cells were harvested, counted and prepared for an additional culture period of 10-13 days (REP). 1-3 times 2x10⁵ live cells/well (1-3 wells/donor) were co-cultured with 5-10x10⁶ irradiated PBMCs pooled from 10 healthy donors in 24 wells, 30ng/ml anti-CD3 antibody (OKT-3) (Miltenyi Biotec) and 3000IU/ml IL-2. Typically, cells were passaged at day 5, 7, 9 and 11 and harvested, washed and counted on day 10-13, based on visual assessment of the T cell proliferation state as above. Cells were either
tested immediately for reactivity, or cryo-preserved in IMDM containing 10% DMSO (Corning) and 30% FBS until further use.

*T cell activation*

After thawing and live cell counting as described above, pre-REP and REP T cells were pre-stained in FACS buffer with anti-CD4 BUV496, anti-CD8 BUV805 (BD Biosciences) for 30 min at 4°C. Cells were washed twice, one time in FACS buffer and then in 20/80 T-cell mixed media (Miltenyi). 100.000 live T cells were co-cultured with 200.000 live tumor digest cells or normal lung digest cells for 6h at 37°C, were stimulated with 10ng/ml PMA and 1ug/ml Ionomycin (SigmaAldrich), or cultured with T-cell mixed media alone. Brefeldin A (Invitrogen) was added after 30 min of co-culture.

*Flow cytometry*

For ex vivo analysis of T cells in the tissue digests, cells were washed in FACS buffer, and then stained in FACS-buffer for 30 minutes at 4°C with the following antibodies: anti-CD3 PerCp-Cy5.5, anti-CD279 FITC, anti-CD56 BV605, anti-CD27 BV510, anti-CD127 BV421, anti-CD103 PE-Cy7 and anti-CD25 PE (all from Biolegend), and with anti-CD8 BUV805, anti-CD45RA BUV737, anti-CD4 BUV496, and anti-CD69 BUV395 (from BD Biosciences). Live/dead fixable near IR APC-Cy7 (Invitrogen) was included for dead cell exclusion. Cells were washed twice with FACS buffer and fixed for 30 minutes with the Perm/Fix Foxp3 staining kit (Invitrogen) according to the manufacturer’s protocol using anti-Foxp3 Alexa647 (Biolegend). Cells were resuspended in FACS buffer and passed through a 70μm single cell filter prior to flow cytometry analysis (LSR fortessa, BD biosciences). The
first six donors were used to optimize the staining panel and cytometer settings, by antibody titration and the use of single stains in combination with the minus one fluorochrome method (MOF).

T cells from pre-REP or REP cultures were stained prior to T cell activation with anti-CD4 BUV496, anti-CD8 BUV805 as described above. After the full activation protocol, cells were washed twice with FACS buffer and stained with anti-CD154 BV510 and anti-CD279 BV421 (Biolegend) and Live/dead fixable near IR APC-Cy7 in FACS buffer for 30 minutes at 4°C, and washed twice in PBS. Cells were then fixed and stained with anti-CD137 PE-Cy7, anti-IFN-γ PE, anti-TNF-α Alexa488 and anti-IL-2 APC (Biolegend) with the Cytofix/CytoPerm staining kit (BD Biosciences) according to the manufacturer’s protocol. Cells were washed in FACS buffer and passed through a 70uM single cell filter prior to acquisition with the LSR fortessa (BD). Flow cytometry settings were defined for each patient material using single stainings for each antibody. To each flow cytometry analysis of patient material, a standardized sample of PBMCs pooled from ten healthy donors that was cryo preserved prior to the start of the study was included. Data analysis was performed with Flowjow Star 10.1.

Statistical analysis
Statistical analyses were performed with Graphpad Prism 7.2. Compiled data are shown as paired data points for each patient, or as single data points with box and whiskers showing maximum, 75th percentile, median, 25th percentile, minimum and mean, unless otherwise indicated in the legend.
Overall significance of differences were calculated with repeated measurement paired one-way ANOVA test. If differences were confirmed (p<0.05), significance between two data points was calculated using paired Student’s t test, with the p value cut-offs of *= p<0.05; **=p<0.01; and ***=p<0.001. If Student’s t test showed p values ≥0.05, p value marking was omitted in panels. Correlations were calculated using Pearson's correlation in combination with linear regression.
RESULTS

High yield of lymphoid cells isolated from NSCLC tumor lesions

We first determined the efficacy of isolating TILs from NSCLC lesions that underwent lobectomy. 23 patients from treatment-naive stage Ib-IVa NSCLC patients suffering from non-squamouse (n=14), squamous (n=5), or from NSCLC not otherwise specified (n=4) were included in this study (Table 1). To evaluate the TIL isolation and expansion procedure from the tumor, we also isolated normal lung tissue from the same patients that was harvested as far away as possible from the tumor lesion.

From the tumor digests, we obtained on average $33.6 \times 10^3$ viable cells/mg tissue, which was comparable to the yield from normal lung tissue digests, with on average $51.2 \times 10^3$ viable cells/mg tissue (Fig. 1A). In line with previous studies (7, 8, 20), we detected high numbers of CD3^+CD56^- T cells in both tumor and normal lung tissues, with 23.7±16.9% and 15.5±13.1% of live cells in the respective tissue digests. This translated into on average $17.8 \times 10^3$ T cells/mg lung tissue and $9.8 \times 10^3$ T cells/mg tumor tissue (Fig. 1B-D). Of note, sufficient cell numbers for TIL expansion and phenotypic analysis could be isolated even from tumors of <1 cm³ (Table 2).

Effective TIL expansion from NSCLC tumors

The expansion efficiency of TILs highly depends on the tumor type and the culture protocol (1, 2). We here determined the efficiency of TIL expansion from 17 tumor digests with the rapid expansion protocol (REP) established for melanoma TILs (1, 2). For comparison, we included 9 normal lung tissue digests. Tumor and lung tissue digests were cultured for 10-13 days in the presence of IL-2 (pre-REP), followed by a
10-13 day restimulation with the α-CD3 antibody OKT-3 and IL-2 (REP). During pre-REP, CD3+ T cells from tumor and lung digest expanded on average 4- and 6-fold, respectively (Fig. 1E). REP cultures expanded CD3+ T cells from the tumor 470-fold and from the lung 570-fold, resulting in an overall expansion of on average 2000-fold for tumor TILs, with a minimum of 540-fold, and 3000-fold lung tissue derived T cells (Fig. 1E). Thus, even though T cells from normal lung tissue may expand slightly better than those from the tumor, the overall efficiency of expanding NSCLC-derived T cells was comparable to recently reported expansions of melanoma-derived TILs (1, 2).

We also measured the percentages of CD4+ and CD8+ T cells in the pre-REP and REP cultures from tumor digests. CD4+ T cells increased from 14.3± 9.6% ex vivo of total live cells to 43.3±26.5% after REP, and CD8+ T cells increased from 10.3±8.3% to 53.4±27.0% (Fig. 1F). CD8+ T cells expanded generally better than CD4+ T cells in these culture conditions, as revealed by the changes over time in the CD8+/CD4+ T cell ratio (Fig. 1G). In conclusion, both CD4+ and CD8+ T cells from NSCLC were effectively expanded with the REP culture conditions.

The majority of TIL products contain tumor-specific T cells

We next examined the tumor reactivity of expanded TILs. As a read-out, we measured the production of the pro-inflammatory cytokine IFN-γ by expanded TILs upon co-culture with autologous tumor digest. This production was compared to co-culture with normal lung tissue digest, or with medium alone. To distinguish the expanded TILs from the high T cell infiltrates within the tissue digests, we pre-stained the expanded TILs with CD4 and CD8 antibodies prior to co-culture.
IFN-γ producing T cells in response to tumor tissue were detected in 16/17 tumor digests (Fig. 2A,B). On average, 6.3±7.8% of the T cells from the TIL products produced IFN-γ when co-cultured with tumor digests (Fig. 2A,B). We detected similar percentages of IFN-γ-producing T cells within the total CD4+ and CD8+ T cell population, with 5.8±6.3% and 6.5±9.4% of each population, respectively (Fig. 2A,B).

Interestingly, we detected 2.3±2.9% IFN-γ-producing T cells in the TIL products upon co-culture with lung digests, which was higher than those of TILs cultured with medium alone (1.1±1.0%; Fig. 2A,B). The reactivity could be due to undetected tumor cell dissemination, distal tumor antigen presentation by dendritic cells, bystander T cells that are not tumor-specific and thus may also respond to normal lung cells (21). The reactivity to normal lung tissue digests could also reflect innate immune responses by T cells that are triggered in a non-antigen specific manner (22, 23). We only considered TIL products tumor-reactive when the percentage of IFN-γ producing T cells was higher in response to the tumor digest than that to the normal lung digest. 13/17 (76%) of the TIL products fulfilled this requirement (see also below). These data therefore indicate that tumor-reactive T cells are highly prevalent in expanded TIL products from NSCLC.

The percentage of T cells producing IFN-γ varied considerably between donors, ranging from 0.3% to 27.5% (Fig. 2B). This variation in tumor-reactivity could result from high heterogeneity of NSCLC tumors and thus depend on the tumor region that was isolated for TIL expansion. However, we consider this possibility unlikely, because the percentage of IFN-γ+ T cells expanded from whole tumor lysates was
comparable to that of TILs that were expanded from three distinct parts of the same tumor (Fig. 2C). Furthermore, all TIL products produced substantial levels of IFN-γ in response to PMA/ionomycin stimulation (65.8±16.4%; Fig. 2D). There was no significant correlation between this potential to produce cytokines and the tumor-reactivity of TIL products (r=0.11, p=0.663). We therefore conclude that the majority of NSCLC-derived TIL products contain tumor-reactive T cells, but that the proportions thereof are variable.

**IFN-γ-producing tumor-reactive T cells express CD137 (4-1BB) and CD154 (CD40L)**

We next interrogated if measuring the production of IFN-γ alone was sufficient to reflect the actual tumor-reactivity of TIL products. We therefore included the analysis of markers that are rapidly induced upon T cell activation, such as the costimulatory molecules CD137 (4-1BB) (24), and CD154 (CD40L) (25). When expanded TILs were co-cultured for 6h with tumor digest, we found a significant induction of CD137 expression in both CD4+ and CD8+ T cells (Fig. 3A, B, top panels). Furthermore, CD4+ T cells - but not CD8+ T cells - showed increased expression of CD154, a marker that was shown to identify antigen-specific CD4+ T cells in the peripheral blood (25) (Fig. 3A, B, middle panels). CD279 (PD-1) expression instead was not induced under these culture conditions (Fig. 3A, B, bottom panels), suggesting that the tumor-reactive T cells are not (immediately) susceptible to negative regulation by immunological checkpoints.

The majority of IFN-γ producing CD8+ T cells were CD137-positive (Fig. 3C,D).

Combining the expression of CD137 and CD154 with the production of IFN-γ in fact
helped to accurately define the presence of tumor-reactive TILs, in particular when the percentage of IFN-γ production was low, and/or the background levels high. However, we also observed that IFN-γ-negative T cells can express CD137 and CD154 (Fig. 3E,F), suggesting that the actual percentage of tumor-reactive T cells may be higher than estimated based on the production of IFN-γ.

**NSCLC-reactive TILs produce TNF-α and IL-2, and can be polyfunctional**

In addition to IFN-γ, other cytokines such as TNF-α and IL-2 play a critical role in effective T cell mediated immunity (26, 27). This prompted us to determine whether TIL products from NSCLC produced these two cytokines. Interestingly, exposure to tumor digest resulted in significant induction of both TNF-α- and IL-2-producing T cells when compared to co-culture with lung digest or medium alone (Fig. 4A, B). This was observed for both CD8+ T cells, and for CD4+ T cells (Fig. 4A, B).

When we used all three cytokines to define tumor-reactive TILs above background levels (i.e. subtracting the response against lung tissue), again 13/17 TIL products (76.5%) were tumor reactive (Fig. 4B, left panel). We next divided tumor-reactive TIL products into three categories. 4/17 (23.5%) expanded TILs did not produce cytokines above background levels. 8/17 (47.1%) TIL products contained up to 5% of cytokine-producing T cells (intermediate responder, Fig. 4B, left panel). Importantly, 5/17 (29.4%) TIL products contained at least 6%, and up to 31% cytokine-producing T cells above background levels (high responder; Fig. 4B, left panel). This finding indicates that TIL products from NSCLC can contain high levels of tumor-reactive T cells.
Polyfunctional T cells that produce more than one cytokine simultaneously are considered the most potent effector T cells against chronic infections (28, 29). This is also observed for T cell responses against tumors (26, 27). We therefore investigated whether NSCLC-derived TIL products contain these polyfunctional T cells. Strikingly, all high responders contained T cells that produced 2 or 3 cytokines (Fig. 4B, right panel). Of the 8 intermediate responders, also two patients, #7 and #11, contained polyfunctional T cells (Fig. 4B, right panel). Polyfunctional responses to tumor digests were observed for both CD4+ and CD8+ T cells (Fig. 4B, right panel). Interestingly, in the high responders, tumor-reactive T cells were equally distributed between CD4+ and CD8+ T cells (Fig. 4B, right panel), and the overall ratio of CD8+ T cells over CD4+ T cells in REP products of high responders was close to 1, whereas this ratio was not skewed to either T cell subset in the intermediate/no responders (Fig. 4C).

In conclusion, most TIL products contained tumor-reactive CD4+ and CD8+ T cells, and in particular high responders contained polyfunctional T cells, indicative for potent anti-tumoral TIL products generated from NSCLC tumors.

**Tumor-specific alterations in T cell composition correlate with tumor-reactivity of TIL products**

We next investigated whether the high variation in tumor reactivity between TIL products related to specific features of T cells in the original tumor tissue *in situ*. Because the composition of T cell infiltrates is highly variable between individuals (see below), we used the normal lung tissue as base line for each patient to define the tumor-specific T cell signature.
Tumor tissues showed a modest, but significant increase in CD8\(^+\) T cell infiltrates compared to lung tissue, a feature that was not observed for CD4\(^+\) T cells (Fig. 5A). Interestingly, tumor lesions contained significantly more FOXP3\(^+\)CD25\(^{\text{high}}\) CD4\(^+\) T cells (Fig. 5B). These FOXP3\(^+\) CD4\(^+\) T cells did not express the IL-7 receptor alpha chain (CD127), identifying them as bona fide regulatory T cells (30) (Fig. 5B).

The percentages of CD8\(^+\) T cells that express both the integrin CD103 and the retention marker CD69 in the tumor lesion significantly exceeded those in normal lung, which were already high (Fig. 5C). For CD4\(^+\) T cells, we instead observed a drop of CD69\(^+\)CD103\(^+\) expressing cells in the tumor when compared to the normal lung tissue (Fig. 5C). As expected, we found that PD-1\(^{\text{hi}}\) expressing CD4\(^+\) and CD8\(^+\) T cells were substantially increased in the tumor lesions compared to normal lung tissue (Fig. 5D). Thus, tumor tissues contain high levels of FOXP3\(^+\)CD25\(^{\text{high}}\) regulatory CD4\(^+\) T cells, PD-1\(^{\text{hi}}\) CD4\(^+\) and CD8\(^+\) T cells, and high levels of CD69\(^+\)CD103\(^+\)CD8\(^+\) T cells.

We next investigated whether a specific *ex vivo* tumor-specific T cell signature correlated with the level of tumor-reactivity detected in the TIL products. To account for the high variation between patients already in the healthy lung (Fig 5), we used the tumor-specific increase of T cell subpopulations for this comparison (delta percentage; \(\Delta\%)\). The tumor-specific infiltration by CD4\(^+\) or CD8\(^+\) T cells did not correlate with the percentage of cytokine producing TILs (Fig. 6A). However, when examining specific T cell subsets, we found that the percentage of CD69\(^+\)CD103\(^+\) CD8\(^+\) T cells positively correlated with the level of tumor-reactivity of expanded TIL products (Fig. 6B). For CD4\(^+\) T cells, a positive correlation with tumor-reactivity was
found for the percentage of PD-1$^{\text{hi}}$ CD4+ T cells (Fig. 6B,C). Intriguingly, even though regulatory T cells are associated with a poor prognosis for NSCLC patients (17, 18), we found that the presence of tumor-specific FOXP3$^{\text{hi}}$CD25$^{\text{hi}}$ CD4$^{+}$ T cell infiltrates correlated positively with cytokine production of expanded TILs (Fig. 6D). Combined, our data suggest that the tumor-specific signature of CD4$^{+}$ and CD8$^{+}$ T cells and a high number of Treg infiltrates is indicative for strong tumor reactivity in the expanded TIL products.
DISCUSSION

Here we show that tumor-specific TILs can be effectively expanded from the majority of NSCLC patients. Importantly, we found that TIL products that contain high percentages of tumor-reactive T cells produce not only IFN-γ, but also TNF-α and/or IL-2. T cells producing more than one cytokine are considered highly potent (26, 27) and vaccination strategies strive to generate these polyfunctional T cell responses (31). We therefore hypothesize that infusion of TIL products containing polyfunctional T cells should be effective against NSCLC.

Recently, it was shown that tumor-reactive TILs could be expanded from 80% of tested melanoma patients that did not respond to anti-PD-1 treatment (32). 2 out of 12 patients that were subsequently treated with TIL therapy showed objective clinical effects (32). These findings thus indicate that melanoma patients who are refractory to anti-PD-1 can still benefit from TIL therapy, and therefore anti-PD-1 and TIL therapy are complementary immunotherapeutic strategies. This could also be the case for NSCLC patients that did not respond to anti-PD-1 treatment (9–13).

Intriguingly, most NSCLC TIL products contain both tumor-reactive CD8+ T cells and CD4+ T cells. The cytolytic function of CD8+ T cells to control and/or eradicate tumors is well established, and tumor-reactive CD8+ T cells were recently expanded by others from NSCLC tumor digests and from peripheral blood (20, 33, 34). In our study the majority of TIL products contained tumor-reactive CD8+ T cells.

Much less is known about tumor-reactive CD4+ T cells. Recently, neo-antigen specific CD4+ T cells were identified in the majority of tested melanoma lesions (35). Furthermore, two patients that suffered melanoma and metastatic cholangiocarcinoma
received TIL therapy with tumor-reactive CD4+ T cells only, and both patients showed clear clinical benefits from this treatment (36, 37). Thus, CD4+ T cells can substantially contribute to anti-tumoral responses. Whether this effect is indirect through promoting effective CD8+ T cell responses, or direct by cytotoxic CD4+ T cells as proposed in murine tumor models (38, 39) is yet to be defined. These findings combined strongly suggest that in addition to the CD8+ T cells, these tumor-reactive polyfunctional CD4+ T cells could be of value for TIL therapy.

The composition of the tumor-infiltrating T cell compartment correlates with patient survival (40–43). Our study now showed that a correlation also exists between this composition and the likelihood that cytokine-producing tumor-reactive T cells are present in expanded TIL products. In line with previous studies that showed that the presence of CD8+ T cells expressing CD103 and CD69 serves as a favorable prognostic marker for survival (40, 41, 44), we describe here that their presence in the tumor lesion also correlated with high percentages of cytokine-producing CD8+ T cells in the TIL products. This was, however not the case for CD4+ T cells, for which high PD-1 expression correlated with the presence of tumor-reactive T cells in the TIL product.

To our surprise, even though the presence of FOXP3+ T cells in tumors is associated with poor survival (42, 43), we found the highest levels of tumor-reactivity in TIL products when the percentage of Tregs in the tumor lesion was also high. One possible explanation could be that Tregs are recruited to quench ongoing T cell responses against the tumor, such that their prevalence would be proportional to the magnitude of the anti-tumor response (45). Irrespective of the mechanism(s) that contribute to the correlation of high Treg numbers with high percentages of tumor-
reactive T cells, it is tempting to speculate that patients with high Treg numbers may profit most from TIL therapy. Furthermore, even though a validation cohort of our findings is warranted, we propose that the presence of CD69+CD103+ on CD8+ T cells, PD-1hi CD4+ T cells, and high levels of Tregs in the tumor lesions may help identify the patients that are most likely to benefit from TIL therapy.

In conclusion, we here demonstrate that most tumor-reactive TIL products generated from NSCLC tumor lesions contain tumor-reactive T cells. We therefore suggest that TIL therapy should be considered as treatment for NSCLC patients.
FIGURE LEGENDS

Figure 1: Effective isolation and expansion of TILs from NSCLC-derived lesions

Single cell suspensions were obtained from distal lung tissue and tumor sections of 23 donors directly after resection. (A) Numbers of life cells/mg tissue. Dead cells were excluded by trypan blue. Each dot represents one donor, box and whisker plots depict minimum, 25th pct, median, 75th pct and maximum values. (B) Gating strategy of CD56-/CD3+ T cells and lymphocytes, and (C) percentage of CD3+ cells within the lymphocyte population in tumor and lung, to calculate (D) the numbers of T cells/mg tissue by flow cytometry, depicted as in (A). (E) Of 9 normal lung tissues (left panel), and of 17 tumor tissues (right panel), single cell suspensions were cultured for 10-13 days with 6000IU/ml IL-2 (pre-REP), followed by 2 weeks of culture with 30ng/ml anti-CD3, feeder mix, and 3000IU/ml IL-2 (REP). The fold change of total cell count was determined from the number of cells used as input. (F, G) The ratio of CD8+ over CD4+ T cells of the CD3+ T cells was determined by flow cytometry in tumor digests directly ex vivo, and after pre-REP and REP cultures. [Paired student’s T test; * p<0.05; *** p<0.001. If no indication, p≥0.05].

Figure 2: Most NSCLC-derived TIL products contain tumor-reactive T cells

(A, B) REP TILs were pre-stained with CD4 and CD8 antibodies, and cultured for 6 hours in the presence of brefeldin A with medium alone, with digests from normal lung tissue, or with autologous tumor digest. (A) Representative dot plots, and (B) compiled data of 17 donors of IFN-γ producing CD3+ T cells (left panel), by CD4+ T cells (middle panel) and by CD8+ T cells (right panel) as determined by flow cytometry. Each dot represents one donor, box and whisker plots depict minimum,
25\textsuperscript{th} pct, median, 75\textsuperscript{th} pct and maximum values. (C) IFN-γ production of CD4\textsuperscript{+} and CD8\textsuperscript{+} T cells compiled in response to tumor tissue from 6 donors extracted from (B), of which three independent tumor regions (A-C) were isolated to compare the outgrowth of tumor-reactive TILs. (D) IFN-γ production by CD4\textsuperscript{+} and CD8\textsuperscript{+} T cells upon activation with PMA-Ionomycin. [Paired student’s T test; * p<0.05; ** p<0.01; *** p<0.001. If no indication, p≥0.05].

**Figure 3: IFN-γ producing TILs express CD137 and CD40L, but not PD-1.**

REP TILs were cultured for 6 hours in the presence of brefeldin A with medium alone, with digests from normal lung tissue or with autologous tumor digest. (A) Representative histograms of CD137 (4-1BB), CD154 (CD40L), and CD279 (PD-1) expression of CD8\textsuperscript{+} and CD4\textsuperscript{+} TILs that were exposed to lung digest (gray), or to tumor digest (black line). (B) Compiled data of 13 TIL products for CD4\textsuperscript{+} T cells (left panel) and CD8\textsuperscript{+} T cells (right panel). (C) Representative dot plots and (D) compiled data of percentage of T cells that co-express IFN-γ and CD137, CD154, or CD279 of CD4\textsuperscript{+} T cells (left panel) and CD8\textsuperscript{+} T cells (right panel). (E) Representative histograms and (F) compiled data of the percentage of CD137, CD154, and CD279 – expressing CD4\textsuperscript{+} and CD8\textsuperscript{+} T cells within the IFN-γ producing (positive) and negative cells. Each dot represents one donor, box and whisker plots depict minimum, 25\textsuperscript{th} pct, median, 75\textsuperscript{th} pct and maximum values. [Paired student’s T test; * p<0.05; ** p<0.01; *** p<0.001. If no indication, p≥0.05].

**Figure 4: NSCLC-derived TIL products can be polyfunctional**

(A) Percentage of TNF and IL-2 production of T cells (left panel), CD4\textsuperscript{+} T cells (middle panel) and CD8\textsuperscript{+} T cells (right panel) as determined by flow cytometry, n=17
donors. Each dot represents one donor, box and whisker plots depict minimum, 25th pct, median, 75th pct and maximum values. (B) **Left panel:** Percentage of cytokine-producing T cells that were cultured with tumor digest, minus the percentage of cytokine production of T cells stimulated with lung digest (n=17). **Right panel:** Total percentage of cytokine producing T cells. The percentage of cytokine producing CD4+ T cells (dark colors) and CD8+ T cells (light colors) is depicted when exposed to tumor digest (top bar), or to the lung digest (bottom bar). Color coding indicates the production of 1 (blue) 2 (purple), or 3 (yellow) cytokines simultaneously. Each patient is indicated with a number. (D) Ratio of tumor reactive CD8+ over CD4+ T cells, in relation to the percentage of cytokine-producing T cells.

**Figure 5: T cell composition is altered in the tumor lesions**

(A-D) Tumor tissue and of normal lung tissue digests were analysed directly *ex vivo* by flow cytometry to determine the expression of CD3+CD4+ (left panels) and of CD3+CD8+ T cells (right panels) within the lymphocyte population (A), of CD4+CD25+Foxp3+ T cells (B), of CD69+CD103+ T cells (C) and of CD279+ T cells (D). Data are shown as representative dot plots and compiled data of all 17 patients. In (B) a representative histogram is shown of CD127 expression on tumor derived CD25+Foxp3+CD4+ T cells versus conventional FoxP3 CD4+ T cells. [Paired student’s T test; * p<0.05; ** p<0.01; *** p<0.001. If no indication, p≥0.05].

**Figure 6: Tumor-specific T cell infiltrates correlate with cytokine production of the TIL product**

Pearson’s correlation of the percentage of cytokine production of expanded T cells as defined in Fig. 4 with tumor-specific infiltrates of (A) CD4+ (left panel) CD8+ (right
panel) CD3+ T cells, (B) CD69+/CD103+ T cell population within CD4+ (left) and CD8+ (right) T cells, (C) with CD279hi CD4+ T cells (left panel) and CD8+ T cells (right panel), and with (D) Foxp3+/CD25high CD4+ T cells. (** p<0.01,*** p<0.001. If no indication, p≥0.05).

**Supplemental Figure 1**

(A) REP TILs were pre-stained with CD4 and CD8 antibodies, and cultured for 6 hours in the presence of brefeldin A with medium alone, with digests from normal lung tissue, or with autologous tumor digest or PMA/Ionomycine. Representative dot plots of the production of TNF-α and IL-2 of one high and one intermediate donor (same as in Fig. 2A) by total T cells as determined by flow cytometry.
REFERENCES


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De Groot et al: Figure 2

A

Live/Dead

Medium   Lung   Tumor

IFN-γ+

0.65%  0.95%  3.53%

1.31%  3.29%  20.2%

B

Total T cells

CD4+ T cells

CD8+ T cells

Stimuli: Medium   Lung   Tumor

% of IFN-γ+ T cells

C

% of IFN-γ+ T cells

D

PMA/Iono

47.4%

70.8%

% of IFN-γ+ T cells
A Lymphocytes

B CD85

C CD69

D CD103

Lung Tumor

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