Research Article

Characterization of Ex Vivo Expanded Tumor Infiltrating Lymphocytes from Patients with Malignant Melanoma for Clinical Application

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Clinical trials of adoptive transfer of autologous tumor infiltrating lymphocytes (TILs) to patients with advanced malignant melanoma have shown remarkable results with objective clinical responses in 50% of the treated patients. In order to initiate a clinical trial in melanoma, we have established a method for expanding TILs to clinical relevant quantities in two steps with an 8 weeks. Further characterization of expanded TILs revealed an oligoclonal composition of T-cells with an effector memory-like phenotype. When autologous tumor was available, TILs showed specific activity in all patients tested. TIL cultures contained specificity towards tumor cells as well as peptides derived from tumor-associated antigens (TAAs) during expansion procedures.

1. Introduction

The incidence of malignant melanoma is increasing worldwide, and upon dissemination has a very poor prognosis [1]. Only two systemic treatments are approved for disseminated disease and encompass IL-2 based immunotherapy (16% response rate and 6% complete responses) [2] and dacarbazine (6%–15% response rate with no improved survival) [3]. However, results from clinical trials of TIL-based immunotherapy conducted at two centres has shown 50% response rates in patients with advanced disease, and responses were long lasting [4, 5]. TILs were reported to be dominated by CD8+ T-cells and mediate specific killing of autologous tumor in most patients [6]. Information on TAA-derived peptide specificities in TIL has mainly shown the occasional large frequency of MART-1 and gp100 specific T-cell populations. On the other hand, results on the clonotypic and phenotypic composition has been scarce; one publication has revealed a mixed clonal content of TIL by FACS analysis [7], and two recent studies report surface markers identical to memory-like effector T-cells from a limited patient material [8, 9].

In our study, we have analysed TIL characteristics from 17 melanoma patients, whereof five have undergone lymphodepletion and TIL-based ACT with low-dose IL-2.

2. Materials and Methods

2.1. Patients. Patients referred to surgery for primary or recurrent stage III-IV malignant melanoma were eligible for the study. The study protocol was approved by the local ethics committee, and all patients were included after signing informed consent. Tumor material from the patients was obtained from the surgically removed tumour within 30 minutes after surgery.

2.2. TIL Bulk Cultures and Rapid Expansion. The TIL culturing method was adapted from Dudley et al. [10] constituting a two-step expansion process: (I) initiating bulk cultures and (II) rapid expansion of selected bulk cultures with a proliferative potential. Following surgical removal of tumor tissue from patients with MM the tumour sample were cut into 1-2 mm fragments. Fragments were subsequently placed...
individually in 24-well culture plates (Nunc, Denmark) and
maintained in 2 mL of culture medium (CM) containing
RPMI1640 (Invitrogen), penicillin, streptomycin, fungizone
(Bristol-Myers Squibb), 10% human serum (Sigma) and
7300 or 6000 IU/mL IL-2 (Aldesleukin, Novartis). Each frag-
ment initiated an individual TIL culture which was main-
tained separately during subsequent expansion and activa-
tion.

Bulk cultures were selected for further expansion accord-
ing to a rapid expansion protocol (REP). TIL were cocul-
tured with irradiated (40 Gy) allogeneic PBMCs serving as
feeder cells in a ratio of 1:200 in a 1:1 mixture of CM and
AIM-V (Invitrogen) initially with 10% HS, and con-
taining 30 ng/mL OKT-3 (Cilag AG, Suisse) and 7300 or
6000 IU/mL IL-2 (Aldesleukin, Novartis) in upright T-flasks.
REPs for preclinical purposes generally were initiated from
6000 IU/mL IL-2 (Aldesleukin, Novartis) in upright T-flasks.

Gradient Gel Electrophoresis (DGGE). RNA was extracted
using the NucleoSpin RNA II (Macherey-Nagel, Germany).
cDNA synthesis and quantitation of cDNA in each sample
was carried out as previously described [11].

For TCR clonotype mapping, cDNA was amplified using a
primer panel covering the 24 BV region families of the
TCR. Resulting PCR products are suited for DGGE
[12, 13]. Amplifications were carried out in a total volume
of 45 μL containing 1xPCR buffer (50 mM KCl, 20 mM Tris
pH 8.4, 2.0 mM MgCl2, 0.2 mM cresol red, 12% sucrose,
0.005% (wt/v) BSA (Boehringer-Mannheim, Mannheim,
Germany)), 2.5 pmol of each primer, 40 mM dNTPs (Pharma-
cacia LKB, Uppsala, Sweden) and 1.25 units of AmpliTaq
polymerase (Perkin Elmer Cetus Corporation, Emeryville,
Calif, USA). Parameters and conditions used for amplifica-
tion were 94°C for 30 sec., 60°C for 60 sec., and 72°C for
60 sec., as described, in [11, 12].

For DGGE 10 μL aliquots were loaded onto a denaturing
gradient gel containing 6% polyacrylamide and a gradient
of urea and formamide ranging from 20% to 80%. Gels were
run at 160 V for 4.5 h in 1x TAE buffer kept at a constant tem-
perature of 56°C. After electrophoresis, the gel was stained
with SYBR Green I (Molecular Probes, Oregon, USA) and
visualized using the FLA-3000 fluorescence detection system
(FUJI film, Science Imaging Scandinavia, Sweden).

2.3. Viability. Cell counting and viability testing were per-
formed by microscopy. Cells were stained with trypan blue
followed by counting of live and dead cells in a haemocy-
tometer.

2.4. Sterility Tests. Bulk and REP cultures were intermittently
sampled for microbiological testing of fungal and bacterial
contamination.

2.5. Peptides. We used the following HLA-A2 restricted
peptides: SUR1M2 (LMLGEPFLKL), HTERT P540 (ILAK-
FLHWL), Cyclin B1 204 (ILIDWLVIQV), MART-1 27–35
(AAGIGILTV ), and NY-ESO 1 157–165 (SLLMWITQC).

2.6. Cell Lines. Autologous tumor cell lines were established
from tumor fragments by outgrowth in 24 well or 6 well
plates (Nunc) in medium consisting of RPMI1640 (Invit-
rogen), penicillin, streptomycin, fungizone, 10% fetal calf
serum (Invitrogen), and SoluCortef (Pfizer).

Tumor cells were cryopreserved in 90% FCS and 10%
DMEM (Hospital Pharmacy, RegionH, Copenhagen, Den-
mark) and stored at −140°C.

2.7. Flow Cytometry. Phenotyping were conducted using a
FACS-Aria with Diva software (from BD) and fluorescence
conjugated monoclonal antibodies (mAb) against CD3
APC-Cy7, CD4 APC, CD8 PerCP, CD25 PE, CD27 PE,
CD45RA FITC, CD45RO PE, CD56 PE (all from BD),
CCR7 FITC (BD Pharmingen), CD16 FITC (Dako), CD28
FITC (Immunotech), CD62Ligand PE (BD Pharmingen),
and CD57 FITC (BD Pharmingen) along with corresponding
isotypes.

2.8. T-Cell Receptor (TCR) Clonotype Mapping by Denaturing
Gradient Gel Electrophoresis (DGGE). RNA was extracted
using the NucleoSpin RNA II (Macherey-Nagel, Germany).
cDNA synthesis and quantitation of cDNA in each sample
was carried out as previously described [11].

2.9. Elispot INFγ Measurement. Antitumor activity
was assessed with Elispot INFγ quantification as described
previously [14]. In brief, nitrocellulose bottomed 96 well
plates (Multiscreen MAIP N45; Millipore, Denmark) were
coated with INFγ capturing antibody (1-DIK; Mabtech,
Sweden) and further washed and blocked with RPMI 1640.
A maximum of 1 × 10^5 effector cells per well were either added
alone when stimulated by peptides, or in coculture with
target cells (1 × 10^4 cells per well) consisting of autologous
tumor cells. After a four-hour or overnight incubation pe-
riod, the medium was discarded and wells washed followed
by application of secondary biotinylated antibody (7-B6-1-
Biotin; Mabtech). The plates were incubated for one hour,
then washed, and avidin–enzyme (Streptavidin; Mabtech)
conjugate, were added to each well followed by one-hour in-
cubation at room temperature. Succeedingly, the wells were
washed and the enzyme substrate NBT/BCIP (nitro blue
tetrazolium/5-bromo-4-chloro-3-indolyl phosphate; Mab-
tech) were added into each well. The plates incubated at
room temperature for 2 to 10 minutes, while emerging pur-
ple spots developed. The reaction was terminated with tap
water. Spots were counted with the ImmuNoSpot Series 2.0
Analyzer (CTL Analyzers) and the frequency of tumor
specific TIL could be calculated from the numbers of spot
forming cells. The assays were preferably done in triplets or
in duplicates in case of low cell numbers.

2.10. Cr Release Assay. A standard Cr51-release assay was
used to quantify the specific cytotoxic ability of selected TIL
cultures. In brief, 5 × 10^3 Cr51-labeled tumor cells (duplicates
or triplicates) were cocultured with TIL (maximum E:T
ratio of 100:1 and titrated) in RPMI containing 10% FCS for a 4 hour incubation period. Thereafter, Cr.sup31-release was measured and percentage of tumor lysis calculated as (#count – Min count)/(Max count – Min count) × 100%.

2.11. Statistical Analysis. We utilized Graphpad Prism statistical software to analyse for statistical differences, using a paired two-tailed t test. P values < .05 were considered significant.

3. Results

3.1. Patients. Tumor material were obtained from 17 patients with either locally advanced or advanced disease from metastasis localized either in lymph nodes (majority of specimens) or subcutaneously. A minimum requirement of 1 cm³ of tumor was needed to ensure sufficient material for TIL expansion. The mean age was 62 years with an equal gender distribution. 12 of the patients had only been treated surgically prior to inclusion, while five patients who were included in our recent established clinical pilot trial had previously received IL-2 and/or DC vaccination based immunotherapies. Patients showed the following distribution of HLA-A types: one HLA-A1+, two HLA-A1/A3+, one HLA-A3+, two HLA-A3/A11+, one HLA-A11+, one HLA-A3+/A2+, four HLA-A2+, one HLA-A2/24+, and four non-HLA-A1/A2/A3/A11/A24.

3.2. TIL Expansion Kinetics. Lymphocytes migrated out from the fragments within two-to-five days and expanded into a confluent layer before splitting the wells. Each well initiated an individual bulk culture and were kept separated from other cultures. TIL bulk cultures expanded to at least 5 × 10⁶ cells were considered sufficiently expanding. This was obtained in 15 out of 17 patients (88%) in 6% to 100% of the bulk cultures (mean 58%) within 3–5 weeks. We found that growth rates varied markedly even between cultures from the same patient, and there was no difference in success rate of TIL growth from LN or SC tumor material, nor between the IL-2 concentrations (data not shown).

We next tested the proliferative potential of a range of bulk cultures from 12 of the 15 patients with sufficiently growing TIL. This rapid expansion procedure (REP) involves the addition of allogeneic feeder cells and a CD3 antibody and has shown to increase TIL expansion rates considerably, in previously reports in both melanoma and head and neck squamous cell carcinoma. Again, the kinetics could vary between cultures from the same patient; however, the procedure could efficiently expand TIL bulk cultures to over a 1000-fold in more than half of the cultures in 2 weeks (Figure 1).

3.3. Phenotypes and Clonal Composition. TIL were visualized in the microscope, showing a blasted morphology related to actively dividing lymphocytes laying either as single cells or in clusters/clones.

In acquisition of cells by flow cytometry, gating of viable cells was performed on the basis of the forward and side scatter dot plots. T-cells (CD3+) predominated the cultures, while NK cells (CD16/56+) (Figure 2) were consistently absent. In bulk cultures, we observed a heterogeneous CD4+ and CD8+ T-cell distribution among cultures inter- and intraindividually. There was, however, an overall skewing towards a CD8+ (mean = 74% ± 24%, range 30%–94%) T-cell predominance in relation to CD4+ T-cells (mean = 19.5% ± 23.5%, range 1%–64%). Next, we investigated the occurrence of surface markers identifying T-cell memory subsets, or alternatively, a differentiation path of effector cells, in the overall CD3+ population, and among CD4+ and CD8+ T-cell subsets in comparison to TIL after two weeks of REP (Figure 2). Overall, there was a distinct predominance of CD45RO+ and CCR-7lo/- T-cell populations before and after REP identifying the cells as T effector memory like. TIL were further characterized by surface markers according to a proposed model of effector CD8+ T-cell differentiation stages by Gattinoni et al. [15]. Expression of the lymphoid homing marker CD62L was significantly reduced after REP in the CD3+ population and the CD8+ subsets and remained unchanged among the CD4+ subsets. Concerning expression of costimulatory markers, we observed a significantly higher expression of CD27 among CD8+ bulk TIL compared to CD4+ cells, while CD4+ cells had sustained higher CD28 expression in bulk cultures and after REP. Although there was a relatively high percentage of CD27/28 double positive cells in a few bulk TIL, they were downregulated after REP. Finally, there was a significant increase in the high-affinity IL-2 receptor (CD25) after REP in the CD4+ population. In conclusion, the CD8+ population express surface markers (CD45RO+, CCR-7Lo, CD62LLo, CD27Lo, CD28Lo, and CD57Lo) resembling intermediate to late-stage effector cells as reported by other groups [8, 9, 16].

Selected expanded cultures were analyzed for the presence of clonally expanded T cells by RT-PCR/DGGE-based TCR clonotype mapping. Analysis revealed the presence of at least 10 different T-cell clonotypes in bulk cultures as well as in rapidly expanded cultures (data not shown). The results support our previous findings in expanded TIL from head and neck cancer patients, that expansion by high-dose IL-2 and CD3 antibody seems to support the continued expansion of bulk T-cell clones.
Figure 2: Phenotypes. FACS determination of phenotypes of TIL cultures pre- (open circles) and post-REP (closed circles) are represented from six patients. The overall T-cell effector memory like phenotype (CD3+CD45R0+CCR-7lo) is preserved after REP with a sustained low expression of CD57 and intermediate CD25 expression. CD28 remains unchanged, while CD62L and CD27 is downregulated, indicating a differentiation towards a later effector stage.

3.4. Sustained Functional Capacity during Expansion. TIL cultures from eight patients were selected to scrutinize the presence of specific T-cell populations in bulk cultures and after REP. Peptides derived from over expressed (Telomerase, Survivin, and Cyclin B1), differentiation (MART-1) and cancer testis antigens (NY-ESO-1) served as known targets, while autologous tumor cell lines presented a panel of unknown antigen specificities. Elispot detection of INFγ release upon antigenic stimulation revealed a sustained functionality of TIL after REP. Due to the high sensitivity of the assay, we
could follow the presence and loss of low-frequency single-peptide-specific T-cell populations (Figure 3(a)), occurring as a consequence of an increase or decrease in cell number of a given specific cell population, during the unspecific expansion procedures provided by IL-2 and anti-CD3. Autologous tumor cell lines were available in four patients, and all patients contained TIL showing antitumor activity in Elispot (Figure 3(b) and data not shown). The presence of autologous tumor-specific T-cell populations was more resistant during REP and showed a sustained functional capacity. Although T-cell-specific antitumor activity was predominating in TIL, we observed LAK/NK cell activity in a few cultures (Figure 3(b)) by unspecific engaging the cell lines K562 and Daudi. Finally, we confirmed a sustained tumoricidal capacity of TIL after REP (Figure 3(c)) indicating that TIL expanded to clinical relevant numbers (2400- and 4000-fold) can engage and kill autologous tumor.

4. Conclusion

We were able to establish sufficiently expanding TIL bulk cultures in five weeks from the majority of included melanoma patients. Further expansion by REP generated a mean expansion fold of 1400 in two weeks, ensuring the feasibility to reach clinical relevant quantities for clinical testing. Based on earlier studies of T-cell therapy of melanoma patients were as low as $1.3 \times 10^9$ infused cells containing 30% MART-1-specific CD8+ T-cells mediated a complete clinical response [17], we estimate that a minimum of $3 \times 10^9$ cells are required to obtain a therapeutic effect. Cell-based analysis revealed an oligoclonal composition of T effector memory cells, predominated by CD8+ cells showing an intermediate to late stage of differentiation after REP. TIL retained the functional capacity measured by INFγ release and lytic activity against autologous tumor. Notably, we did not find differences between the two doses of IL-2 used during TIL.
culturing, and even further lowering of IL-2 dose to 3000 IU/mL is now the standard used in TIL expansion at other centres. Finally, there were no significant influence on TIL expansion kinetics or phenotypes by pretreatment, age or performance status of the patients.

5. Perspectives

In a recently initiated clinical trial of TIL-based ACT, low-dose IL-2, and lymphodepletion preconditioning, one out of five treated melanoma patients has obtained an ongoing partial response (+13 months). We are currently screening the TIL cultures for the occurrence of tumor associated antigen (TAA) specificities by measuring INFγ in Elispot. This enables us to identify the specific combination of TAA specificities in each patient, which potentially can be identified during immune monitoring of the patient samples. In addition, we are establishing and validating a flow cytometry-based method of identifying TAA-specific T-cell populations and obtain information on the kinetics of T-cell memory and effector stages before and after treatment. Information providing more insight into the prognostic values of adoptively transferred TIL.

Conflict of Interests

The authors state no potential conflict of interests.

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References


