



REVIEW

Cancer gene and immunotherapy: recent developments

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Gene and immunotherapeutic approaches to treat human malignant tumors are reviewed. Special attention is given to the different strategies of cancer gene therapy and to recent aspects of cytokine-supported tumor immunotherapy or tumor-specific vaccination. The limitations of these therapy approaches are critically discussed especially with respect to immune escape mechanisms.

Keywords: tumour; gene therapy; immunotherapy; cytokines; escape; melanoma; review

Gene therapy of cancer

In recent years various gene therapeutic approaches to the treatment of human cancer have been developed and over 300 clinical protocols have already been approved.^{1,2} Three main categories of somatic cell gene therapy can be identified.² First, cells are removed from the body or obtained from tissue culture, transfected *ex vivo* with a vector and reinfused after gene engineering into the patient.³ Second, the vector is placed directly *in situ* into affected tissues, eg the lung or a tumour mass, of the patient.² Third, is an *in vivo* therapy, whereby the vector is injected directly into the blood stream. At present there are only clinical examples of the first two categories. In these studies, the genes were transferred into target cells either by lipofection, electroporation or naked DNA technique using classic eukaryotic expression vectors, eg based on SV40 or cytomegali virus (CMV), or viral vectors based on RNA (eg MuLV), or DNA viruses (eg of adeno or poxvirus families).²

Using these tools, gene cancer therapy tries to intervene at different molecular or physiological levels of the malignant tumour cell development. One approach attempts to infect cancer cells with intact tumour suppressor (eg p53) or anti-oncogenes, and to use, respectively, antisense oligonucleotides or intracellular recombinant single chain antibodies to inhibit activated oncogenes (eg *c-myc*, *erbB2*).⁴⁻⁷ Other strategies try to modify cellular drug resistance (eg by wild type MDR cDNA) or to introduce 'suicide' genes, eg herpes simplex virus thymidine kinase (HSV-tk), into the tumour cells.^{8,9} A further approach, especially used in leukaemia and lymphoma patients, introduces gene markers (eg β -galactosidase, neomycin resistance) into blood or bone marrow cells to identify efficacy of purging and origin of relapse following autologous stem cell transplantation.^{10,11} The most commonly used concept (64/144 clinical protocols worldwide), however, tries to augment the immune system's capacity to eliminate autologous cancer cells by different approaches.

This is attempted by cytokine-supported immunotherapy, enhancing the tumour immunogenicity or tumour-specific vaccination and by combinations of the different approaches (see reviews of recent clinical protocols^{1,2}).

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Cytokine-supported immunotherapies

The notion that cytokines such as interferons and interleukins can enhance the immunogenicity and induce reproducible responses also to spontaneous tumours led to increasing investigations and later clinical use.¹²⁻¹⁴ Clinical trials were started with systemic applications of interferons and interleukin-2 (IL-2).^{15,16} After administration of high doses of recombinant IL-2 to 283 consecutive patients (Surgery Branch at the National Cancer Institute: 1985-1992), complete responses were seen in patients with metastatic renal cell cancer (9%) or melanoma patients (7%). However, systemic IL-2 administration was associated with substantial toxicity (eg capillary leak syndrome).¹⁴ These side effects, including treatment-related mortality, could be diminished by cardiac screening of patients and the aggressive use of prophylactic antibiotics, but they remain a problem.¹⁴

The mechanism(s) by which cytokines support an anti-tumoural immune response are manifold.^{12,13} It was proposed that local rather than systemic cytokine effects mimic the natural cytokine physiology and are more likely to promote T cell responses against weak immunogenic tumour antigens, to activate non-specific killing (NK), lymphokine activated killer (LAK) cells, Mo/M Φ , or to enhance tumour antigen presentation. This hypothesis was supported by rejection of tumours after repeated local administration of exogenous cytokines in the vicinity of draining lymph nodes or directly into the tumour.¹⁷ In addition, numerous studies have shown that weakly immunogenic tumour cells could be altered to become targets for specific immune rejection by transfection with cytokine genes. The first experimental models and clinical applications carrying cytokine gene transfected malignant cells focused on IL-2.¹⁸⁻²⁷ In most experimental systems, the expression of IL-2 by weakly immunogenic tumour cells resulted in growth inhibition of the modified tumour cells. In several cases, a systemic immune response against the parental tumour, leading to an immune memory against a challenge with parental tumour cells was also observed.¹⁹⁻²¹ The inhibitory effect was dose dependent and the degree of suppression of growth correlated directly with the amount of IL-2 produced by the tumour cells. In vaccination experiments, however, intermediate doses of IL-2 achieved highest cure rates in mice.^{28,29} Clinical trials treating advanced cancer patients with IL-2 secreting

tumour cells showed a variable outcome of the therapy. Thus, tumour regression, stabilisation of the disease, mixed types of response with simultaneous evidence of regressing and non-responding lesions in the same patient or progressive tumour growth were noted.³⁰⁻³³ The evaluation of mechanisms involved in treatment with transfected tumour cells showed that IL-2 release *in situ* leads to a dense infiltration of the tumour involving mainly CD4⁺ and CD8⁺ T cells as well as NK cells. It was demonstrated that CD8⁺ T cells are essential for complete tumour rejection and long term protection against the original tumour^{19,34,35} and that IL-2 can bypass CD4⁺ T helper function in the generation of an anti-tumour response¹⁹ or can overcome cytotoxic T lymphocyte (CTL) anergy.^{36,37} The *in vivo* role of NK cells remains unclear but might be important especially in tumours with down regulated MHC class I molecules or in the absence of killer cell inhibitory receptors (KIR).^{38,39,40,41}

Several other interleukins (IL-4, IL-7, IL-12, IL-13), interferons (IFN γ , IFN α), haematopoietic growth factors, granulocyte-macrophage colony-stimulating factors (GM-CSF, G-CSF, M-CSF), chemokines (MCP-1, TCA-3), and inflammatory mediators – tumour necrosis factor (TNF), IL-1, IL-6 – transfected in tumour cells showed similar tumouricidal effects mediated by different effector mechanisms.^{12,13} It was also demonstrated that cytokines can be protective against tumour growth if transfected into normal fibroblasts or when secreted by allo or xenogeneic cells with respect to the tumour and the host animal.^{13,38,42} The concept of using cytokine-transfected allogenic and xenogenic tumour cells or fibroblasts instead of autologous/syngenic cells in the prevention and treatment of tumours was developed by several groups, eg by P Kourilsky at the Institut Pasteur in Paris.¹² It included the transfection of different murine tumour cells with murine or human cytokines (IL-2, GM-CSF, IL-4, and IFN γ) or allogenic major histology complex (MHC) class I molecules (H2-Kb).^{12,38,43} Using allo or xenogenic cytokine-secreting cells, it was demonstrated that this specific immunotherapeutical strategy was effective to control tumour growth in several mouse and rat models.^{12,38,43} This anti-tumour effect was mainly mediated by NK-cells.³⁸ The results led to the initiation of a controlled study more closely mimicking the biology of human tumours. Treatment of spontaneous fibrosarcomas in cats and melanomas in dogs with xenogenic IL-2 transfected fibroblasts, the green monkey cell line

(Vero-IL-2) (Transgène, Strasbourg) resulted in substantially improved disease free and overall survival.⁴²

Although many questions remain to be answered, and only in some experimental systems was rejection of established disease achieved,^{42,44-49} the promising results obtained in animal studies have established the basic biological value of gene therapy with cytokine-secreting cells and these approaches are already widely used in clinical trials.^{1,13} More than 60% (26/43) of recent 'cytokine-supported' clinical cancer gene therapy trials use IL-2, whereas others utilise IL-4, IL-7, IL-12, IFN γ , GM-CSF, TNF or combinations thereof.¹ One of the first cancer gene therapy trials initiated in Switzerland in January 1996 used the 'xenogenic approach'.^{12,42} In this study, three cohorts of three successive patients with advanced solid tumours accessible to CT or ultrasound-guided injection were treated intra-tumourally repeatedly with different doses of xenogeneic monkey fibroblasts secreting high doses of human IL-2 (Vero-IL-2). Endpoints of the study were feasibility, toxicity, and clinical and biological effects of this novel approach to immunotherapy of cancer. Treatment was well tolerated and toxicity consisted of transient fever in one patient and short-lived, mild itching and erythema in two others. One patient with soft tissue sarcoma showed a more than 90% and more than 50% reduction of the volume of two distant, non-injected metastases, lasting for 29+ and 26 months, respectively. Four other patients showed stabilisation of their disease for 3-9 months, among whom was a patient with melanoma who developed marked vitiligo. Repeated injection of up to 5×10^7 Vero-IL-2 cells was thus feasible and safe in heavily pretreated patients with advanced solid tumours and showed signs of clinical and biological activity.³³ Histopathological, immunological and molecular analyses were performed on biopsy specimens of tumours and blood samples obtained before, during and after treatment. Only a slight but statistically significant ($P < 0.05$) increase of serum IL-2 could be observed on day 5 of treatment which returned to base line values by day 15. Also on day 5, a significant increase ($P < 0.05$) in CD3 mRNA was detected in tumour biopsies by RT-PCR, indicating transient infiltration of injected sites by T lymphocytes.³³ No expression of exogenous IL-2 mRNA (ie of the transgene) could be detected by RT-PCR in tumour biopsies and peripheral blood of days 1, 5, and 15, most likely reflecting rapid destruction of Vero cells by patient's NK cell-like killer cells.

Melanoma antigens: recognition and vaccination

An important prerequisite for the induction of a tumour-specific immune response is the MHC restricted T lymphocyte recognition⁵⁰ of small antigenic peptides^{50,51} from tumour-specific (TSA) or tumour-associated (TAA) antigens in tumour cells.^{52,53,54}

To identify the antigenic peptides recognised by tumour-specific CTL on MHC class I molecules, mainly four methods have been used: transfection of recombinant tumour DNA libraries into cells expressing MHC presenting molecules,⁵⁴ biochemical purification of peptides eluted from MHC molecules of the tumour cells,⁵⁵ designing of consensus anchor motifs carrying candidate peptides from proteins known to be overexpressed or mutated in tumour cells⁵⁶ or screening of cellular immune response to SEREX-defined (serological analysis of tumour antigens by recombinant cDNA expression cloning) tumour antigens.^{57,58}

Based on the pattern of expression of the parent protein and tumour specificity, the antigens can be classified into five major groups:

- i) antigens resulting from mutations,^{59,60}
- ii) viral antigens,⁶¹
- iii) tumour-specific shared antigens of the 'cancer testis' gene family,^{62,63}
- iv) over expressed antigens^{64,65}, and
- v) tissue-specific differentiation (TSDA) antigens.^{53-55,66}

The number of tumour-specific mutated neoepitopes (eg in melanoma, renal cell cancer) or viral antigenic peptides (eg in cervical cancer, hepatomas) recognised by human CTL clones is growing continuously.^{60,61,67,68} In various cancer patients bearing the appropriate MHC haplotypes, however, an important number of CTL clones is directed against peptides from molecules (MAGE, BAGE, GAGE and RAGE gene family) of the so-called 'cancer testis' antigen group.^{53,54,66} These peptides are able not only to stimulate CTL specific to melanoma cells *in vitro*⁶⁹⁻⁷¹ but also to induce tumour regression *in vivo*.⁷²⁻⁷⁴ Another group of antigenic peptides recognised by a large number of CTL clones from melanoma patients⁷⁴⁻⁷⁸ derives from TSDAs (Pmel-17/gp100, MART-1/Melan-A, tyrosinase, and tyrosinase-related proteins, TRP-1 and 2) that are also expressed in normal melanocytes.⁷⁹ The role of CTL against these 'autoantigenes' in melanoma rejection is not clear. However, sometimes a depigmentation, a

so-called vitiligo, can be observed in treated melanoma patients. The observed depigmentation which is probably caused by destruction of normal melanocytes and the reported association of vitiligo with prolonged survival and spontaneous tumour regression support a possible role of anti-TSDA CTL in tumour regression.^{53,54} Most of the melanoma antigenic peptides characterised until now are restricted to HLA-A1, HLA-A2 or HLA-A31 molecules,^{53,54,58} expressed in about 26%, 44% or 6% of Caucasian melanoma patients, respectively.^{78,80} Therefore, clinical applications of antigenic peptides for tumour vaccination or therapy monitoring (eg by ELISPOT) are limited to patients of these MHC haplotypes.^{69,73,74,76,81,82} The first attempts to immunise melanoma patients with MAGE-1, MAGE-3, tyrosinase, gp-100 or MART-1 peptides showed objective tumour regression in a few patients. In some cases the clinical response was accompanied by the presence of peptide-specific CTL in the blood.^{69,76,81,83,84} A more 'MHC-independent', melanoma-specific vaccination was obtained in a more recent study,⁷³ using autologous dendritic cells (DC) pulsed with tumour cell extracts, supported by an additional 'non-specific' CD4 helper cell stimulation with keyhole limpet haemocyanin (KLH) which resulted in objective clinical responses in 5/16 patients. The tumour regression was accompanied by a peripheral immune response, as detected by delayed-type hypersensitivity skin reaction to melanoma peptide or tumour cell lysate pulsed DCs.⁷³ The application of somatic cell hybrids (SCH)⁸⁵⁻⁸⁷ of syngenic or autologous tumour cells with professional APCs (dendritic cells, M Φ , or activated B-cells)⁸⁸ has been recently applied in rat and mouse tumour models as another successful therapeutic approach for a more 'MHC-independent' vaccination.⁸⁵⁻⁸⁶

A third way, the combination of 'non-specific' (eg IL-2 supported) stimulation and specific vaccination (gp-100 peptide) also resulted in objective cancer responses but was accompanied by decreased circulating CTL in the blood of 13/31 patients.⁷⁴ This disappearance of peripheral CTL was explained by their possible invasion of tumour sites. In contrast to the other vaccination strategies, a clinical response to this treatment, however, was only found in those patients who had also received systemic IL-2, indicating synergistic effects of the two components. Together with the results of our phase I study, these data support our recently initiated randomised, French-Swiss multi-centre phase II trial using two different doses of Vero-

IL-2 cells injected repeatedly in patients with melanoma.

A simultaneous monitoring of the treatment seems to be important for understanding the mechanisms of different treatment strategies. Skin testing,⁷³ somatic cell hybrids^{85,86} or the use of autologous antigen presenting cells (APC) pulsed with peptides, tumour extracts or infected with recombinant viruses encoding tumour antigens^{89,90} could become useful tools for monitoring peripheral anti-tumour responses to detect CTL precursor cells recognizing epitopes restricted to various MHC haplotypes *in vitro*. Additionally, these approaches also allow the detection of MHC class II restricted tumour antigen presentation.^{84,89,90}

Immune escape mechanisms

Induction and detection of T-cell response to tumour antigens, however, is frequently limited by the tumour cells themselves. Malignant transformation is often associated with genetic alterations providing tumour cells with mechanisms to escape from immune surveillance leading to selection of less immunogenic tumour cell variants.⁹¹⁻⁹⁵ This can be observed in primary or metastatic tumour lesions *in vivo* but can also be detected after oncogenic transformation *in vitro*. At the molecular level, the defective signalling of tumour cells could be attributed to

- i) down regulation or loss of major histocompatibility complex (MHC) molecules,^{96,97}
- ii) alteration of antigen processing pathways, resulting in an inability to present tumour-specific antigens to T cells,^{98,99}
- iii) absence of costimulatory or adhesion molecules that are essential for activation of the host immune system,^{100,101} or
- iv) production of factors modifying host immune responses.¹⁰²

Presentation of antigenic peptides on MHC class I or II molecules is essential for induction or reactivation of a specific immune response.^{50,103-106} Downregulation or loss of certain MHC haplotypes or melanoma antigens enables tumour cells to escape immune surveillance.¹⁰⁷⁻¹⁰⁹ Therefore, an altered expression of these molecules might represent an important parameter restricting the chances of a successful gene/immune therapy.^{107,110,111} However, the opposite might also be true. A partial human leukocyte antigen (HLA) loss

can lead to tumour cell killing by KIR expressing CTL if the loss-variant downregulates respective MHC molecules (eg HLA-Cw7) recognised by the KIR (eg p58.2), whereas original tumour cells will not be lysed.³⁹ The analysis of the involved mechanisms, eg β 2M loss, 'defective' expression of transporter associated with antigen presentation (TAP) or proteasome subunits,^{94,109,110,112} will also show whether reversible (eg by IFN γ stimulation) or irreversible changes in the tumour cells are responsible for defective tumour antigen expression.^{94,109,110,112}

Outlook

Immune therapy is one possibility of cancer treatment. Since the detection of tumour-specific or associated antigens in spontaneous human tumours, rapidly progressing strategies to treat cancer patients have been developed. Successful treatment of patients, however, is furthermore narrowed by our limited understanding of relationships between the components of the immune system and tumour cells. In the case of gene therapy, in 1996, experts of the US Department of Health, NIH, evaluated results of the first 7 years of treatment. In conclusion, they noted that gene therapy has been a feasible and until now also very safe treatment, but unequivocal proof of clinical efficacy failed in many of the more than 100 clinical trials. The most important consequence of the report was the request to 'go back to the bench' as well as to develop more effective viral and non-viral gene therapy vectors. In the meantime more than 300 clinical protocols with more than 3000 patients are open worldwide but the assessment of the situation has not changed dramatically. The main problems are still the failure of vectors to transduce efficiently their targets *in vivo*, the lack of tumour specificity of available application systems, the increasing switch-off of transcription elements *in vivo*, even after successful transfer of the transgenes, and the incomplete understanding of the molecular pathology of tumour development and progression.

W French Anderson, one of the pioneers of gene therapy expects the first unequivocal clinical results within the next 5 years, most probably in the field of genetic originative diseases.² In the same period the first tumour-specific vectors will be applied in clinical practice and tissue-specific expression of transgenes will be realised. Among others, one result of the terminated 'Human Genome Project' will be an exponential growth of various gene therapy studies in the subsequent decade. At the end of this period, the first

intravenous applicable vectors will be used in clinical practice, resulting in an efficient expression of transgenes in a sufficient number of tumour cells. When induction of tumour remission by gene or immune therapy will be possible remains speculation. Probably gene and/or immune therapy at most will act as part of a multimodal strategy in oncological therapy, combining surgical, radiotherapeutical, cytostatical and other strategies.

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