



**Karolinska  
Institutet**

**Institutionen för Onkologi-Patologi**

## **Vaccination in gastrointestinal cancer**

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*To my family*



## Abstract

Advances in immunology have increased the possibility to develop therapeutic cancer vaccines (TCV), as a complementary approach to standard treatment. The goal of a successful cancer vaccine is to induce a potent long-lasting immune response against the tumour with limited toxicity on normal cells. Most tumour cells express tumour-associated antigens (TAA), which can act as targets for the immune system. However, most TAAs evade recognition by the immune system to avoid auto-immunity, as many TAAs coexist in normal tissues. Commonly expressed TAAs in gastrointestinal malignancies are Carcinoembryonic antigen (CEA) and telomerase which both have been used as targets in cancer immunotherapy.

The aim of this thesis was to explore the immunogenicity and safety of a CEA based protein and DNA TCV in patients with colorectal cancer (CRC) in the adjuvant setting and telomerase vaccination (GV1001) in patients with advanced pancreatic adenocarcinoma (PC).

A long-term follow-up of CRC patients immunized with recombinant (rCEA) ± Granulocyte-macrophage colony-stimulating factor (GM-CSF) was conducted. Induction of anti-CEA IgM, IgA and IgE antibodies was monitored from 36 months after start of immunization. GM-CSF significantly augmented the anti-CEA response for all three classes ( $p < 0.05$ ). A significant correlation between survival and high IgA anti-CEA titers was noted ( $p = 0.02$ ). Anti-CEA IgA antibodies could lyse CEA positive cells in antibody-dependent cellular cytotoxicity (ADCC) and complement-dependent cytotoxicity (CDC) assays.

The type, severity and duration of side-effects of CEA66-DNA vaccination in combination with cyclophosphamide and GM-CSF, was evaluated in 10 CRC patients. CEA66-DNA was delivered by a needle-free device system (Biojector). Adverse events (AE) were mild and transient, without any grade 3 or 4 AEs. No clinical signs of autoimmunity were seen.

In an explorative study using CEA66-DNA (producing unglycosylated CEA) and wild type (tetwt)-CEADNA (producing glycosylated CEA) for immunization in combination with cyclophosphamide and GM-CSF immune responses (proliferation assay, ELISPOT, cytokine secretion assay) were analyzed in the adjuvant setting of CRC patients. 10 patients received intradermal (i.d.) or intramuscular (i.m.) CEA66-DNA by Biojector at weeks 0, 2 and 6 (part 1). 10 patients; (part 2), received tetwt-CEADNA 400 µg i.d. by needle followed by electroporation at weeks 0 and 12. Part 3 (n=6) included patients primed with CEA66-DNA and boosted with tetwt-CEADNA. GM-CSF and cyclophosphamide was also included. In total, 16 out of 20 (80%) patients mounted a single assay cellular response; 10/10 (100%) in part 1 and in 6/10 (60%) of the patients in part 2 ( $p = 0.025$ ). Immune responses were weak but durable.

We also assessed the safety and immunogenicity in advanced PC patients using a 16 aa telomerase peptide (GV1001) for vaccination in combination with GM-CSF and gemcitabine as first line treatment. Three different vaccine treatment schedules (groups A, B, C) were used. In groups A and B, differing only in the dose of GM-CSF, a total of 67% of the patients showed an induced telomerase response. An induced ras (antigenic spreading) specific immune response was noted. All responses were weak and transient. A significant decrease in regulatory T cells over time was noted in patients in groups A and B.

In conclusion, durable weak anti-CEA immune responses were seen following rCEA and CEA-DNA vaccination in CRC patients in the adjuvant setting. Weak and transient anti-telomerase responses following peptide vaccination were induced in patients with advanced PC. To develop a therapeutic concept of clinical significance measures have to be taken to optimize vaccine strategies.

Keywords: CEA, hTERT, immunotherapy, cancer vaccine

## List of publications

- I. **Staff C**, Magnusson CGM, Hojjat-Farsangi M, Mosolits S, Liljefors M, Frödin JE, Wahrén B, Mellstedt H, Ullenhag G J. Induction of IgM, IgA and IgE antibodies in colorectal cancer patients vaccinated with a recombinant CEA protein. *J Clin Immunol.* 2012, 32:855-865.
- II. **Staff C**, Mozaffari F, Haller K, Wahren B, Liljefors M. A phase I safety study of DNA immunization targeting carcinoembryonic antigen in colorectal cancer patients. *Vaccine.* 2011, 29:6817-6822.
- III. **Staff C**, Mozaffari F, Haller KB, Frödin JE, Wahren B, Mellstedt H, Liljefors M. DNA immunization targeting carcinoembryonic antigen in colorectal cancer patients. (*Manuscript to be published*)
- IV. **Staff C**, Mozaffari F, Frödin JE, Mellstedt H, Liljefors M. Telomerase-peptide vaccination (GV1001) together with gemcitabine in advanced pancreatic cancer patients. (*Manuscript to be submitted*)



## List of abbreviations

|                   |  |
|-------------------|--|
| 5-FU              | 5-Fluorouracil                                   |
| ADCC              | Antibody-dependent cellular cytotoxicity         |
| AE                | Adverse events                                   |
| Ag                | Antigen  |
| AJCC              | American Joint Committee on Cancer               |
| APC               | Antigen presenting cell                          |
| Bev               | Bevacizumab                                      |
| CDC               | Complement-dependent cytotoxicity                |
| CEA               | Carcinoembryonic antigen                         |
| Cet               | Cetuximab  |
| CFC               | Cytokine flow-cytometry                          |
| CR                | Complete remission                               |
| CRC               | Colorectal carcinoma                             |
| CT                | Computed tomography                              |
| CTL               | Cytotoxic T lymphocyte                           |
| CTLA-4            | Cytotoxic lymphocyte antigen 4                   |
| Da                | Dalton   |
| DC                | Dendritic cell                                   |
| DTH               | Delayed type hypersensitivity                    |
| ELISA             | Enzyme linked immunosorbent assay                |
| ELISPOT           | Enzyme-linked immunosorbent spot                 |
| EGFR              | Epidermal growth factor receptor                 |
| EP                | Electroporation                                  |
| EpCAM             | Epithelial cell-adhesion molecule                |
| ER                | Endoplasmatic reticulum                          |
| F(ab)             | Fragment antigen binding                         |
| FasL              | Fas ligand                                       |
| GM-CSF            | Granulocyte-macrophage colony-stimulating factor |
| HD                | Healthy donor                                    |
| HIV               | Human immunodeficiency virus                     |
| HLA               | Human leukocyte antigen                          |
| hTERT             | Human telomerase reverse transcriptase           |
| i.d.              | Intradermal                                      |
| IFN               | Interferon                                       |
| Ig                | Immunoglobulin                                   |
| i.m.              | Intramuscular                                    |
| IL                | Interleukin                                      |
| KRAS              | Kirsten rat sarcoma viral oncogene               |
| LPS               | Lipopolysaccharid                                |
| mAb               | Monoclonal antibody                              |
| mCRC              | metastatic colorectal cancer                     |
| MDSC              | Myelo-derived suppressor cells                   |
| MHC               | Major histocompatibility antigen                 |
| MRI               | Magnetic resonance imaging                       |
| NF- $\kappa\beta$ | Nuclear factor $\kappa\beta$                     |
| NK                | Natural killer cell                              |
| NKT               | Natural killer T cells                           |
| NO                | Nitric oxide                                     |

|              |   |
|--------------|---|
| OIRR         | Overall induced immune response         |
| OS           | Overall survival                        |
| Pan          | Panitumumab                             |
| PBMC         | Peripheral blood mononuclear cell       |
| PFS          | Progression free survival               |
| PD1          | Programmed death-1                      |
| PDL1         | Programmed death ligand-1               |
| PHA          | Phytohemagglutinin                      |
| PPD          | Purified protein derivate of tuberculin |
| PSA          | Prostate specific antigen               |
| rCEA         | Recombinant carcinoembryonic antigen    |
| RNA          | Ribonucleic acid                        |
| SAE          | Serious adverse events                  |
| s.c.         | Subcutaneous                            |
| SD           | Stable disease                          |
| SFU          | Spotforming units                       |
| TAA          | Tumor associated antigen                |
| TAM          | Tumor infiltrating macrophage           |
| TCR          | T cell receptor                         |
| TCV          | Therapeutic cancer vaccine              |
| tetwt        | Tetanus wild type                       |
| Th           | T helper                                |
| TGF- $\beta$ | Transforming growth factor- $\beta$     |
| TIL          | Tumor infiltrating lymphocytes          |
| TLR          | Toll-like receptor                      |
| TME          | Total mesorectal excision               |
| TNF          | Tumor necrosis factor                   |
| TT           | Tetanus toxoid                          |
| TTP          | Time to progression                     |
| VEGF         | Vascular endothelial growth factor      |

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**PAPERS I-IV**

# 1 THE IMMUNE SYSTEM

*“Den som är väldigt stark måste också vara väldigt snäll” Ur Känner du Pippi Långstrump*

The immune system is crucial to human survival and can be defined as our defence towards pathogens. The word immune stems from the Greek word *immunitas* meaning “liberated or excluded from” for example the effects of a specific pathogen. In brief, two groups of the immune system coexist and interact constantly; the innate (non-adaptive) and the acquired immune system. The innate is quick to respond to pathogens, the acquired is slow but on the other hand it gives rise to a specific memory towards the pathogen – rendering a quicker and more precise immune response at the next encounter. *Thus, the innate immune system carries the memory of the species whereas the acquired immune system carries the memory of the individual.*

The *innate responses* use phagocytic cells (neutrophils, monocytes and macrophages), cells able to release inflammatory mediators (basophils, mast cells, and eosinophils), natural killer (NK) cells and natural killer T cells (NKT cells). Besides NK cells, interdigitating dendritic cells (DCs) comprise essential links between innate and acquired immunity <sup>[1]</sup> The molecular compartments of innate responses include complement, acute-phase proteins and cytokines. Cytokines are proteins made by cells that affect the behaviour of other cells and bind to specific receptor on their target cells. Cytokines made by lymphocytes are often called interleukins (IL).

*Acquired responses* involve the proliferation of antigen-specific B and T cells, which occurs when the surface receptors of these cells bind to the cognate antigen. Specialized antigen-presenting cells (APCs) display the antigen to the major histocompatibility complex (MHC) molecules and cooperate with them in response to the antigen. B cells respond to the immunogens by secreting immunoglobins (Ig), the antigen-specific antibodies mainly responsible for elimination of extracellular agents. There are two main types of T cells: helper T cells (Th) (CD4+) and cytotoxic T cells (CD8+). Although T cells can help the B counterpart to make antibodies, they also contribute to the eradication of intracellular pathogens by activating macrophages and directly killing infected cells.

## 1.1 Antigen presentation to T cells

The immune system recognizes antigens presented by two types of MHCs, Class I and II. In humans, the MHC Class I molecule corresponds to the human leukocyte antigen (HLA)-A,- B, and C- molecules, and MHC Class II molecules correspond to HLA-D molecules.

MHC Class I molecules are expressed on all nucleated cells and in general present 8-11 amino acid-long peptides derived from *intracellular proteins* digested in the proteasome complex. These MHC class I peptide complexes are recognized by CD8+ T cells. If a CD8+ cell finds a MHC Class I peptide -complex, with its unique T cell receptor (TCR), it undergoes clonal expansion and differentiates into mature CD8+ and memory CD8+ cells. Mature CD8+ cells migrate through the body, searching for cells that possess peptide-

complexes, to which the TCR will bind and destroy. Memory CD8+ cells persist and will multiply if they are re-exposed to the same MHC peptide complex again.

MHC Class II molecules are mainly expressed on APCs, such as dendritic cells (DCs), macrophages and activated B cells. These cells engulf proteins and present small peptides, in conjunction with MHC Class II to CD4+ T cells. These peptides are longer, usually 13 to 17 amino acids (or more), and derive from *exogenous proteins* endocytosed into the cell. If a CD4+ cell recognizes such a MHC class II-peptide complex, it will divide and mature producing many cytokines and express many surface elements, thereby regulating other aspects of the immune system.

## 1.2 Activation of naïve T cells

All types of acquired immune responses are started by the activation of antigen-specific T cells and the whole process of activation and differentiation occurs in the immediate nursing environment of APCs. Two signals are necessary for the initial activation of naïve T cells. The “*two signal model*” predicts that when an antigen is presented by an MHC molecule (signal 1) together with costimulatory molecules (signal 2) expressed by APCs, an immune response will be generated [2, 3]. Several costimulatory molecules have been described; one important complex is CD28 on T cells binding to B7’s on the APC. According to the danger theory, signal 1 in the absence of signal 2 leads to tolerance irrespective if the antigen is self or non-self. In the absence of danger signals, the APCs are not induced to express costimulatory molecules and the result of such presentation will be tolerance rather than activation. Any cell, however, when stressed and in danger, sends alarm signals to activate APCs.

The not-ever present expression of B7’s on APCs ensures that naïve T cells do not respond to their specific antigens in the absence of danger. This provides a mechanism for peripheral tolerance that prevents naïve T cells with receptor that bind self-antigens from being activated and differentiate into auto reactive effector T cells.

## 1.3 CD4+ Th1 and Th2 cells

Towards the end of their proliferation, activated T cells, acquire the capacity to synthesize the proteins they need to perform the specialized functions as effector T cells.

For CD4+ cells these proteins comprise cell-surface molecules and soluble cytokines that activate and help other types of cells to participate in the immune response. Because of its facilitating functions, CD4+ cells are called T helper (Th) cells. Defining the range of behaviour are two types of helper cells called CD4+ Th1 or CD4+ Th2. The main cytokines secreted by Th1 cells, IL-2 and INF- $\gamma$ , promote cytotoxic T cell lymphocytes (CTL) generation and activate NK cells. They may also secrete Granulocyte-macrophage colony-stimulating factor (GM-CSF), which promote a more efficient antigen presentation. The main cytokines secreted by Th2 cells, IL-4, IL-5, IL-6 and IL-10, mainly lead to B cell differentiation. Thus an immune response biased towards Th1 cells is described as being *cell-mediated immunity*, dominated by CD8+ T effector cells whereas an immune response biased towards Th2 cells is dominated by antibodies, described as a component of the *humoral immunity*.

## 1.4 CD8+ T cells

CD8+ T cells (CTL) are the principle effector cells of the acquired immune response, which mediate antigen-specific, MHC-restricted, cytotoxic effects. Antigen-specific CD8+ T cells become activated by the TCR-MHC Class I peptide interaction on an APC, together with help from activated T cells. The activation of CTL occurs either via direct recognition of antigen on the tumor cell, or by presentation of tumor antigens on APC's (cross presentation) (see Section 1.6), which subsequently prime CTL. This leads to clonal expansion of antigen-specific CTL that will lyse cells that express the same peptide-MHC Class I complex. CTL mediated cytotoxicity is mediated primarily by cytotoxins (perforin, granzymes) or via interactions between cell surface molecules on the CTL (Fas ligand) and the target cell (Fas molecules) <sup>[4]</sup>.

## 1.5 B cells

Each B cell has a receptor molecule with a single specificity called B cell receptor (BCR) or immunoglobulin in its secreted form. Five major Ig classes exist; IgM, IgD, IgG, IgA and IgE and these determine the functional activity of the antibody. After developing B cells leave the bone marrow, they begin to recirculate between the blood, secondary lymphoid tissue and the lymphatic vessels. At this stage the B cells are not fully mature, expressing surface IgM and IgD. Naïve B cells (that have not yet met their specific antigen) passing through the secondary lymphoid tissue pick up, (via surface immunoglobulins), process and present their specific antigens if present there. Upon meeting Th2 cells at the same site, the T cell receptor screen the peptides presented by the MHC Class II molecule on the B cell. Th2 cells then provide signals that activate the B cell to proliferate and differentiate. Some differentiate into plasma cells (the effector cell), which secrete IgM in the secreted form rather than the surface-bound form. Other activated B cells differentiates and undergo isotype switching and hypermutation, producing other classes of antibodies. Cytokines secreted by Th2 cells – IL-4, IL-5 and transforming growth factor- $\beta$  (TGF- $\beta$ ) influence B cells to switch their isotype. Some cells will develop into resting memory B cells, capable of a quicker and stronger immune response at the next encounter with the antigen. In humans IgG can be further subdivided into four subclasses of human IgG (IgG1, IgG2, IgG3 and IgG4), whereas IgA antibodies exist as two subclasses, (IgA1 and IgA2). IgG1 and IgG3 are the most potent mediators of antibody dependent cellular cytotoxicity (ADCC) and complement-dependent cytotoxicity (CDC). Circulating antibodies can bind to antigens on microorganisms or transformed cells and destroy them either directly or by ADCC and CDC. Some antigens, notably components of bacterial cell walls and capsules, are capable of inducing a rapid antibody response that does not require T cell help.

## 1.6 Cross-presentation and cross-priming

As a general rule, exogenous antigens enter the MHC Class II processing pathway and are presented to CD4+ T cells. By a process known as cross-presentation exogenous antigens are delivered into the MHC Class I processing pathway and presented to CD8+ T cells <sup>[5]</sup>.

Cross-presentation is involved in responses to viral infections, transplanted organs and in tumors<sup>[5, 6]</sup>. TAAs, released by tumor cell or administered as a therapeutic cancer vaccine (TCV), are taken up by APCs, which process and present them on APC cell surface restricted by their own MHC Class I or II molecules.

APCs such as DCs can then efficiently prime naïve T cells if they display MHC-antigen complexes (signal 1) together with costimulatory molecules (signal 2). When an immune response is initiated by cross-presentation this is known as a cross-priming of the immune response<sup>[6]</sup>.



## 2 THE INTERACTION BETWEEN THE IMMUNE SYSTEM AND CANCER

### 2.1 Pioneers of modern oncoimmunology

Immunotherapy for cancer was initially used over a century ago. In the 1800s German physician Paul Ehrlich and the American physician William Bradley Coley proposed that vaccination might be used against cancer. Ehrlich delivered the idea of “the magic bullet” that would specifically kill malignant cells but failed to demonstrate that weakened cancer cells, injected in patients, may generate antitumor immunity<sup>[7]</sup>. On the other hand, inspired by cancer patients who underwent regression following streptococcal fevers, Coley became convinced he could use bacteria to kill cancer cells. Coley created a mixture of heat-killed bacteria (Coley toxin) in 1896 that mediated potent antitumor effects in patients<sup>[8,9]</sup>. One of the major hurdles against development of anticancer vaccines, the “self/non-self” dichotomy, was later theorized by Frank Macfarlane Burnet in 1949 and in fact hampered further development for quite some time<sup>[10]</sup>. According to this theory, tumors – constituting self-tissues – are non-immunogenic and therefore insensitive to immunotherapeutic interventions. In 1994, Polly Matzinger however presented a theory that proposed that the immune system do not react to non-self (while sparing self) constituents, but would rather respond to situations of “danger”, either of exogenous (non-self) or endogenous (self) origin i.e. irrespective of the origin<sup>[11]</sup>. Hence, conditions that have long been thought to be immunologically silent, including cancer and trauma, are now thought to be able to activate the immune system<sup>[12-14]</sup>.

### 2.2 Tumor immunosurveillance and immunoediting

The long-standing theory of “*immunosurveillance*” suggests that cells are constantly monitored by an ever-alert immune system and that this surveillance is responsible for recognizing and eliminating incipient cancer cells. Accordingly, solid tumors have somehow managed to avoid detection by the immune system or have been able to limit the extent of immunological killing, thereby evading eradication<sup>[15]</sup>.

The developments of carcinogen-induced tumors were assessed in mice deficient for various components of the immune system. It was observed that tumors arose more frequently and grew more rapidly in immunodeficient mice compared to immunocompetent controls. Tumor incidence was, in particular, increased in mice with deficiencies in the development or function of CD8<sup>+</sup>, CD4<sup>+</sup> Th1 helper cells, or natural killer (NK) cells. Results from this experimental mode demonstrated that both the cellular arms of the innate and the acquired immune system contributed significantly to immune surveillance and tumor eradication<sup>[16,17]</sup>.

Spontaneous antitumor immunity that eliminates tumors or delays their growth involves production of INF- $\gamma$ , as well as the generation of CTLs<sup>[18,19]</sup>. Mice deficient in interferon (IFN- $\gamma$ ) are more susceptible to induced cancers, as are mice lacking T cells and perforin, a key effector protein in T cell cytotoxicity<sup>[18,19]</sup>.

A number of observations from humans indicate that functional immunosurveillance do exist. Patients with ovarian and colon cancer that are heavily infiltrated by CTLs and NK have a better prognosis than those who lack such tumor infiltrating lymphocytes (TILs) [20-22]. The hypothesis is further strengthened by data that indicate that patients indeed develop immune responses to tumor associated antigens [23, 24] (see Section 3.3.1) and also by the striking evidence of certain cancers in immunocompromised individuals [25].

According to the currently accepted model of immunoediting, neoplasms acquire the ability to develop and grow in spite of the immune system in three sequential steps. Initially, the growth of cancer is efficiently controlled due to robust tumor-specific immune responses (elimination). As the elimination phase is normally unable to completely eradicate malignant cells, some of them may acquire alterations that either reduce their immunogenicity or increase their resistance to the cytotoxic functions to the immune system (equilibrium). Such cells eventually grow uncontrolled (escape), leading to a clinically manifest neoplasm [18].

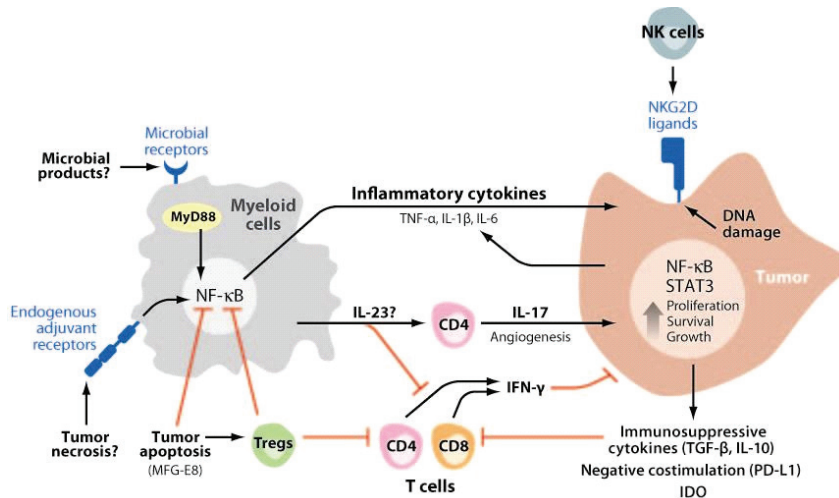
Editing refers to the changes that occur spontaneously as a tumor develops under the influence of an intact immune system and the understanding of immunoediting mechanisms has important implications for cancer immunotherapy in humans, and *avoiding immune destruction* has recently been proposed as an emerging hallmark of cancer by Hanahan and Weinberg [15].

### 2.3 The immune response may promote tumorigenesis

The immune system interacts with tumors throughout their development, and the consequences of this interaction have implications for cancer therapy. Some host responses may inhibit tumor growth, as discussed previously, but immune responses can also *promote* cancer by provoking chronic inflammation and drive growth, survival and angiogenesis (see Figure 1).

The proinflammatory cytokine tumor necrosis factor (TNF- $\alpha$ ), largely produced by macrophages and mast cells, is implicated in early cancer development as well as in infections. TNF- $\alpha$  promotes cell growth, survival and angiogenesis and the recruitment of immune effector cells. The events downstream of TNF- $\alpha$  are not well known, but nuclear factor- $\kappa$ B (NF- $\kappa$ B) family transcription factors appear to be linked to TNF- $\alpha$  and cancer [26].

NF- $\kappa$ B controls the transcription of several proteins involved in cell survival, division, growth and is important for the production of many inflammatory cytokines, including TNF- $\alpha$  itself. In an animal model for colon cancer, ablation of NF- $\kappa$ B in immune cells led to reduction in tumor growth and ablation in the colonic epithelium decreased tumor incidence [27]. For further details, see Figure 1.



**Figure 1.** Complex interactions with the immune system shape tumor development. Chronic inflammatory response can be initiated by microbial products or endogenous adjuvants released from necrotic cells. These signals activate nuclear factor- $\kappa$ B (NF- $\kappa$ B) in myeloid cells, leading to the production of inflammatory cytokines (TNF- $\alpha$ , IL-1 $\beta$ , and IL-6), which in turn activate NF- $\kappa$ B in the tumor. Tumor-intrinsic NF- $\kappa$ B activation promotes growth, survival, and proliferation. IL-23 produced by myeloid cells can promote the generation of IL-17-secreting T cells, which can further support tumor growth. Genotoxic stress in tumor cells can activate NK cell ligands, which can synergize with endogenous tumor-specific CD4<sup>+</sup> and CD8<sup>+</sup> T cells that produce IFN- $\gamma$  and restrict tumor development. Tumors can suppress nascent immune responses through a variety of mechanisms, including immunosuppressive cytokines (TGF- $\beta$  and IL-10) and metabolites [indoleamine 2,3-dioxygenase (IDO)] and the expression of negative costimulatory molecules [programmed death ligand 1 (PD-L1)]. Tumors can also promote T<sub>reg</sub> recruitment and differentiation, in part through the recognition of apoptotic cells by the MFG-E8 (milk fat globule epidermal growth factor 8) pathway. (Reprinted with permission from Copyright Clearance Center. Annual Review of Immunology. Dougan, Dranoff<sup>[26]</sup> Copyright 2014).

## 2.4 Tumor immune escape mechanisms

Tumors themselves might develop several strategies allowing them to escape immune surveillance and destruction. Tolerance induction is one of the major mechanisms involved and is induced by several steps.

- Mutation or down-regulation of MHC molecules, particularly MHC Class I, has been documented in several tumor types<sup>[28]</sup>. In many cases, individual HLA alleles are selectively lost, but complete MHC Class I loss has been observed in tumors. The most common mechanism for total MHC Class I loss is mutations in the  $\beta$ 2-microglobulin genes. Loss of MHC Class I is an escape mechanisms for CTL recognition.
- Tumor cells often have altered expression of molecules involved in antigen presentation and processing, due to down modulation. Such transporter proteins associated with antigen processing (TAP1), low molecular mass-protein 2 (LMP2) and LMP1 proteasome components, results in suboptimal peptide delivery to MHC Class I<sup>[28, 29]</sup> and impaired antigen presentation.
- Expression of anti-apoptotic molecules and down-regulation or mutation of pro-apoptotic molecules renders tumor cells resistant to apoptosis<sup>[30]</sup>. Due to acquired

defects in Fas signal transduction, cancer cells may be resistant to the Fas ligand (FasL)-mediated cytotoxicity of antitumor T and NK cells. Fas resistance enables the tumor cells to express their own FasL, which delivers an apoptotic death signal to activated T and NK cells, thereby inhibiting lymphocyte infiltration into FasL-expressing tumor nests.

- Tumor cells themselves or tumor stroma can secrete immune suppressive cytokines such as IL-10, TGF- $\beta$  and vascular endothelial growth factor (VEGF). For example, TGF- $\beta$  inhibits the activation, proliferation and differentiation of T cells, and suppress the activity of CTLs and DC, while inducing differentiation of regulatory T cells ( $T_{regs}$ )<sup>[31-33]</sup>. Tumors also express the enzyme indoleamine 2, 3-dioxygenase (IDO) which prevents proliferation of CD8+ cells and induce apoptosis of CD4+ T cells<sup>[34, 35]</sup> (see Figure 1).

## 2.5 Suppressive tumor infiltrating leucocytes

Evidence suggests that, unlike cells found in lymphoid organs that respond to acute infections, immune cells in tumors are dysregulated and functionally impaired and use immune-regulatory patterns to generate an *immune-suppressive environment*. Tumor infiltrating leukocytes subsets can play strikingly antagonistic functions<sup>[36]</sup>.

The suppressive microenvironment of tumors is further established through the activity of *myeloid derived suppressor cells (MDSC)*, *activated macrophages*, and  $T_{regs}$ <sup>[37-40]</sup>.

*MDSCs* are described as a population of CD11b<sup>+</sup> Gr-1<sup>+</sup> cells in mice with the ability to suppress CD8+ T cell antitumor activity<sup>[41, 42]</sup>. GM-CSF is important in driving the expansion of these cells<sup>[43]</sup>. Studies have shown that MDSCs are a heterogeneous population of cells consisting of cells of monocytic as well as granulocytic origin<sup>[43]</sup>. In humans, the markers for MDSCs are not clearly defined, but the phenotype resides within a population of LIN-HLA-DR-CD33+ cells<sup>[43]</sup>.

MDSCs express high levels of two enzymes called inducible nitric oxidase synthase (iNOS) and arginase, leading to depletion of, for T cells, essential nutrients<sup>[43, 44]</sup>. Arginase depletes L-arginine from the tumor environment and induces cell cycle arrest in T cells<sup>[45]</sup>. iNOS leads to increased production of nitric oxide (NO). High levels of NO block specific signal transduction in T cells, leading to suppressed production of IL-2<sup>[43, 44]</sup>. Other mechanisms include sequestration of cysteine, an essential amino acid for T cells and secretion of IL-10 which has suppressive properties. The ability to skew the differentiation of CD4+ T cells to regulatory cells has also been suggested<sup>[46]</sup>. Taken together, MDSCs exercise many functions that blunt effector T cell responses.

*Tumor associated macrophages (TAM)*, are divided into two subsets, M1 and M2. INF- $\gamma$  drives the polarization towards M1, IL-4 polarizes macrophages towards M2 profile<sup>[36]</sup> promoting tumor evasion<sup>[40, 47, 48]</sup>. M2-skewed macrophages produce lower levels of proinflammatory cytokines, such as IL-1 $\beta$ , TNF- $\alpha$  and IL-12 and higher levels of immunosuppressive cytokines, such as IL-10, TGF- $\beta$  and VEGF. Dysfunctional macrophages have an impaired ability to mediate direct lysis of malignant cells in comparison to M1-type cells<sup>[49]</sup>. The most frequent TAM phenotype seems to be M2<sup>[36]</sup>.

*Regulatory T cells* ( $T_{\text{regs}}$ ) also infiltrate many tumors<sup>[50]</sup>, their immune suppressive profile was established in a series of experiments showing that CD25+ T cell depletion improved antitumor immunity in mouse models<sup>[51-53]</sup>. Characteristics of CD4+CD25<sup>high</sup>FOXP3+ T cells ( $T_{\text{regs}}$ ) are their anergic state and their capacity to inhibit CD8+, T cells, DCs, NK cells, NKT cells and B cells<sup>[54]</sup>.  $T_{\text{regs}}$  constitute 5-10% of human CD4+ T cells and have a pivotal role in maintenance of immunologic self-tolerance and avoiding auto-immunity<sup>[55]</sup>. They are subdivided into “natural”  $T_{\text{reg}}$  ( $nT_{\text{reg}}$ ) cells.  $nT_{\text{reg}}$  develop in the thymus and act through contact-depending mechanisms, maintaining self-tolerance and preventing autoimmunity<sup>[56]</sup>, “induced”  $T_{\text{reg}}$  ( $iT_{\text{reg}}$ ) cells, are induced to differentiate in the periphery and mediate suppression by contact-independent mechanisms, including production of TGF- $\beta$ <sup>[56, 57]</sup>.  $T_{\text{regs}}$  also secrete suppressive IL-10 creating an environment that blunts effector antitumor responses by CD4+, CD8+ and NK cells and by acting as a competitive sink regarding IL-2, due to high affinity for IL-2, in the tumor environment<sup>[58, 59]</sup>. Activated T cells become exquisitely sensitive to the lack of IL-2, leading to cell death under conditions of IL-2 deprivation<sup>[60]</sup>.

The transcription factor FoxP3 is critical for the development of the functional characteristics of regulatory T cells<sup>[61, 62]</sup>. FoxP3-knockout mice, and humans with homozygous mutation of FoxP3 (encoded on the X chromosome) develop autoimmune syndromes involving multiple organs<sup>[57]</sup>. An increase of  $T_{\text{regs}}$ , both in the periphery and within tumors, is seen in human cancer<sup>[63, 64]</sup> and the presence of these cells correlates with poor prognosis in ovarian cancer, breast cancer and hepatocellular carcinoma<sup>[50]</sup>.

## 2.6 Immunosuppressive checkpoints

Cytotoxic T-lymphocyte-associated antigen 4 (CTLA-4), is an important immune-checkpoint receptor that normally prevents excessive and uncontrolled immune responses. It is expressed exclusively on T cells where it primarily regulates the amplitude of the early stages of T cell activation.

CTLA-4 counteracts the activity of the T cell co-stimulatory receptor CD28<sup>[57]</sup>. CD28 does not affect T cell activation unless the TCR is first engaged by cognate antigen. Once antigen recognition occurs, CD28 signalling strongly amplifies TCR signalling to activate T cells. CD 28 and CTLA-4 share identical ligands; CD80 (also known as B7.1) and CD86 (also known as B7.2). As CTLA-4 has a much higher affinity for both these ligands, it has been proposed that it's expression on the surface of T cells dampens the activation of T cells by outcompeting CD28 binding to CD80 and CD86<sup>[57]</sup>, thereby acting *immunosuppressive*. Even though CTLA-4 is expressed by activated CD8+ effector T cells, the major physiological role of CTLA-4 seems to be through distinct effect on two major subsets of CD4+ T cells; downmodulation of helper T cell activity and enhancement of the immunosuppressive activity of regulatory  $T_{\text{regs}}$ <sup>[57]</sup>.

In contrast to *CTLA-4* the major role of *Programmed death 1 (PDI)* is to limit the activity of T cells in peripheral tissues at the time of an inflammatory response to infection and to limit autoimmunity. PDI thus predominantly regulates effector T cell activity within tissue and tumors, whereas CTLA-4 predominantly regulates T cell activation<sup>[57]</sup>. PDI is highly expressed on  $T_{\text{regs}}$  where it may enhance its proliferation in the presence of its ligand<sup>[57]</sup>. PDI is also expressed on B cells and NK cells – limiting their lytic activity<sup>[57]</sup>. PDI is

normally transiently induced following immune activation but chronic antigen exposure, such as occurs with chronic viral infection or cancer, can lead to high levels of persistent PD1 expression, which induces a state of exhaustion or anergy among cognate antigen-specific T cells <sup>[57]</sup>.

*The PD1 ligands* are commonly upregulated on the tumor cell surface in many human cancers <sup>[65]</sup>. The major PD1 ligand that is expressed is programmed death ligand 1 (PDL1) <sup>[57]</sup>. PDL1 (present or not present) status in tumors has been correlated with poor prognosis, better prognosis or without correlation to prognosis, probably due to differences in analyses, tumor types and stages <sup>[57]</sup>.

## 3 IMMUNOTHERAPY IN CANCER

### 3.1 Background

#### General Components

Immunotherapy against cancers includes active, passive or immunomodulatory strategies. While passive immunotherapy administers exogenously produced components such as antibodies, active immunotherapies augment the ability of the patients own immune system to mount an immune response able to recognize and eliminate malignant cells. Immunomodulatory agents are not targeted at specific antigens but enhance general immune responsiveness.

### 3.2 Passive immunotherapy

Impressive clinical responses to monoclonal antibodies (mAb) therapy have already been achieved and several are part of routine treatment, both in the adjuvant and palliative setting. Several murine, chimeric as well as humanized mAbs have been approved. Of these, no less than 14 mAbs, including naked reagents as well as mAbs coupled to antibiotics or radioactive isotopes, are nowadays authorized for use in cancer patients<sup>[66, 67]</sup>. mAbs that potentially exert antineoplastic effects belong to one of six classes<sup>[68]</sup>:

- 1) mAbs that target cancer cell-intrinsic prosurvival signal transduction cascades, e.g. cetuximab (Cet) targeting epidermal growth factor receptor (EGFR) (in CRC)
- 2) mAbs that interrupt thropic interaction between malignant cells and stroma, e.g. bevacizumab (Bev) targeting VEGF (in several adenocarcinomas)
- 3) mAbs that recognize antigens expressed on surface of tumor cells and initiate ADCC as well as CDC, e.g. Rituximab targeting CD20 (in lymphoma)
- 4) Bispecific mAbs, who can bind two antigenic targets, while retaining ability to exert immune effector functions e.g. catumaxomab, an anti CD3 and anti-epithelial cell-adhesion molecule (EpCAM) mAb (in EpCAM+ tumors)
- 5) Immunoconjugates e.g. <sup>90</sup>Y-ibritumomab (radionucleotide coupled) anti CD20 mAb (in lymphoma)
- 6) Immunostimulatory mAbs, that facilitate immune response by balancing immunosuppressive mechanisms, e.g. ipilimumab targeting CTLA-4 (in melanoma).

### 3.3 Active immunotherapy

#### General considerations

Active specific immunotherapy principally involves the use of TCV, with the aim to evoke a tumor specific immune response in cancer patients.

The major challenge in developing a successful TCV strategy is shifting the balance from tolerance to self-antigens towards a long-lasting therapeutic anti-tumor immunity, without inducing autoimmune toxicity to normal cells.

Animal models have played a crucial role in understanding the mechanisms of tumor immunology and different modalities of immunization and experimental models have been employed to determine whether TAAs can induce immune responses that are able to hamper the growth of a clinically manifest cancer.

The promising results have driven the attempts to move these applications from bench to bedside. To this day, however, only three vaccines have been approved by the Food and drug administration (FDA) for use in humans; Cervarix® and Gardasil® (in fact constituting preventative measures towards development of cervical carcinoma) and sipuleucel-T (Provenge®), a cellular preparation for therapy of metastatic hormone-refractory prostate cancer. In the veterinary field an immunotherapeutic DNA vaccine encoding human tyrokinase, Oncept™, has been approved as TCV for malignant melanoma in dogs. Oncept™ generates an effective antibody response against the dog tyrokinase and prolongs the lives of dogs even with advanced stage disease.

### 3.3.1 Tumor associated antigen for therapeutic cancer vaccines

Virtually any mutant, overexpressed or abnormally expressed protein in cancer cells, can serve as a target for cancer vaccines. Since the pivotal work done by pioneers in oncoimmunology, characterization and identification of hundreds of TAAs has been conducted and additional insights into the mechanisms whereby TAAs can break self-tolerance and elicit an immune response has been provided. Simply put, TAAs can be classified into four classes, see Table 1.

**Table 1.** Tumor antigens

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#### Classes of tumor antigens <sup>[69]</sup>

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- (1) Truly exogenous, non-self TAAs (invariably of viral origin)
  - (2) Unique, mutated TAAs (stemming from cancer cell-specific genetic alterations)
  - (3) Idiotypic TAAs (the unique way whereby the B cell receptor expressed by some clonal hematopoietic malignancies is rearranged)
  - (4) Shared TAAs (which are also expressed by normal cells, mostly at lower levels)
- 

Shared TAAs include cancer-testis and differentiation antigens that are either silent or expressed at only low levels in normal tissue but are transcriptionally activated in certain tumors e.g. CEA in CRC and human telomerase reverse transcriptase (hTERT) in pancreatic cancer.

Optimally designed TCV should combine the best tumor antigens and delivery strategies to achieve a clinical result. A National Cancer Institute (NCI) immunotherapy workshop was held in 2007 to rank agents with high potential to serve as immunotherapeutic drugs. The ranking was based on the likelihood for efficacy. A priority-ranked list of cancer antigens weighed the criteria for the “ideal” cancer antigen to provide focus for prioritized translational research <sup>[70]</sup>. The resulting criteria were as follows: (1) therapeutic function, (2) immunogenicity, (3) role of the antigen in oncogenicity, (4) specificity, (5) expression level and percentage of antigen-positive cells, (6) stem cell

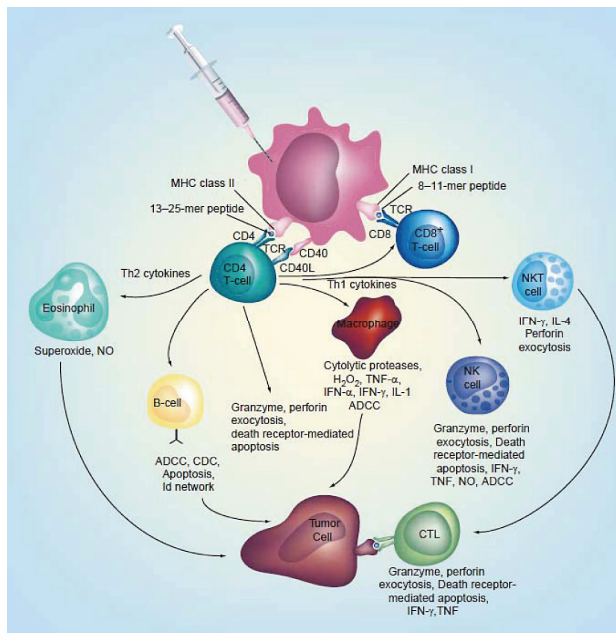


expression, (7) number of patients with antigen-positive cancers, (8) number of antigenic epitopes and (9) cellular location of antigen expression.

### 3.3.2 Desired mechanisms of action of TCV

The processes required to mount an effective anti-tumour response can be subdivided into four different steps [71]. In the first, TAAs must be captured, processed and presented by DCs (or directly presented by tumour cells). The second step requires suitable activation/maturation signals that allow DC to differentiate migrate to lymph nodes and present TAAs to naïve T cells. The third step involves expansion of T cells in sufficient numbers capable of recognizing and eliminating tumour cells. In the final step, antigen-educated T cells must leave the lymph node, traffic to infiltrate the tumour and persist long enough to kill the malignant cells.

Thus CD4+, CD8+ T cells, and B cells can all be activated in the immune response against cancer. For intracellular TAAs, CD8+ CTLs may be the primary effector cell, but the activity and longevity of these will likely be enhanced by CD4+ Th1 cell activation. For cell surface antigens, *both CTL and humoral responses may mediate antitumor activity*, again attesting to the importance of activating both CD8+ and CD4+ T cells [72].



**Figure 2.** Summary of potential effector mechanisms involved in antitumor immunity induced by vaccine. Immature or intermediate dendritic cells acquire the antigen(s) at the site of vaccination. In response to maturation and activation signals, dendritic cells migrate to draining lymph nodes, upregulate costimulatory molecules, such as B7, and present tumor-associated antigen-derived peptides (13-25-mer) to cognate CD4+ T-cells in the context of MHC class II molecules. The interactions of additional costimulatory signals, such as CD40-CD40L, further promote dendritic cell maturation providing help for efficient priming and activation of CD8+ T-cells. [3]. Dendritic cells can present 8-11-mer peptides derived

from endogenous antigens (e.g., viral vector-encoded tumor-associated antigens) or exogenous antigens (e.g., recombinant protein vaccine) in the context of MHC class I molecule to CD8+ T-cells. CD4+ T-cells producing Th1 cytokines, such as IL-2, further contribute to the clonal expansion of CD8+ T-cells [73, 74]. Tumor-specific CD8+ T-cells migrate to the sites of tumor metastasis where they encounter peptide-MHC complexes presenting the tumor antigen on tumor cells. CTLs are able to kill tumor cells by perforin-mediated cell lysis or apoptosis mediated through granzymes. CTLs can also kill target cells through death receptor mediated apoptosis via Fas ligand or TNF-related apoptosis inducing ligand (TRAIL). CTLs secreting IFN- $\gamma$  and TNF- $\alpha$  (or TNF- $\beta$ ) may elicit direct or indirect cytotoxic activity [73]. [5]. Activated CD4+ T-cells may also kill tumor cells by using similar pathways as CTLs. CD4+ T-cells

producing Th1 cytokines may stimulate effector cells of the innate immune system, such as macrophages, NK and natural killer T-cells, which might exert antitumor effects by several mechanisms. CD4+ T-cells secreting Th2 cytokines may attract and activate eosinophils releasing their cytotoxic granule content.

Th2 cells may also activate B-cells producing antibodies, which may activate B-cells producing antibodies, which may contribute to tumor cell destruction by ADCC and CDC<sup>[74, 75]</sup>. Antibodies may also induce an idiotypic network cascade or tumor-cell apoptosis.

ADCC: Antibody-dependent cellular cytotoxicity; CDC: Complement-dependent cytotoxicity; CTL: Cytotoxic T-lymphocyte; IFN: Interferon; IL: Interleukin; MHC: Major histocompatibility complex; NK: Natural killer; NO: Nitric oxide; TCR: T-cell receptor; Th: T-helper; TNF: Tumor necrosis factor.

(Reprinted with permission from Expert Review of Vaccines. Mosolits et al.<sup>[76]</sup> Copyright 2014).

### 3.4 Adjuvants

Adjuvants are substances or interventions that, combined with an antigen, enhance its immunogenicity and the desired type of immune response. The preferred type of response in cancer is one in which IFN- $\gamma$  producing Th1 lymphocytes and cytotoxic T lymphocytes predominate. Adjuvants discussed here fall into one of the following categories: biological or chemical adjuvants and cytokines/chemokines.

- Biological adjuvants are recognized as non-self and known to induce migration of APCs at the site of delivery and hence the induction of an inflammatory response. APCs may then capture and process TAA at the same site. The most commonly used biological adjuvant is the bacillus Calmette-Guerin (BCG), diphtheria toxin and tetanus toxoid.
- Chemical adjuvants function similarly to biological ones and some, for example, aluminium hydroxide (alum) and incomplete Freud's adjuvant (IFA) provide a matrix that sequesters antigen at the delivery site, allowing a timed release of antigen to APCs.
- Cytokines used for decades include GM-CSF and IL-12, known for inducing recruitment, migration, stimulating cross-presentation of DCs and promoting Th-1 differentiation. GM-CSF is known to augment both humoral and cellular immunity, although immune suppression may be seen at high doses<sup>[77, 78]</sup>. GM-CSF also induces the expansion of NKT cells that may contribute to immunity against tumors<sup>[79]</sup>.
- Microbes often elicit immune responses by activating pattern-recognition receptors such as members of the Toll-like receptor (TLR) family. TLR-signalling play an important role in both the innate and acquired immune system and TLR agonist has been evaluated as adjuvants. TLR agonists, such as unmethylated CpG-motifs (CpG) and lipopolysaccharid (LPS) derivate monophosphoryl lipid A (MPL) have been in the focus for evaluation lately. However, the use of TLR agonists is often associated with severe toxicity, resulting from non-specific activation of lymphocytes<sup>[80]</sup>.

### 3.5 Therapeutic cancer vaccines approaches

A plethora of active vaccination strategies exist with attributed advantages and disadvantages.

Discussed in this thesis:

- Whole cell-based vaccines
- Dendritic cell vaccine strategies
- Peptide and protein vaccines
- DNA vaccines

### **Whole cell-based vaccines**

Whole cell-based vaccines have been investigated for decades. They have the advantage to deliver a diverse panel of TAA and to simultaneously provide both CD8+ and CD4+ epitopes, and particularly autologous tumor-cell derivative potentially include unique TAAs. Autologous or allogeneic tumor cells are processed (lysates or irradiated cells) to optimize the release of their antigens, and are injected together with adjuvants or haptens. However, it is difficult to obtain and individually prepare vaccines for each patient. To avoid this problem, tumor cell vaccines have been prepared from allogeneic tumor cell lines. This is feasible as many tumors have overlapping antigen expression<sup>[81, 82]</sup>. This is attractive as tumor antigen-specific immune responses can be initiated by cross-priming, by-passing the need to match the MHC haplotype of the patient to the vaccine platform<sup>[83]</sup>.

### **Dendritic cell vaccine strategies**

Dendritic cell vaccine strategies are attractive as activated and mature DCs express high levels of both MHC Class I and II molecules for priming of CD8+ and CD4+ T cells, and additionally have co-stimulatory signals for T cell activation. By providing DCs with TAA it is possible to induce tumor-specific immune responses. Vaccination strategies based on DCs can be divided into three categories.

The 1<sup>st</sup> group encompasses strategies where DCs are generated by culturing patient-derived hematopoietic progenitor cells or monocytes with specific cytokine combinations in the presence of adjuvants and loaded with TAAs *ex vivo*. The DCs are then re-infused to the patient, most often *i.d.*, together with an adjuvant. *Ex vivo* DC vaccines exhibit a good safety profile and can induce TAA-specific CD8+ and CD4+ T cells, as seen with the approval of CD-based vaccine Sipuleucel-T for use in patients with prostate cancer<sup>[84]</sup>.

The 2<sup>nd</sup> group of promising DC vaccine strategies comprise TAA delivery to DC *in vivo* and is achieved by coupling TAAs to monoclonal antibodies or other vectors that recognize specific DC surface receptors<sup>[85]</sup>. Encouraging results for *in vivo* – targeted vaccination came from the use of engineered lentiviral vectors encoding the human melanoma antigen NY-ESO-1<sup>[86, 87]</sup>.

The 3<sup>rd</sup> group includes approaches based on DC-derived exosomes<sup>[85]</sup>. DC derived exosomes (small membrane-surrounded vesicles originating from fusion of plasma-membrane and multivesicular bodies, released by DCs) are 100-fold more enriched with MHC Class II molecules than DCs. They have shown to be capable of inducing immune responses once loaded with TAAs and inoculated *in vivo* in animal models<sup>[85]</sup>.

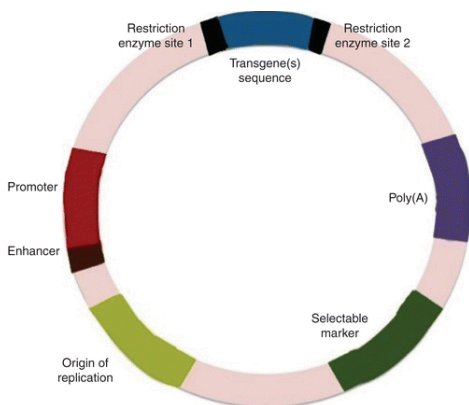
### **Protein and peptide vaccine**

Protein and peptide vaccine (Papers I and IV) strategies are based on the administration of high doses of the TAA(s) in order to load empty MHC molecules on APCs.

Tumor-derived *peptide* epitopes constitute the simplest cancer vaccine formulation delivered i.d. together with an adjuvant<sup>[88]</sup>, but this strategy requires prior knowledge to the precise definition of MHC Class I and II epitopes and are thus HLA-restricted. Other drawbacks include the weak immunogenicity of short peptides. Many immunodominant epitopes from TAAs have a low to intermediate binding affinity for the MHC molecule and are subdominant epitopes recognized by low affinity T cells that have escaped central tolerance. The tumor may also have ceased to express the epitope (antigen-variant escape mutants). The immunogenicity may be enhanced by introducing lipid, carbohydrate or phosphate groups, by introducing protease-resistant peptide bounds and thereby prolonging the peptides half-life in vivo<sup>[89]</sup>. On the other hand, manufacturing on a large scale is affordable and peptide vaccinations are safe with no potential for recombination. The greatest advantage of immunizing with MHC Class II peptides is the concomitant generation of CD4+ cells T cells, particularly the generation the Th1 subtype, essential for the generation of a robust CTL response. It is anticipated that the peptide is intracellularly processed to yield shorter CD8+ epitopes<sup>[12]</sup>. These are presented by APCs on MHC Class I molecules (cross-presentation) to produce a CTL response, ensuring a maximal immune response.

### DNA vaccines

Unlike other vaccines, **DNA vaccines** (Papers II and III) do not contain the antigens themselves but are simple vehicles for in vivo transfection and antigen production. The circular DNA plasmid is made of the plasmid backbone, containing the origin of replication, the antibiotic resistance gene, a transcriptional unit that contains the eukaryotic promoter, the transgene(s), and a polyA tail that aids in the stability and translation of the messenger ribonucleic acid (mRNA)<sup>[90]</sup>. A schematic picture is shown in Figure 3.



**Figure 3.** Basic design of plasmid DNA vaccine vector. Therapeutic and prophylactic DNA vaccines for Human immunodeficiency virus (HIV-1). (Reprinted with permission from Copyright Clearance Center (Expert Opinion of Biological Therapy. Ramirez et al<sup>[90]</sup>. Copyright 2014)

These plasmids are delivered in the form of circular DNA or within appropriate delivery vectors to the intradermally, subcutaneously or intramuscularly by various delivery methods.

In all scenarios, using the host cellular machinery, the plasmid enters the nucleus of transfected local cells, including resident APC<sup>[91]</sup>. Here, gene expression from the plasmid

is followed by generation of the transgenes(s) <sup>[91]</sup>. DNA vaccines induce protein synthesis in vivo increasing the possibility to utilize both the MHC Class I and II antigen presenting pathways to induce humoral as well as CD4+ and CD8+ T cell responses <sup>[91, 92]</sup>. The relevant epitopes do not need to be defined, nor is the vaccine limited to particular HLA genotypes, as the whole protein will be produced and processed in the host cell.

However, whereas professional APCs are very efficient at direct presentation, myocytes and keratinocytes are not, as they express detectable yet rather low levels of MHC Class I and co-stimulatory molecules <sup>[92, 93]</sup>. The induction of immunity following TAA expression must then proceed via *cross-presentation* (see Section 1.6).

As compared to cell-based and recombinant preparations many advantages with DNA plasmids exist. They are stable at room temperature, can thus be stored without effort and are rapidly produced in bacteria in high amounts at low cost <sup>[94]</sup>. Additionally, DNA vaccines are inherently immunogenic due to CpG motifs in the DNA backbone, able to bind and activate specific TLR who constitute part of the innate immune system <sup>[89, 90]</sup> and as mentioned previously DNA plasmids need not be HLA specific to elicit an immune response as the whole protein will be produced in the host cell.

DNA vaccines are also highly flexible and can be designed to encode both TAA and immunological components that may enhance the immune response. They can be engineered to express non-self-proteins (in additions to the TAA gene) that exert adjuvant effects, as fragments of tetanus toxin <sup>[95]</sup>, pseudomonas aeruginosa exotoxin <sup>[96]</sup> or other immunostimulatory factors, such as cytokines, including IL-2, IL-12 and GM-CSF <sup>[69]</sup>. Furthermore, the plasmids can be engineered to alter the intracellular routing of the TAAs, resulting in the preferential activating of humoral (when TAAs are targeted to the endoplasmatic reticulum where folding and secretion can occur by adding a gene coding for a signal sequence) or cellular (if TAAs are targeted to the cytosol or – by adding genes encoding for ubiquitin aimed to enhance degradation and peptide production in the proteosome) immunity <sup>[97, 98]</sup>.

A major safety concern has been mutagenesis, which may trigger oncogenes. Studies using DNA vaccines has however not exhibited mutagenesis <sup>[99]</sup>. No association between plasmid DNA and genomic DNA were seen in mice following DNA vaccination, towards HIV-1 <sup>[100]</sup>. Vaccine-related adverse events in humans were mild and tolerable using the same DNA vaccine <sup>[101]</sup>.

For plasmid immunization, the aim is to rapidly induce an immune response, while for safety reasons; the plasmid should preferably be cleared rapidly from the body. During the efforts to prove the safety of plasmid DNA, the biodistribution of plasmids have been studied in several animal species, and the results all indicate that the plasmid is rapidly cleared from the body <sup>[100]</sup> and found exclusively at the site of injection at later time points. Additionally, studies with DNA vaccines have shown that even after multiple immunizations, anti-DNA antibodies are not produced <sup>[91]</sup>.

The vast majority of clinical trials with DNA vaccines to date have utilised naked DNA plasmids.

### 3.6 Delivery route and delivery systems

Induction of tolerance or activation of immune cells may depend on the nature of the TAA, but also the formulation, dose, route, delivery system and schedule of the vaccine. In particular, delivery of DNA vaccines has been the focus for attention for a long time. Intramuscular injections were commonly used during early tests with large animals and humans resulting in relatively poor efficacy although robust immune responses were seen in mice <sup>[69]</sup>. This probably stems from the fact that the efficacy of DNA vaccines administered i.m. strictly depends on the injected volume. A high hydrostatic pressure not only augments the uptake of the DNA vaccine by myocytes and resident APCs but also promotes tissue damage, resulting in the release of danger signals <sup>[69]</sup>. Scaling up the volume needed for DNA vaccines i.m. to humans is not always feasible why alternative routes have been sought for. Some DNA vaccine delivery methods have focused on delivering DNA to skin due to the presence of large quantity of immune cells there, including APCs and lymphocytes, in comparison to muscle <sup>[90]</sup>.

**Electroporation (EP)** (Paper III) has been used, to enable molecules to be delivered intracellularly *ex vivo*, for several years and has recently been tested *in vivo* to aid in transfecting cells. EP consists in the electrical stimulation of skin or muscle immediately after delivery of naked DNA. EP delivers a local controlled electrical field to cells, unstablising the cellmembranes and thereby allowing molecules, such as plasmids, to easily penetrate the cells and enhancing transfection <sup>[102, 103]</sup> While the electrical field is applied transient pores appear after which the cell can either heal-closing off the pores, or die due to lysis depending on the length and amount of the electrical field exposure <sup>[90]</sup>. I.m. EP is associated with an increase in transfection efficacy (enabling less injected volume) and local tissue injury resulting in danger signals <sup>[69]</sup>.

EP i.m has been the most commonly used technique in preclinical models for practical reasons <sup>[104]</sup> but could prove difficult in translational human studies due to painful muscle contractions. In mice skin electroporation was performed after prostate-specific antigen (PSA) DNA administration i.d. A robust PSA-specific CD8+ T cell response was induced after one single DNA injection <sup>[104]</sup>. Laddy et al conducted a head-to-head comparison of EP-augmented i.m. and i.d. delivery of equal amounts of influenza virus-encoding plasmids in monkeys. Results revealed that i.m. EP induced the highest levels of cellular immune responses, whereas i.d. EP was superior for induction of antibodies <sup>[105]</sup>.

The introduction of *in vivo* electroporation for increasing the uptake and immunogenicity of plasmid vaccines raised concerns that the method might affect the persistence of plasmid DNA in the target tissue and pose a risk of integration into the host genome. In mice, immunized and EP-augmented with a HIV DNA plasmid, no association between plasmid DNA and genomic DNA could be detected <sup>[100]</sup>.

Details about a **needle-free jet-injection device**, Biojector (Paper II, III), will also be discussed within the scope of this thesis. Biojector, is a CO<sub>2</sub>-propelled device that injects plasmids as a highly focused liquid stream and is distributed in the i.d. or i.m. depending on the amount of pressure used <sup>[106]</sup>. This has been shown to enhance antigen expression as compared to conventional needle, most probably due to a larger area and thus larger number cells being targeted <sup>[107]</sup>. In a mice model, a plasmid CEA66-DNA vaccine

construct, was safe and induced both CEA-specific CD4+ and CD8+ T cell responses (IFN- $\gamma$ ) when delivered i.d. by Biojector, however humoral responses were weak<sup>[108]</sup>.

**Gen-gun immunization** is carried out by propelling DNA-coated gold particles into the cells of the skin and Langerhans cells, which were shown to migrate rapidly to regional lymph nodes<sup>[109]</sup>, generating CD8+ T cell responses.

The potency of the gen-gun approach was shown in a clinical trial in humans; patients had not generated immunity to the licensed recombinant HBsAg vaccine but did produce antibodies following gene gun-delivered hepatitis B surface antigen (HBsAg) DNA<sup>[110]</sup>. Newly developed i.d. **tattoo DNA vaccine delivery** consist of fine metal needles that oscillate at a high frequency and puncture the skin, leading to transfection of skin associated cells. Robust T cell responses have been seen in a phase I melanoma studies<sup>[111]</sup>.

### 3.7 Immunomodulating strategies

For TCV to be successful, both sides of the adaptive system must be manipulated. The effector system should be stimulated and the suppressor system inhibited.

Combination therapies may produce synergistic antitumor responses and certain chemotherapeutic agents, rather than being immunosuppressive can, under certain conditions act as strong adjuvants for active immunotherapy<sup>[112, 113]</sup>. Drugs may induce immunogenic tumor cell death resulting in emission of danger signals and TAA<sup>[114]</sup>. Secondly, chemotherapy may indirectly stimulate immune components, by inducing transient lymphodepletion, by affecting immunosuppressive mechanisms or by stimulating immune effectors<sup>[115]</sup>.

Cyclophosphamide, a nitrogen mustard alkylating agent, is used for treatment of a variety of tumors. However, the biological activities of cyclophosphamide are dose-dependent<sup>[115]</sup>. Although regarded as immunosuppressive, cyclophosphamide has been shown to act as a strong immunomodulator for active immunotherapy when used with carefully defined doses<sup>[112, 113, 116]</sup>. Historically, 300 mg/m<sup>2</sup> has been used for enhancing immunotherapies in phase III trials<sup>[117]</sup>. Treatment with low-dose cyclophosphamide has been shown to transiently reduce T<sub>reg</sub> levels and enhance tumor-reactive T cell responses when used alone or in combination with active immunotherapy<sup>[118, 119]</sup>.

Novel monoclonal antibodies can directly antagonize immune suppressive check points, (CTLA-4 and PD1), they target lymphocyte receptors or their ligands in order to enhance the antitumor activity<sup>[57]</sup>.

Two fully humanized antibodies; ipilimumab and tremelimumab, exist. Ipilimumab is approved for treatment of advanced melanoma since 2010. In combination with TCV, a randomized three-arm trial of patients with advanced melanoma showed survival benefit for ipilimumab  $\pm$  vaccine. Patients received either: a peptide vaccine of melanoma-specific gp100 alone; the gp100 vaccine plus ipilimumab; or ipilimumab alone. There was a 3.5 month survival benefit for patients in both groups receiving ipilimumab compared to the group receiving gp100 alone<sup>[120]</sup>.

Clinical experience with PD1 antibodies is less extensive than CTLA-4, but initial results look promising<sup>[121]</sup>.

## 4 IMMUNOTHERAPY TARGETING CEA AND hTERT

### 4.1 Carcinoembryonic antigen

CEA, a member of the immunoglobulin family of molecules, is a cell-surface-expressed 180 kDa glycoprotein. Many functions have been linked to CEA. CEAs primary role is in cellular adhesion and it dimerizes through a unique two-point intracellular adhesion mechanism and can interact with several integrins in the intercellular matrix, suggesting that it may promote malignant cell proliferation and metastasis. It has been shown to inhibit cell death induced by loss of anchorage to the extra cellular matrix. Also, it promotes cells entering G0-like state, thus facilitating accumulation of additional oncogenic events [122-124]. CEA has been shown to induce cellular secretion of cytokines that promote cellular adhesion, thereby increasing the malignant potential of cells overexpressing this protein [125].

In the adult human body, CEA protein expression is low. CEA is normally expressed to a low level in gastrointestinal epithelium and to a higher extent during fetal development. CEA expression in colorectal tissues indicated a CEA tissue content below 300 ng/mg protein in healthy donors, less than 2000 ng/mg protein in normal mucosa of cancer patients, and 200 to >10 000 ng/mg of protein in tumor biopsies [125]. It is overexpressed in nearly all colorectal cancers and to a high extent overexpressed in pancreatic adenocarcinomas, breast cancer and non-small-cell lung cancers [126]. In cancer patients, significantly augmented cell-surface expression of CEA has been associated with more advanced disease and increased relapse rates compared to patients expressing lower levels of CEA. Neoplastic tissues can secrete CEA into the bloodstream, where it can be used as a serological circulating marker, mainly in colorectal cancer [125].

The CEA molecule is synthesized as a precursor protein with N- and C-terminal signal peptides. These signal peptides target the protein through the endoplasmic reticulum (ER), where it is heavily glycosylated, and subsequently to the cell membrane [125]. More than 50% of the molecule consists of N-linked oligosaccharides on 28 potential sites of N-glycosylation [127]. Both signal peptides are removed during post-translational processing. The mature protein consist of a 107-amino acid N-terminal Ig V-like region; three 178 amino acid C2 Ig-like repeating units called A1 B1, A2 B2 and A3 B3, respectively; and finally, the C-terminal signal peptide has been replaced by a glycoposphatidylinositol (GPI) membrane anchor [126] [72]. Furthermore, CEA may have different molecular weights in normal vs. cancer cells due to different N-glycosylation pattern in cancer cells [128] where hypoglycosylation is associated with tumorigenesis.

CEA protein is processed and presented on major histocompatibility complex (MHC) proteins for multiple alleles, including HLA A2, A3 and A24. T lymphocytes from healthy volunteers and cancer patients can recognise processed epitopes of CEA and can become activated to lyse CEA-expressing tumors.

CEA has as a TAA been identified as an attractive target for vaccination approaches against multiple types of cancer due to its pattern of expression and its likely function in



tumorigenesis. In a well-vetted priority-ranked list, composed to prioritize TAA for translational research based on predefined criteria, CEA ranked no.13 of 75 [70].

## 4.2 Therapeutic cancer vaccines targeting CEA

Several studies have shown that CEA can be used as a target for cancer immunotherapy protocols. It has previously been demonstrated that tolerance to CEA could be interrupted by mouse immunization with recombinant human CEA without leading to autoimmune disease [129-131]. Among the diverse CEA-based cancer vaccines in humans, DCs pulsed with agonist epitopes of CEA and recombinant virus-based vaccines have been most successful [128].

### 4.2.1 Dendritic based vaccines

Matsuda et al performed a pilot study on eight patients with advanced gastrointestinal malignancy [132]. HLA-A24-positive patients received CEA peptide-pulsed DCs subcutaneously (s.c.) every two or three weeks. One patient had a dramatic decrease of CEA-levels. The other patients experienced disease progression, except three who had disease stabilization lasting for 3, 4 and 5 months respectively. In a phase I-II study, the effect of DCs transfected with RNA encoding CEA was evaluated [133]. Twenty-nine patients with advanced cancer expressing CEA were included. Two patients had minor response, 3 patients showed stable disease (SD) lasting more than two years and 18 underwent progression.

### 4.2.2 Recombinant virus-based vaccines

PANVAC is a cancer vaccine delivered through the viral vectors rV and vF, both of which include transgenes for epithelial mucin 1 (MUC1) and CEA in combination with specific costimulatory molecules (TRICOM). Early clinical trials have demonstrated PANVACs safety and ability to activate antigen-specific T cell responses [134]. PANVAC in combination with GM-CSF was evaluated for efficacy in 25 patients with metastasized CRC, lung, breast gastric or ovarian carcinomas. Multiple previous chemotherapy treatments and short time since last chemotherapy treatment correlated with lack of immune response. Three patients had prolonged SD or improvement and partial response (PR) was seen. Several patients had prolonged survival [135]. A second clinical trial evaluated PANVAC in combination with GM-CSF in 26 patients with metastatic breast or ovarian cancer [136]. SD was seen in four patients, with progression free survival (PFS) of 4-6 months and overall survival (OS) 16.1-40.5 months. Minor response and on case of complete response (CR) was seen. A significant reduction in circulating T<sub>regs</sub> and increased CD4+ T cell responses were observed in the patient with CR and in another patient with SD.

In a set of Phase I trials, patients with stage III/IV pancreatic cancer who received PANVAC showed a slight increase in survival [134]. This led to a Phase III trial in which PANVAC was compared to best supportive care or second-line chemotherapy in patients who had failed first-line chemotherapy with gemcitabine [134]. The trial did not meet its primary endpoint of improved OS, possibly due to poor trial design that is – administering vaccine as a monotherapy to an inappropriate patient population [134]. An ongoing Phase II

trial is evaluating PANVAC in patients with CRC after complete resection of liver or lung metastasis.

#### 4.2.3 Recombinant protein vaccines

Patients vaccinated with a recombinant CEA protein in combination with GM-CSF mounted a humoral and cellular immune response of a significantly higher magnitude than patients in the non-GM-CSF group. A positive correlation between the anti-CEA IgG titer and overall survival was suggested<sup>[137, 138]</sup>.

#### 4.2.4 Yeast vaccines

Non-pathogenic yeast-strain has been shown to be effective vectors for TCV and can stimulate antigen-specific cellular and humoral immune responses without inducing neutralizing antibodies<sup>[139, 140]</sup>. They can be administered multiple times to increase immune response with no toxicity and are easy to produce<sup>[141]</sup>. Twenty-five patients with metastatic CEA-expressing carcinomas were treated with recombinant yeast designed to express CEA. Five patients had SD > 3 months and five patients showed evidence of CEA-specific T cells<sup>[142]</sup>.

### 4.3 hTERT

Telomeres are specialized structures at the end of human chromosomes composed of 1000-2000 non-coding repeats of TTAGGG nucleotide DNA sequences. They serve as protective “caps” at the end of chromosomes protecting them from DNA degradation and unwanted repair<sup>[143-145]</sup>. In normal human cells telomeres shorten with each successive cell division, and upon reaching critical lengths they elicit DNA-damage responses, thus activating cell cycle check points, leading to cell apoptosis. In contrast, cancer cells which develop chromosomal aberrations show activation or re-activation of telomerase upon exposure to DNA damage signal, thereby bypassing cell cycle checkpoints and leading to uncontrolled growth and proliferation<sup>[145, 146]</sup>.

Telomerase is a human ribonucleoprotein reverse transcriptase (hTERT) composed of two main subunits: the catalytic protein hTERT and the ribonucleoprotein template hTER<sup>[143, 144, 147]</sup>. Telomerase synthesizes telomeric DNA by continually adding single stranded 3' end of telomere in the 5' to 3' direction<sup>[143, 145, 148]</sup>. Telomerase, and specifically its catalytic subunit hTERT, is overactive in 85-90% of most cancers<sup>[147]</sup> and has become a widely accepted target for anticancer therapeutics. In normal non-malignant cells telomerase is present in embryonic, male germline and some adult stem cells. In most somatic cells telomerase is present in very low or almost non-detectable levels and is less active compared to cancer cells<sup>[149]</sup>.

One of the advantages of targeting telomerase is that rapidly progressive cancer cells have shorter telomeres compared to normal somatic cells and stem cells, that have not yet reaches critical lengths, due to end replication problem that occurs as a result of aging<sup>[150, 151]</sup>. By de novo synthesising TTAGGG repeats, telomerase can maintain cancer cell telomeres at stable length at all times, ensuring their immortal capacity. The difference in telomere length and activity in normal cells and cancer cells explains therapeutic

cytotoxicity on cancer cells while having minimal impact on normal cells <sup>[152]</sup>. Chronic lymphocytic leukemia (CLL) patients have been shown to exhibit spontaneous T cells recognizing GV1001, (a hTERT-peptide, see Section 4.4.1) which could lyse autologous telomerase expressing leukemic cells <sup>[153]</sup> cells. In a well-vetted priority-ranked list, composed to prioritize TAA for translational research based on predefined criteria, hTERT ranked no.23 of 75 <sup>[70]</sup>.

#### 4.4 Therapeutic cancer vaccines targeting hTERT

There are two general strategies of telomerase targeting in cancer treatment. One is targeting directly by inhibiting the activity of its catalytic subunit (hTERT) or its RNA template (hTER), leading to inhibition of telomerase activity, telomere shortening and inhibition of cell proliferation. Another strategy is to target the telomerase subunit indirectly via G-quadruplex stabilizers, tankyrase or HSP90 inhibitors, thus blocking telomerase access to telomeres or inhibiting binding of telomerase-associated proteins leading to telomere uncapping and cell apoptosis <sup>[154, 155]</sup>. The latter do not constitute immunotherapeutic modalities. Targeting of hTERT from an immunological standpoint will be discussed in this section.

Since telomerase is present in most cancers, its peptides are considered shared TAAs. They are however capable of producing strong immune response by eliciting both CD4+ and CD8+ T cell responses, potentially leading to tumor cell lysis <sup>[155-157]</sup>.

##### 4.4.1 GV1001- hTERT peptide-based vaccine

GV1001 is a 16 amino acid HLA Class II restricted hTERT-peptide, which consists of amino acids 611-626 of the hTERT active site <sup>[158-160]</sup> and is capable of binding to molecules encoded by multiple alleles of all three loci of HLA Class II <sup>[161]</sup>. These characteristics of the hTERT peptide might virtually enable all patients, irrespective of HLA-type, to present one or more immunogenic epitopes to effector cells. GV1001 is administered as an MHC Class II peptide, which is endogenously processed to yield a MHC Class I peptide producing both CD4+ and CD8+ responses. The activity of CD4+ T cells leads to a secretion of IFN- $\gamma$  and IL-2, further stimulating CD8+ CTL's and NK cells, which may help to increase the infiltration and the retention of CD8+ T cells into the tumor sites leading to an up regulation and re-expression of MHC Class I molecules. This may have a therapeutic advantage in patients with advanced cancers where loss of MHC Class I is seen <sup>[162, 163]</sup>.

Several strategies are employed in the development of vaccines that may induce hTERTs immunogenicity and eliminate the issue of self-tolerance, such as the use of adjuvans like GM-CSF or TLR-7 agonist (imiquimod), used in GV1001 vaccine strategies <sup>[160, 162]</sup>. This may prevent the rapid degradation and elimination of anticancer vaccine peptides before recognition by APCs, which may occur due to self-tolerance to self-peptides <sup>[160, 164]</sup>.

GV1001 has completed several phase I and II studies conducted in patients with advanced stage melanoma, hepatic cellular carcinoma and in patients with pancreatic cancer <sup>[149]</sup>. Forty-eight patients with unresectable pancreatic adenocarcinoma were given

i.d. injections of GV1001 at three dose-levels along with GM-CSF for 10 weeks followed by monthly boosters. Delayed-type hypersensitivity (DTH) and in vitro T cell proliferation was measured. Of the 27 evaluable patients, median survival for the intermediate group (300 nmole GV1001) was 8.6 months, which was significantly longer than the low and high-dose groups. One-year survival in the intermediate group was 25% [165].

Two other phase III studies with GV1001 in patients with nonresectable pancreas cancer were Primo Vax and Telo Vac. The Primo Vax trial examined vaccine monotherapy versus gemcitabine but was terminated due to lack of survival advantage. The Telo Vac had three arms; sequential gemcitabine/capecitabine; and concurrent gemcitabine/capecitabine and vaccine. The results are pending. Another study is evaluating radiation therapy, tadalafil, sargramostim, gemcitabine and GV1001 in patients with unresectable pancreatic cancer and is ongoing, Clinical Trials.gov identifier NCT01342224.

## 4.5 Immune monitoring

Technologies available for measuring immune responses are numerous. It is critical to select an assay that can accurately measure therapy-induced changes in the frequency and/or function of immune cells. Briefly, phenotypic markers, functional assessments or cellular product may be assayed [166]. Phenotypic assays include measurement of absolute cell numbers and cell frequencies by flow-cytometry and tetramer binding. Functional assays include DTH, proliferation, cytotoxicity assays (ADCC, CDC) and enzyme-linked immunosorbent spot (ELISPOT). Assays for cellular products include enzyme linked immunosorbent assay (ELISA) (immunoglobulin levels, cytokines) and chemokine/cytokine levels (Multiplex).

In patients treated with TCV, three single-cell assays for detection of antigen-specific effector T lymphocytes can be used for monitoring: ELISPOT, cytokine flow-cytometry (CFC) and tetramer binding.

Each of these is based on T cell receptor recognition of cognate peptides presented by MHC Class I/II molecules on the surface of APCs to the responder T cells. They have in many ways replaced “bulk assays” as cytotoxicity assays and proliferation assays. However, no consensus exists as to which of these three should be selected to best monitor vaccination results [166]. ELISPOT measures function of individual responder cells by identifying those that produce and secrete the measured cytokine, CFC measures cytokine expression in a cell and not its actual secretion [166]. Tetramer binding detects peptide-specific T cells expressing the relevant TCR but may also bind non-functional T cells [167]. However, CFC and tetramer binding are flow-cytometry based and can provide information about identification of responding cells.

Simultaneous assessment of multiple biomarkers, for example multiplex assay may provide the investigator with an “immune profile”. Multiplex assays have all but replaced ELISA detection for cytokines allowing simultaneous measurement of, for example, suppressive cytokines, proinflammatory cytokines and Th1 vs Th2 cytokines [166]. For further details see Table 6. Polychromatic flowcytometry enables further examination of cells. mAbs labelled with various chromophores with various excitation wavelengths is used to identify and determine levels of expression of surface or intracytoplasmic markers.

**Table 6.** Assays used for immune monitoring in vaccine clinical trials

| Assay                           | Brief description   | Advantages  | Disadvantages  |
|---------------------------------|---|---|--|
| 1 DTH                           | Ag as soluble protein is injected i.d. and the diameter of erythema or induration is measured after 48-72 h   | <ul style="list-style-type: none"> <li>• only <i>in vivo</i> measure available</li> <li>• easy to perform</li> </ul>  | <ul style="list-style-type: none"> <li>• no standardized cut-off for a positive response</li> <li>• no standardized dose for DTH testing</li> <li>• Ag-specificity is questionable</li> <li>• Mainly mediated by CD4+ effector memory T cells</li> </ul>   |
| 2 ELISPOT                       | Lymphocytes are cultured with the Ag studied in a micro-titer plate coated with a monoclonal Ab to a specific soluble factor (e.g. IFN- $\gamma$ , IL-4, IL-10, TNF- $\alpha$ ). The cells and the Ag are then washed from the wells and replaced with secondary antibody conjugated to a detection reagent. The plate is developed with a chromogen and spots appear where there was a cell secreting the soluble factor being investigated.                               | <ul style="list-style-type: none"> <li>• functional assay</li> <li>• allows to measure individual soluble factors secreted by activated T cells and identify the pathway of the immune system activated by the vaccine</li> <li>• Lowest limit of detection (1/100000 Ag-specific T cells)</li> <li>• considerably reliable</li> <li>• relatively rapid</li> </ul>  | <ul style="list-style-type: none"> <li>• provides no information on cell phenotype</li> <li>• Responder-stimulator interactions may result in unacceptable high background spots</li> </ul>  |
| 3 Tetramer staining             | Tetramers are composed of four MHC-I molecules, each bound to the epitope of interest. The tag is a fluorescent label, which allows to measure the binding of the tetramer to the TCR at flow-cytometry.  | <ul style="list-style-type: none"> <li>• sensitive (1/100000 Ag-specific T cells)</li> <li>• T cell subset analysis is optimal</li> <li>• allows to identify the peptide sequence or epitopes that bind to the highest number of TCR in a naïve individual</li> <li>• allows to identify the phenotype of the T cell to which the tetramer binds</li> <li>• allows to measure the change in the number of T cells displaying a particular TCR before and after vaccination</li> </ul> | <ul style="list-style-type: none"> <li>• requires knowledge of the epitope</li> <li>• requires availability of the tetramer for the respective epitope/HLA allele</li> <li>• unable to distinguish between functional and dysfunctional T cells</li> </ul>   |
| 4 Cytokine Flow Cytometry (CFC) | Lymphocytes are cultured with the Ag studied and the presence of intra-cellular cytokines is detected by fluorescein-labelled mAb. The phenotype of the lymphoid cells (CD4+, CD8+, etc) is identified with a second set of fluorescein-labelled mAbs.  | <ul style="list-style-type: none"> <li>• functional assay</li> <li>• sensitive (1/10000 Ag-specific T cells)</li> <li>• provides additional information on cell phenotypes</li> <li>• relatively rapid</li> </ul>   | <ul style="list-style-type: none"> <li>• non-specific background staining</li> </ul>   |
| 5 Lympho-proliferation assay    | Lymphocytes are cultured with the Ag studied. $^3\text{H}$ thymidine is added to the culture medium. Proliferating (dividing) cells incorporate $^3\text{H}$ thymidine, which is quantitated using a beta scintillation counter   | <ul style="list-style-type: none"> <li>• easy to perform</li> <li>• reliable</li> <li>• sensitive</li> <li>• reproducible</li> </ul>  | <ul style="list-style-type: none"> <li>• can be influenced by the non-specific immune function of the patient</li> <li>• can be influenced by the <i>in vitro</i> stimulation procedure</li> <li>• not qualitative</li> <li>• not quantitative</li> <li>• no information about responding cell population (CD4+, CD8+, etc)</li> </ul> |
| 6 Cytotoxicity assay            | Lymphocytes previously sensitized to the Ag present on the target cells are cocultured with the target cells. Percentage of lysis of target cells is quantitated by $^{51}\text{Cr}$ release assayed by Flow Cytometry  | <ul style="list-style-type: none"> <li>• functional assay</li> <li>• measures the ability of direct tumor lysis</li> </ul>  | <ul style="list-style-type: none"> <li>• low sensitivity</li> <li>• often involves multiple <i>in vitro</i> stimulations</li> <li>• not quantitative</li> <li>• often other targets than autologous tumor are used, which may not reflect the capability of effector cells to lyse autologous tumor cells <i>in vivo</i></li> </ul>    |
| 7 Multiplexing for cytokines    | A mixture of an infrared dye with a red dye is incorporated into polystyrene beads. The concentration ratio of both dyes is varied to produce beads which emit a well-defined fluorescence spectrum upon excitation. Each bead is identified by its unique colour, and can be coupled with a capture molecule specific for a ligand of interest, e.g. like a protein. The mixture of the beads is quantitatively analyzed in a FACS-like fluorescence-activated bead sorter | <ul style="list-style-type: none"> <li>• Used as cytokine profiling Th1/Th2</li> <li>• Small sample volume fluid</li> <li>• High throughput</li> </ul>  | <ul style="list-style-type: none"> <li>• Sera measurement may be less useful than <i>in situ</i> measurement</li> </ul>  |

Adapted from [Palma et al. 2007<sup>[168]</sup>]

## **4.6 Clinical endpoints**

The gold-standard for confirmatory trials is OS but evaluation of OS in clinical trials is a lengthy process, suggesting the need to identify surrogate endpoints for markers of efficacy. A number of surrogate end-points for OS exist including time to progression (TTP), PFS and disease-free survival (DFS), but such endpoints may not accurately predict the outcome of OS. This was exemplified in the study concerning Sipuleucel-T, previously mentioned above. The primary end-point of TTP was not met, but there was a trend towards improved OS in the treatment group <sup>[169]</sup>.

Furthermore, in order to show clinical efficacy, measurements of tumor progression is often made by using the Response Evaluation Criteria in Solid Tumors (RECIST), which defines the response to treatment according to size <sup>[170]</sup>. However, this may not always reflect the efficacy of immunotherapy. Anecdotal data suggest that increase in tumor size may be due to increased inflammatory infiltrates. For example, the administration of mAb targeting the immunosuppressive receptor CTLA4 has been shown to double the survival of Stage IV melanoma patients in the absence of early tumor shrinkage <sup>[120]</sup>.

## 5 COLORECTAL CANCER

### 5.1 Epidemiology

CRC is the third most common cancer in both men and women, and the fourth leading cause of cancer deaths in the world <sup>[171]</sup>. The estimated worldwide incidence of CRC is 1.2 million per year <sup>[171]</sup>. In Sweden approximately 6000 patients are diagnosed with a colorectal cancer every year, see Table 3 <sup>[172]</sup>.

**Table 3.** Colorectal cancer incidence (total number) in Sweden 2011

| Tumour site | Female | Male | Total |
|-------------|--------|------|-------|
| Colon       | 2102   | 2081 | 4183  |
| Rectum      | 821    | 1158 | 1979  |
| Total       | 2923   | 3239 | 6162  |

Mean age at diagnosis was 72 years for colon cancer and 70 years for rectal cancer in patients diagnosed during 2000-2007 <sup>[173]</sup>. A screening program for early detection commenced in 2008 in the Stockholm-Gotland region, and plans are to further expand this nationwide in late 2013. The vast majority of CRCs, about 95% constitute of adenocarcinomas.

### 5.2 Clinical staging

To select the optimal therapeutic strategy for a patient with CRC, a correct clinical and pathological staging is of great importance. Furthermore, the clinical and pathological staging at diagnosis is a crucial prognostic factor <sup>[174]</sup>.

Preoperative standard staging involves endoscopy of the rectum and colon, computer tomography (CT) of the chest and abdomen, and in rectal cancer, an additional magnetic resonance imaging (MRI) of the pelvis.

The diagnosis of an adenocarcinoma in the colon or rectum is made by biopsy of the tumor. In order to select the optimal treatment strategy for CRC patients, the results of the staging procedure are discussed at a multidisciplinary team (MPT) conference <sup>[175]</sup>.

#### 5.2.1 Tumor-Node-Metastasis staging

Collected data is condensed according to a staging system, based on the depth of extension of the carcinoma through the bowel wall (T), the presence or absence of lymph node metastases (N) and the presence or absence of distant metastasis (M). Historically, Duke's classification system has been used, but the TNM staging system by American Joint Committee on Cancer (AJCC) and International Union Against Cancer (UICC) is now recommended. Details on TNM staying system and comparison to Duke classification system are shown in Table 4. In the revision of the TNM system to the 7<sup>th</sup> edition, further subclassifications were added <sup>[176]</sup>. In our studies we have used the Dukes classification system in Paper I and AJCC TNM 5<sup>th</sup> Edition in Papers II and III. In our studies we used

Duke's classification system in Paper I and AJCC TNM 5<sup>th</sup> Edition in Papers II-III, see Table 4.

Table 4.

| <i>TNM Stage 7<sup>th</sup> Edition</i> |        |        |     | <i>TNM Stage 5<sup>th</sup> and 6<sup>th</sup> Edition</i> |       |       |    | <i>Duke's</i> |
|---|--------|--------|-----|--|-------|-------|----|---------------|
| <i>Stage</i>                            | T      | N      | M   | <i>Stage</i>   | T     | N     | M  |               |
| <b>0</b>                                | Tis    | N0     | M0  | <b>0</b>   | Tis   | N0    | M0 | –             |
| <b>I</b>                                | T1-2   | N0     | M0  | <b>I</b>   | T1-2  | N0    | M0 | A             |
| <b>IIA</b>                              | T3     | N0     | M0  | <b>IIA</b>   | T3    | N0    | M0 | B             |
| <b>IIB</b>                              | T4a    | N0     | M0  | <b>IIB</b>   | T4    | N0    | M0 | B             |
| <b>IIC</b>                              | T4b    | N0     | M0  |  |       |       |    | B             |
| <b>IIIA</b>                             | T1-2   | N1/N1c | M0  | <b>IIIA</b>  | T1-2  | N1    | M0 | C             |
|   | T1     | N2a    | M0  |  |       |       |    | C             |
| <b>IIIB</b>                             | T3-T4a | N1/N1c | M0  | <b>IIIB</b>  | T3-4  | N1    | M0 | C             |
|   | T2-T3  | N2a    | M0  |  |       |       |    | C             |
|   | T1-T2  | N2b    | M0  |  |       |       |    | C             |
| <b>IIIC</b>                             | T4a    | N2a    | M0  | <b>IIIC</b>  | Any T | N2    | M0 | C             |
|   | T3-T4a | N2b    | M0  |  |       |       |    | C             |
|   | T4b    | N1-N2  | M0  |  |       |       |    | C             |
| <b>IVA</b>                              | Any T  | Any N  | M1a | <b>IV</b>  | Any T | Any N | M1 | D             |
| <b>IVB</b>                              | Any T  | Any N  | M1b |  |       |       |    | D             |

#### Tumor-Node-Metastasis Classification

| <i>AJCC 7<sup>th</sup> Edition</i> |   | <i>AJCC 5<sup>th</sup> and 6<sup>th</sup> Edition</i>                                      |  |
|------------------------------------|---|--|--|
|                                    |   | <i>T (primary tumor)</i>   |  |
| <b>TX</b>                          | Primary tumor cannot be assessed  | Primary tumor cannot be assessed   |  |
| <b>T0</b>                          | No evidence of primary tumor  | No evidence of primary tumor   |  |
| <b>Tis</b>                         | Carcinoma in situ: intra epithelial or<br>Invasion of lamina propria  | Carcinoma in situ: intra epithelial or<br>Invasion of lamina propria                       |  |
| <b>T1</b>                          | Tumor invades submucosa   | Tumor invades submucosa  |  |
| <b>T2</b>                          | Tumor invades muscularis propria  | Tumor invades muscularis propria   |  |
| <b>T3</b>                          | Tumor invades through the muscularis propria into<br>pericorectal tissues   | Tumor invades through the muscularis propria into<br>pericorectal tissues                  |  |
| <b>T4</b>                          | Tumor directly invades other organs or structures<br>and/or perforates visceral peritoneum  | Tumor directly invades other organs or structures<br>and/or perforates visceral peritoneum |  |
| <b>T4a</b>                         | Tumor penetrates to the surface of the visceral<br>peritoneum   |  |  |
| <b>T4b</b>                         | Tumor directly invades or is adherent to other<br>organs or structures  |  |  |
| <i>N (Regional lymphnodes)</i>     |   |  |  |
| <b>NX</b>                          | Regional lymph nodes cannot be assessed   | Regional lymph nodes cannot be assessed  |  |
| <b>N0</b>                          | No regional lymph node metastases   | No regional lymph node metastases  |  |
| <b>N1</b>                          | Metastases in 1 to 3 regional lymph nodes   | Metastases in 1 to 3 regional lymph nodes  |  |
| <b>N1a</b>                         | Metastases in one regional lymph node   |  |  |
| <b>N1b</b>                         | Metastases in 2 to 3 regional lymph node  |  |  |
| <b>N1c</b>                         | Tumor deposit(s) in the subserosa, mesentery, or<br>non peritonealized pericolic or perirectal tissues<br>without regional nodal metastases |  |  |



|            |  |  |
|------------|--|--|
| <b>N2</b>  | Metastases in 4 or more regional lymph nodes | Metastases in 4 or more regional lymph nodes |
| <b>N2a</b> | Metastases in 4-6 regional lymph nodes       |  |
| <b>N2b</b> | Metastases in 7 or more regional lymph nodes |  |

**M (Distant metastases)**

|            |  |                                       |
|------------|--|---------------------------------------|
| <b>MX</b>  | Distant metastases cannot be assessed  | Distant metastases cannot be assessed |
| <b>M0</b>  | No distant metastases  | No distant metastases                 |
| <b>M1</b>  | Distant metastases   | Distant metastases                    |
| <b>M1a</b> | Metastases combined to one organ or site (for example lung, liver, ovary, non-regional node) |                                       |
| <b>M1b</b> | Metastases in more than one organ/site or the peritoneum                                     |                                       |

The TNM system defines the stage, see Table 4, and the 5-year survival for patients with CRC is directly related to the clinical and pathological staging at the time of diagnosis. It varies between 95-100% for stage I, 75-90% for stage II, 45-60% for stage III and about 10% for stage IV, reported in a recently published Swedish follow-up study<sup>[177, 178]</sup>.

## 5.3 Treatment

### 5.3.1 Surgery

Surgery is the primary treatment in CRC and the goal is to remove the tumor together with its regional lymphatic drainage and blood supply. The total mesorectal excision (TME) approach for rectal cancer, described in 1982, has reduced the rates of local recurrence considerably<sup>[179]</sup>.

Patients in the palliative patients are offered surgery of the primary tumor if complications such as bleeding, perforation or obstruction may or do occur.

### 5.3.2 Non-surgical treatment of CRC

#### Adjuvant treatment/Colon cancer

The aim of adjuvant treatment is to prevent local recurrence, distant metastases and to prolong survival. Postoperative chemotherapy has been advocated for colon cancer in Sweden since the mid nineties. Currently, 6 months of adjuvant chemotherapy is recommended, although ongoing studies are investigating shorter periods of treatment as well as administering part of the chemotherapy preoperatively. Treatment might commence within 8 weeks after surgery<sup>[180-182]</sup>.

Controversy still exists regarding the role of standard adjuvant chemotherapy for **stage II colon cancer** disease<sup>[183]</sup>. Factors associated with a higher risk of recurrence include T4 stage, perforation at presentation and inadequate node sampling (less than 12 nodes). Tumor grade and lymphovascular invasion are more controversial in this regard. All these factors are associated with a higher risk of recurrence but they are not predictive of a benefit from adjuvant chemotherapy, (i.e they are not prognostic markers). The TNM staging system has its limitations. For example, stage II patients who display mismatch repair deficiency (microsatellite instability, which is not included in the TNM system),

have an excellent prognosis following radical surgery and do not appear to benefit from adjuvant 5-fluorouracil (5-FU) <sup>[184]</sup>. Regarding the use of oxaliplatin in stage II patients, there is no evidence for the addition of this to 5-FU <sup>[185]</sup>. Stage II patients need to be considered carefully for chemotherapy based on all available clinical individual data.

In *stage III colon* cancer, combination of oxaliplatin and 5-FU is recommended for most patients. However, the benefits of oxaliplatin are less clear when the risk of recurrence is low (e.g. low grade, T1/2 or N1 disease) or the treatment related risks are high, such as those with comorbidity, elderly or poor performance status <sup>[185-187]</sup> and capecitabine monotherapy is a reasonable alternative in these situations <sup>[188]</sup>. Irinotecan, widely used in metastatic colorectal cancer (mCRC), has failed to improve survival in the adjuvant setting <sup>[189, 190]</sup>.

Likewise, biological agents have been disappointing and are not routinely used in the adjuvant setting. The addition of the antiangiogenic monoclonal antibody targeting VEGF, Bev has not improved 3-year disease free survival (DFS) <sup>[191]</sup>.

Adding the anti-EGFR-targeted monoclonal antibody Cet to adjuvant chemotherapy has also proven disappointing in both Kirsten rat sarcoma viral oncogene (KRAS) wild-type and KRAS mutant tumors <sup>[192]</sup>, thus the identification of a predictive/prognostic biomarker has not yet affected adjuvant therapy selection as it has for mCRC.

### 5.3.3 Curative treatment of rectal cancer

Neoadjuvant therapy in rectal cancer often involves radiation therapy. The introduction of more precise surgery, TME and the use of pre or postoperative radiation therapy or radiochemotherapy have contributed both to much lower local recurrence rates and improved survival and more accurate preoperative staging with MRI and a series of trials have established radiotherapy as standard care <sup>[193, 194]</sup>. Radiation therapy, alone or with chemotherapy is given not only to the 10-15% non-resectable but also to much less advanced tumors, and in Sweden the standard is to give RT before surgery.

Two different preoperative approaches are mainly used. The short-course consists of 25 Gy in 5 Gy fractions given during 5 days, followed by surgery the following week. The long-course consist of 45-50.4 Gy given in 1.8-2.0 Gy fractions concomitant with 5-FU, either given as bolus or as capecitabine <sup>[193, 195, 196]</sup>.

Patients are grouped according to their MRI-staging results and considered for radiation therapy. Tumors labelled as “good” receive no radiation therapy before surgery, tumors labelled as “bad” receive short-course radiation therapy and tumors labelled as “ugly” receive long-course radiation therapy, see Table 5 <sup>[197]</sup>.

**Table 5.**

|                   | <b>Good</b>    | <b>Bad</b>      | <b>Ugly</b>         |
|-------------------|----------------|-----------------|---------------------|
| <b>Low tumor</b>  | T1-2, N0,mrf-  | T3,N1-2,mrf-    | T4,T3 mrf+          |
| <b>High tumor</b> | T1-T3b,N0,mrf- | T3c-d,N1-2,mrf- | Lateral lymphnodes+ |

Cut-off between low/high tumors is at app. 8 cm from the anal verge  
mrf; Mesorectal fascia  
mrf+ Tumors growing adjacent to the mesorectal fascia  
mrf- Tumors not growing adjacent to the mesorectal fascia  
Extramural vessel invasion (not mentioned above) is labelled as at least “bad”

Adapted from Blomqvist, Glimelius. Acta Oncologica, 2008; 47:5-8.

Opinions on adjuvant chemotherapy vary a great deal in the literature. Some authors advocate that chemotherapy ought to be given applying the same criteria as in colon cancer, whereas some advocate that although surgery and preoperative (chemo)radiotherapy was adequately performed – the value of adjuvant chemotherapy has not been established <sup>[198]</sup>. Swedish National Guidelines suggest the following recommendations:

- Adjuvant chemotherapy is recommended to patients with highrisk-criteria rectal cancer stage II and stage III patients to whom preoperative radiation therapy was not given.
- Adjuvant chemotherapy may be considered in patients with highrisk-criteria stage II and stage III patients who received short-course radiation therapy and surgery the following week.
- Adjuvant chemotherapy may be considered in patients with highrisk-criteria stage II and stage III patients operated 4-6 weeks after preoperative short-course radiation therapy.

The uncertainty is even greater regarding patients who receive long-course radiation therapy and 5-FU preoperatively. The addition of oxaliplatin in the adjuvant setting has not been evaluated. A small retrospective study compared adjuvant 5-FU-based chemotherapy with no treatment in patients with T3-4 rectal cancers <sup>[199]</sup>. The results showed no significant impact on adjuvant chemotherapy on PFS or OS, although a difference seemed to emerge at approximately, respectively, 2 and 5 years after start of preoperative treatment. Patients in whom no down staging was seen did not benefit from adjuvant chemotherapy, indicating that the same prognostic factors may drive both tumor sensitivity for the primary treatment and long-term benefit from further adjuvant chemotherapy.

An alternative approach is to administer the CT preoperative, as is currently evaluated in the Rapido trial, Clinicaltrial.gov identifier NCT01558921. Primarily “ugly” tumors are included. To guarantee control of the rectal tumor short course radiation therapy is given followed by systemic chemotherapy (capecitabine and oxaliplatin) before surgery. The investigators hypothesize that with this protocol both the local tumour and possible micro metastases are effectively treated and that this will result in an increased survival. The investigators will compare this with the standard treatment of long-course radiation therapy

followed by TME surgery and adjuvant chemotherapy (capecitabine and oxaliplatin). I.e all patients in the standard arm receive adjuvant chemotherapy.

### 5.3.4 Metastatic CRC

In selecting the optimal treatment for mCRC, the initial consideration is whether the aim is curative (initially or potentially resectable disease) or noncurative.

At the time of diagnosis approximately 15% of the patients with stage IV CRC have metastatic disease in the liver and less than 10% have pulmonary metastases that are resectable [200, 201]. The majority of patients with mCRC and liver metastases are considered unresectable at presentation, although a minority of these patients may eventually be able to be treated with curative intent if a good response to chemotherapy is achieved. Complete resection of limited liver metastases and/or lung metastases is the only potentially curative treatment for mCRC with a survival rate after hepatectomy of 40-60% if patients are carefully selected [202, 203].

Affecting the choice of treatment are factors relating to the patient (age, performance status, comorbidities, patients preferences) and those relating to the tumor (potential resectability, disease burden, symptoms, rate of progression, prognostic biomarkers and prior treatment history).

Doublet chemotherapy is recommended if an aggressive approach is indicated, i.e to stabilize rapidly progressive disease, in order to reduce metastasis where surgery is possible and in first-line resectable patients. 5-FU is the backbone in doublet chemotherapy. There are little differences in efficacy between doublets of 5-FU-infusion/capecitabine and oxaliplatin (FOLFOX and XELOX) or 5-FU-infusion/capecitabine and irinotecan (FOLFIRI and CAPIRI) regimens [174]. Single chemotherapy (5-FU) is recommended in non-rapidly progressive nonresectable disease.

Patient outcome is further improved by the addition of a biological agent to these chemotherapy regimens, why this may be added to the treatment of potentially resectable mCRC patients and non-resectable patients.

The addition of Bev to both irinotecan and oxaliplatin based doublet chemotherapy significantly improves PFS and with irinotecan, OS in mCRC patients, but with smaller effects on resection rates [204, 205].

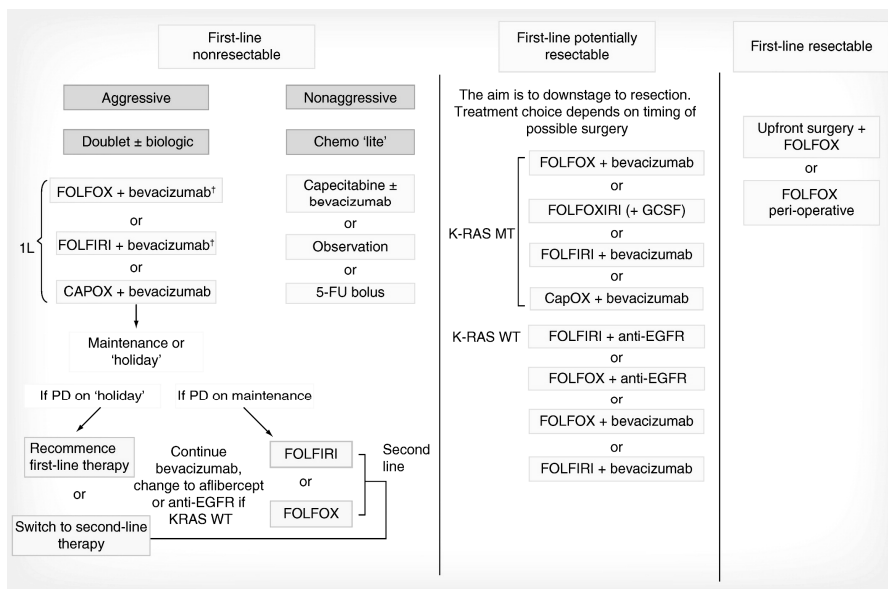
The benefit of EGFR-antibody (Cet or panitumumab (Pan)) treatment to KRAS wildtype (WT) tumors are highly variable in trials. Unlike antiangiogenics, a predictive marker exists for EGFR-targeted therapy. KRAS gene mutation (MT) is predictive of nonresponse to EGFR-targeted therapy either as a single agent or in combination with irinotecan and oxaliplatin based chemotherapy and in standard care EGFR-antibodies are only recommended to KRAS WT patients. KRAS mutations in codon 12 are most common, but mutations in codon 13 constitute approximately 15% of all KRAS mutations [174]. Conflicting data exists on whether KRAS MT in codon 13 are eligible for treatment with EGFR-targeted therapy [174]. Furthermore, CRC tumors, defined as KRAS WT may harbour other oncogenic alterations (BRAF, NRAS) in which the benefit of EGFR-targeted therapy is yet unknown [174]. However, large Phase III studies have shown OS benefit provided by Cet and PFS benefit for both Cet and Pan in KRAS WT patients. EGFR-

targeted therapy with Cet or Pan also generally improves RR in KRAS WT patients by over 10% when added to doublet chemotherapy [206, 207].

The use of triplet chemotherapy, 5-FU/irinotecan and oxaliplatin, (FOLFOXIRI) for first line mCRC has been investigated in Phase III studies with an unclear survival benefit compared with double chemotherapy. FOLFOXIRI significantly increased resection rate, complete resection rate after surgery, PFS and OS compared [208] to FOLFIRI in one study, while in another study, FOLFOXIRI was not superior to FOLFIRI [209]. Toxicity is a major concern and FOLFOXIRI is therefore not part of standard treatment, but may be considered in selected patients.

### 5.3.5 Second-line treatment and beyond

For most patients, progression following an oxaliplatin doublet will result in a switch to irinotecan-based therapy and vice versa. PFS is shorter than with first-line therapy [210]. If single chemotherapy was given, registry data suggest oxaliplatin based doublet chemotherapy [211]. KRAS WT patients, not previously exposed to Cet or Pan, may receive this as third-line therapy alone or in combination with irinotecan. The choice of biological agent in second line therapy is an area of active research with ongoing Phase III studies. Afibercept in second-line treatment combined with FOLFIRI and regorafenib in late-stage disease has shown survival gains in recently reported trials and may become part of standard treatment [174]. Summary of proposed management for mCRC is seen in Figure 4.



**Figure 4.** Proposed chemotherapy management algorithm for metastatic colorectal cancer. †If aim is downstaging or bulky and symptomatic disease is present in KRAS WT patients, anti-EGFR should also be considered. EGFR: EGF receptor; MT: Mutation; PD: Progressive disease; WT: Wild-type.

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## 6 PANCREATIC CANCER

### 6.1 Epidemiology and prognosis

Pancreatic cancer is currently one of the deadliest of the solid malignancies. The annual incidence in Sweden is approximately 11 cases per 100 000 inhabitants. In Sweden, 1580 people died due to pancreatic cancer in 2011 <sup>[172]</sup>. Approximately 85% of the patients have advanced spread disease at the time of diagnosis and only 15-20% of the patients are eligible for curative surgery. Following curative surgery, 90 % of the patients will relapse. Despite decades of effort, the 5-year survival rate remains at only 5% and the median survival reported for resected pancreatic cancer patients ranges from 17-27 months <sup>[212]</sup>.

In the sections that follow, “pancreatic cancer” (PC) will refer to invasive ductal adenocarcinoma of the pancreas.

### 6.2 Risk factors

Both environmental and inherited factors contribute to the development of PC. The most common risk-factor is smoking. Studies has shown that current smokers have a 2.2 fold increased risk of PC than never smokers and approximately 25% of PC are attributable to smoking <sup>[213, 214]</sup>. Long standing type 2 diabetes of >10 years duration have a 1, 5 foldincrease compared with nondiabetics <sup>[215]</sup>. Increased body mass index (BMI) is an independent risk factor <sup>[216]</sup>. Heavy alcohol consumption and chronic pancreatitis also elevates the risk of PC, the latter 2.7-fold. Individuals with a family history of PC have a 1.9-13-fold increased risk, based on case-control and cohort studies <sup>[212]</sup>. Inherited (germline) mutations in the BRCA2 gene, PALB2 and Lynch syndrome are all associated with higher risks <sup>[212]</sup>.

### 6.3 Staging

A contrast-enhanced thin-slice CT scan of the thorax, adomen and pelvis is performed in the staging procedure. Optimal treatment depends on careful staging and all patients should be discussed in a multidisciplinary team. In the absence of metastatic disease, the relationship of the tumor to the adjacent major vessels defines resectability. Resectable stages include stage I and II, and the subset of stage III that is defined as borderline resectable. Less than 180-degree involvement of the celiac axis or superior mesenteric artery is considered stage III borderline resectable. Greater than 180-degree involvement, is considered locally advanced or unresectable PC <sup>[217, 218]</sup>. AJCC/UICC and TNM definitions for pancreatic exocrine cancer are shown in Table 6.

**Table 6.** AJCC/UICC stage groupings and TNM definitions for pancreatic exocrine cancer

| AJCC/UICC stage | TNM                              |
|-----------------|----------------------------------|
| 0               | Tis,N0,M0                        |
| IA              | T1,N0,M0                         |
| IB              | T2,N0,M0                         |
| IIA             | T3,N0,M0                         |
| IIB             | T1,N1,M0<br>T2,N1,M0<br>T3,N1,M0 |
| III             | T4,any N,M0                      |
| IV              | Any T, Any N, M1                 |

TNM Classification for Exocrine pancreatic cancer. Adapted from AJCC Cancer Staging manual. 7<sup>th</sup> ed, NY: Springer, 2010, pp 241-9

#### Primary tumor (T)

|     |  |
|-----|--|
| TX  | Primary tumor cannot be assessed   |
| T0  | No evidence of primary tumor   |
| Tis | Carcinoma in situ  |
| T1  | Tumor limited to the pancreas, less than 2 cm in greatest dimension  |
| T2  | Tumor limited to the pancreas, more than 2 cm in greatest dimension  |
| T3  | Tumor extends beyond pancreas but without involvement of the celiac axis or the superior mesenteric artery |
| T4  | Tumor involves the celiac axis or the superior mesenteric artery   |

#### Regional lymph nodes (N)

|    |   |
|----|---|
| NX | Regional lymph nodes cannot be assessed |
| N0 | No regional lymph node metastasis       |
| N1 | Regional lymph node metastasis          |

#### Distant metastasis (M)

|    |                       |
|----|-----------------------|
| M0 | No distant metastasis |
| M1 | Distant metastasis    |

## 6.4 Treatment

### 6.4.1 Surgery

Patients with Stage I/II disease are recommended immediate surgical resection followed by adjuvant chemotherapy. The low chance of long-time survival after surgery, mentioned above, and considerable morbidity in 40-60% of the patients following major surgery needs to be considered when selecting patients <sup>[212]</sup>. Complete recovery can take 2-3 months even in the absence of complications.

### 6.4.2 Neoadjuvant therapy

The advantage with neoadjuvant chemotherapy is 1; down-staging of some locally advanced tumors, (conversion rates vary between 15-40%) <sup>[219]</sup>, 2; it spares the 15-35% of patients who develop metastatic disease during treatment the risks of a major operation <sup>[212]</sup> and 3; guarantees that almost all patients receive some form of chemotherapy or radiation

therapy as postoperative complications may prevent postoperative treatment. In resectable patients neoadjuvant chemotherapy delays the curative surgery and systemic chemotherapy is therefore given postoperatively.

Stage III borderline patients should be recommended neoadjuvant therapy prior to surgery. Patients with stage III locally advanced disease should be treated with chemotherapy or chemoradiotherapy – a few of these patients may be considered for surgery.

Similar neoadjuvant regimens are often used for locally advanced and unresectable disease, i.e. Stage III with greater vessel involvement. The combination of 5-FU, LV, irinotecan and oxaliplatin (FOLFIRINOX), the combination of gemcitabine, docetaxel and capecitabine and gemcitabine alone are used, typically followed 5-FU infusion, capecitabine or gemcitabinebased chemoradiation to 45-54 Gy in 1,8 to 2,5 Gy fractions or 36 Gy in 2,4 Gy fractions <sup>[212]</sup>. Surgery is performed within 6-8 weeks following completion of neoadjuvant therapy. Neoadjuvant chemotherapy to patients with Stage I/II is controversial <sup>[212]</sup>.

### 6.4.3 Adjuvant therapy

Six months of adjuvant chemotherapy with either 5-FU or gemcitabine is recommended <sup>[212]</sup>. When gemcitabine was compared with observation in PC patients, DFS and OS were 6,9 months and 20,5 months for the observation arm, and 13,4 months and 24,2 months for the treatment arm <sup>[220]</sup>. A randomized trial between adjuvant elongated bolus 5-FU or gemcitabine showed similar OS of 23 months, with higher but acceptable rates of diarrhea in the 5-FU group and haematological toxicity in the gemcitabine group <sup>[221]</sup>. Chemoradiation adjuvant to the tumor bed is being explored, but has not reached standard practice.

Other adjuvant regimens are being evaluated. The combination of interferon-alfa, cisplatin and 5-FU with adjuvant chemoradiation had a median OS of 32 months compared to 28,5 months in the treatment arm with 5-FU alone <sup>[212]</sup>. Erlotinib has been studied and was safe to combine with gemcitabine as well as with concomitant capecitabine and chemoradiation adjuvant <sup>[212]</sup>.

In a phase II setting immunotherapy was combined with a pancreatic cancer vaccine of irradiated GM-CSF transfected allogenic whole-cell tumor lines. The median OS was 24, 8 months and patients who demonstrated a CD8+ T cell response to mesothelin had a higher likelihood of remaining disease-free <sup>[222]</sup>.

### 6.4.4 Metastatic disease

Since 1997, gemcitabine has been the standard treatment in the metastatic patient and hence tested with various other agents in doublet chemotherapy regimens <sup>[223]</sup>. Capecitabine coupled with gemcitabine has not shown superiority over gemcitabine alone, but may be considered in frail patients not eligible for single gemcitabine <sup>[224]</sup>. Nor has gemcitabine plus oxaliplatin or cisplatin shown significantly improved OS in phase III trials <sup>[212]</sup>. Adding docetaxel and capecitabine to gemcitabine has albeight in small studies, demonstrated good disease control rates and may be considered for patients with good performance status <sup>[225]</sup>.



Gemcitabine has also been tested with targeted therapy, the combination of gemcitabine and erlotinib improved OS minimally<sup>[226]</sup>. The addition of Cet to gemcitabine, or of Bev both demonstrated no survival benefit and added toxicity<sup>[227,228]</sup>.

FOLFIRINOX has emerged as a new promising regimen for fit patients in comparison to gemcitabine. Disease control rate and median OS were 70% and 11.1 months in the FOLFIRINOX arm, compared to 51% and 6, 8 months in the gemcitabine arm with intact quality of life<sup>[229]</sup>.

As for second line, patients treated with gemcitabine should be considered for a 5-FU based treatment, either alone or coupled with oxaliplatin or irinotecan<sup>[212]</sup>.

## **7 AIMS**

The overall aim of this thesis was to investigate immunogenicity of rCEA-protein, CEA-DNA and hTERT-peptide vaccination.

- To analyse the induction of IgM, IgA and IgE anti-CEA response in CRC patients without macroscopic disease, immunized with rCEA ± GM-CSF.
- To determine the safety of CEA66-DNA vaccination delivered by needle-free injection in combination with GM-CSF in CRC patients.
- To investigate the cellular and humoral immune response following CEA66-DNA vaccination delivered by Biojector or tetwtCEA-DNA-vaccination followed by electroporation as adjuvant treatment in CRC patients.
- To analyse the cellular and humoral immune response after priming with CEA66-DNA vaccination and boosting with tetwtCEA-DNA vaccination in CRC patients.
- To explore the immunogenicity and safety of GV1001 vaccination combined with GM-CSF and gemcitabine as first-line treatment in patients with advanced pancreatic adenocarcinoma.

## 8 MATERIAL AND METHODS

### 8.1 Patients

All patients included in this thesis were recruited and treated at the Department of Gastrointestinal Cancer at the Karolinska University Hospital, Sweden.

#### 8.1.1 Healthy donors

Healthy donors were recruited from the staff at the laboratory at Cancer Centre Karolinska (CCK). In Paper I, four healthy donors were analysed in the flow cytometry, the CDC and the ADCC analysis. No healthy donors were included in Paper II.

In Paper III and IV, nine healthy donors respectively were analysed in the proliferation assay, ELISPOT assay and flow-cytometry.

### 8.2 Methods

#### 8.2.1 Immune assays

Blood sampling was performed in the clinic and handled in a standardized fashion. All assays were carried out at the adjacent CCK laboratory – except for ELISA in Paper I and III. In Paper I, ELISA was performed at the Department of Clinical Chemistry at Ängelholm Hospital and in Paper III ELISA was performed at the Swedish Institute for Infectious Control. Used immune assays are displayed in Table 7.

Table 7

| Assays used | Prolif. | ELISPOT (INF- $\gamma$ ) | Cytokine secretion | DTH | ELISA | Flowcytometry | CDC | ADCC |
|-------------|---------|--------------------------|--------------------|-----|-------|---------------|-----|------|
| Paper I     |         |                          |                    |     | X     | X*            | X*  | X*   |
| Paper III   | X       | X                        | X                  |     | X     | X**           |     |      |
| Paper IV    | X       | X                        | X                  | X   |       | X**           |     |      |

\*In one patient only. \*\*T cell subsets, NK cells, NKT cells and MDSC.

#### 8.2.2 Proliferation assay

Peripheral blood mononuclear cell (PBMC),  $1 \times 10^5$ /well, were cultured with the rCEA protein (Paper I), or GV1001, the ras-peptide and the HIV-peptide (Paper IV), purified protein derivative of tuberculin (PPD) (Papers III, IV), phytohemagglutinin (PHA) (Papers III, IV) and tetanus toxoid (TT) (Paper III) in 96-well culture plates for 6 days. During the last 18 hours of incubation,  $1 \mu\text{Ci/well}$  [ $^3\text{H}$ ] thymidine was added. Cells were harvested and incorporated radioactivity was measured by a beta-counter. Results are shown as a stimulation index (SI) calculated by dividing mean radioactivity (cpm) of 6 replicates of experimental wells by that of the background value (cells with medium alone). Cells

stimulated with PPD, PHA, TT served as positive controls. Cells stimulated with HIV served as a negative control.

A change in ratio  $\geq 2$  on at least two occasions, compared to baseline, was considered significant [230].

### 8.2.3 ELISPOT (IFN- $\gamma$ )

PBMCs were cultured in 48-well plates with rCEA (Paper III), a pool of CEA peptides (Paper III), PHA and PPD and TT (Papers III, IV) for 5 days. A millipore 96-well filter plate was coated with anti-IFN- $\gamma$  antibody. Cultured PBMC were transferred to the coated plate and incubated for 20 h with the antigens as above. Cells were washed and incubated with a secondary biotinylated anti-IFN- $\gamma$  antibody for 2 hours at room temperature. After washing, Streptavidin-ALP conjugate was added to the cells and incubated for 1 hour at room temperature. Cells secreting IFN- $\gamma$  were developed by adding substrate BCIP/NBT plus and incubated at room temperature for 5 min. The reaction was stopped at the appearance of dark spots. Spots were counted by an automatic ELISPOT assay reader.

Paper III: A vaccine induced IFN- $\gamma$  response in ELISPOT was defined as an increase in delta value (experimental value minus background) at least two-fold compared to baseline [231].

Paper IV: A vaccine induced IFN- $\gamma$  response was defined if all of the following criteria were fulfilled: i) Spotforming units (SFU) of stimulated wells significantly higher ( $p < 0.05$ ) than of unstimulated wells (background) and at least twice that of the background, ii) SFU of cells stimulated with the control peptide not significantly ( $p > 0.05$ ) higher than background; iii) SFU of a post-vaccination test at least twice that of the pre-vaccination test [230]. The non-parametric Mann Whitney two-tailed rank sum test was used for comparison of unstimulated vs stimulated cells in Paper IV.

### 8.2.4 Cytokine secretion assay

Supernatants were collected (20  $\mu$ l/well) after 24 and 120 hours of incubation from the proliferation assay and stored at  $-70^{\circ}\text{C}$  until analyzed. The volume was replaced with complete medium. IL-4, IL-10, IFN- $\gamma$ , TNF- $\alpha$  and GM-CSF were analyzed (Paper III, IV). Standard concentration curves were generated. The coefficient of variation (mean  $\pm$  SD of the individual analytes) of PHA stimulated cells was calculated.

Cytokines concentration (pg/ml) in supernatants of antigen stimulated cells divided by that of cells alone using the highest value at 24 or 120 hours culture periods respectively was used. Post vaccination ratios divided by prevaccination ratios at different time points are shown. A ratio  $\geq 1, 5$  (Paper III) and  $\geq 2$  (Paper IV) (relative increase) was considered an antigen induced specific response [232].

### 8.2.5 DTH

GV1001 (0.112mg) in 0.1ml saline (groups A/B) was injected intradermally in the volar part of the forearm. GV1001 (0.105mg) in 0.22 ml saline (group C) was injected intradermally in the lower abdominal wall. The skin test was read, by the patient, after 48 hours by measuring the diameter of induration (mm). A positive DTH response was defined as  $\geq 5 \times 5$  mm of induration (Paper IV).

### 8.2.6 ELISA

The levels of IgM, IgA and IgE antibodies against rCEA were assayed by conventional isotype-specific sandwich ELISA (Paper I) <sup>[233]</sup>. Wells were coated with recombinant human CEA (2 ug/ml) and serum samples were assayed at 1:500 (IgM), 1:200 (IgA) and 1:10 (IgE) dilutions. Following overnight incubation in the wells antibodies were detected with alkaline phosphatase (ALP)-conjugated rabbit polyclonal isotype-specific F(ab')<sub>2</sub>-fragments (IgM and IgA) and biotinylated anti-IgE mAb/ALP-conjugated streptavidin (19) after 2h/2h incubations, respectively.

The antibody levels were calculated from standard curves established with chimeric IgM, IgA2 and IgE anti-NIP (5-iodo-4-hydroxy-3-nitro-phenacetyl acid) hapten antibodies <sup>[233, 234]</sup> using BSA-NIP conjugate (10 ug/ml) as coating antigen. This procedure permits comparison of results between runs in a semi-quantitative way where 1 AU (arbitrary ELISA unit) approximates to 1 ng of antibody as previously described <sup>[233]</sup>. The sensitivities of the IgA, IgM and IgE antibody assays in serum are 0.1 ug/ml, 2 ug/ml and 0.4 ng/ml, respectively.

In Paper III, conventional isotype-specific sandwich ELISA was also performed. Plates were coated with 0.1 µg per well of rCEA or purified hCEA diluted in 0.05 M Na<sub>2</sub>CO<sub>3</sub> (pH 9, 6) and incubated at room temperature overnight followed by 24 h in 4°C. After washing in ELISA buffer (0.05% Tween20, 0.15 M NaCl in distilled water), the plates were blocked by 5% milk in PBS for 2 h. Sera from immunized patients were diluted in 2.5% milk in PBS, and added to the plates (1:4 – 1:10000). Following incubation over night at room temperature, excess serum was removed and the plates were washed with ELISA buffer. To detect CEA-specific IgG antibodies, a rabbit anti-human HRP conjugate (1:3000 in 1.25% milk) was added to the plates and incubated at 37°C for 2 h. Plates were washed with ELISA buffer and developed by adding O-phenylene diamine substrate activated with H<sub>2</sub>O<sub>2</sub>. The reaction was stopped by adding 2.5 M H<sub>2</sub>SO<sub>4</sub>. Optical density (OD) was read at 490 and 650 nm. A net absorbance of 0.5 was used as a cutoff. A monkey anti-CEA serum served as a positive control.

A patient was considered to have developed a significant response if IgM, IgA or IgE (Paper I) and IgG (Paper III) antibodies could be detected at at least two different time points and was at least twice that of the pre-immunization value for that patient.

### 8.2.7 Flowcytometry

#### *Indirect immunofluorescence using flow cytometry*

In paper I, SW48 (a human cancer cell line expressing CEA) and A549 (a human lung cancer cell line not expressing CEA) were used for surface staining of a patient serum, pooled sera from healthy donors (HD) and a non-relevant IgA isotype control (22). Briefly, 10<sup>6</sup> cells were washed in PBS containing 0.01% sodium azide and incubated with the IgA isotype control, HD and the patient serum (1:10–1:1000) at 4°C for 1 h followed by 3 washings. A polyclonal rabbit anti human IgA serum (1:100) in washing buffer was then added and incubated at 4°C for 1 h. The cells were finally washed 3 times and fixed by adding 1% paraformaldehyde in PBS. Cells were analyzed by flow cytometry.

### **Cellular staining and flow cytometry**

Briefly conjugated antibodies were added to cells ( $5 \times 10^5$  cells per tube) and incubated for 30 min. Intracellular staining was performed using a T-reg staining kit. After a final wash, cells were resuspended in PBS and events acquired using LSRII and analyzed by the FlowJo software (Paper III, IV).

### **8.2.8 Complement dependent cytotoxicity (CDC)**

Stored sera ( $-70^\circ\text{C}$ ) from a patient who had developed high IgA anti-CEA antibody titers were used (Paper I). Two sampling times were pooled to obtain a sufficient amount of serum. IgA from the patient and pooled sera from healthy donors were isolated using peptide M/Agarose affinity column chromatography yielding highly purified IgA.

Briefly, as previously described [235], SW48 (a human cancer cell line expressing CEA) and A549 cells (a human lung cancer cell line not expressing CEA) ( $5 \times 10^4$ ) were plated in V-bottomed microtiter plates in 100  $\mu\text{l}$  RPMI-1640 containing 10% FCS (complete medium). Cells were incubated with 20  $\mu\text{g/ml}$  of purified IgA, from the patient and HD, respectively for 30 min at room temperature followed by twice washing with RPMI1640. After washing, 20% normal human serum in complete medium was added to the cells and incubated at  $37^\circ\text{C}$  in humidified air with 5%  $\text{CO}_2$  for 1 h. Finally, cells were collected, washed twice with 1xPBS and resuspended in 100  $\mu\text{l}$  of 1x binding buffer. Five  $\mu\text{l}$  of propidium iodine was added to the cells, vortexed and incubated at room temperature in the dark for 15 minutes. 100  $\mu\text{l}$  of 1x binding buffer was added to the cells which were analyzed by flow cytometry.

Results are presented as percent lysis (mean  $\pm$  SEM), in four different groups; 1; cells alone, 2; cells incubated with human complement, 3; cells plus isolated IgA from the immunized patient + human complement and 4; cells plus isolated IgA from HD  $\pm$  human complement. Statistical calculations were done comparing groups by unpaired t-test.

### **8.2.9 Antibody dependent cellular cytotoxicity (ADCC)**

ADCC was performed as previously described [236]. In brief, cell lines (same as used in CDC) were labeled with 2.8 MBq sodium  $^{51}\text{Cr}$  for 2 h. After 3 washings with DMEM medium 10000 cells in 100  $\mu\text{l}$  medium were added to each round-bottomed microtiter well (Nunc) and Ficoll-isopaque isolated PBMC of healthy donors to yield target:effector cell ratios of 1:25 and 1:50 to a final volume of 200  $\mu\text{l}$ . The test was run in sixuplicate. After 24 h at  $37^\circ\text{C}$  the reaction was stopped by centrifugation.  $^{51}\text{Cr}$  release was measured.

The percentage of target cell lysis was calculated according to the following formula: % specific lysis = (experiment cpm-spontaneous cpm) / (maximal cpm-spontaneous cpm) x 100. Maximal  $^{51}\text{Cr}$  release was determined by adding RIPA lysis buffer (150 mM NaCl, 25 mM Tris PH=7, 5 mM EDTA, 0.5 % SDS and 1 % Triton X-100) to target cells and spontaneous release was measured in the absence of antibodies and effector cells (Paper I). In each cell line, results were compared between three groups; 1; Isolated IgA from immunized patient, 2; Isolated IgA from HD and 3; no antibody. Statistical calculations were done comparing groups by unpaired t-test.

### 8.3 Follow-up

#### Paper I

The first patient was included in Dec 1994 and the last in April 1996. Patients were contacted in July 2007 and their medical history was taken. Data was cross-checked with patient's journals. Data on deceased patients were taken from patients journals. All patients were accounted for. Patients were monitored immunologically for 36 months and clinically for 154 months.

#### Paper and II and III

Systemic medical events occurring after the injection were assessed by the investigators and considered as product-related events only if their relation to the product was judged possible, probable or definite. AE were graded as to their seriousness, severity and relationship to the immunization by the investigators. NCI Common toxicity criteria (version 3.0) grades were applied (<http://ctep.cancer.gov/reporting/ctc.html>). Serious adverse events (SAEs) were defined as life threatening AE's, or AE's that resulted in significant or persistent disability, hospitalization, or death and were collected throughout the study period.

Injection site and systemic symptoms were recorded in a memory aid up to the next scheduled visit. Patients were asked to measure erythema and oedema (only Paper II). Standard urine analysis and extensive laboratory tests were taken.

The study period was 72 weeks during which the patients were monitored for safety, in total seven doctors visits. This was followed with clinical follow-up visits every 6 months, over a 5-year period, from the first immunization for long-term adverse events with special attention to autoimmune diseases, second malignancies, DFS and OS.

#### Paper IV

During the study, patients were regularly checked for performance status, routine blood haematology and chemistry analyses and serum tumour markers. AEs were assessed once weekly throughout the study period.

AE were assessed according to the National Cancer Institute Common Toxicity Criteria versions 2.0 (Groups A/B) and 3.0 (Groups C/D) resp. and considered related to treatment if a relationship was reported as possible or probable. SAEs were defined as life threatening AE's, or AE's that resulted in significant or persistent disability, hospitalization, or death and were collected throughout the study period.

### 8.4 Criteria for immune response

#### Paper I

ELISA: A patient was considered to have developed a significant response if IgM, IgA or IgE antibodies could be detected at at least two different time-points and was at least twice that of the pre-immunization value for that patient.

### Paper III and IV

- A patient was considered to have a *single assay immune response* if a response in one of the assays (proliferation, ELISPOT, Th-1 like cytokine secretion) was noted at one time point only against rCEA and/or CEA derived peptides (Paper III) or telomerase and/or ras (Paper IV). DTH was performed and included in overall induced immune response (OIIR) in Paper IV. Results of the ELISA assay were used descriptively (Paper IV).
- A patient was considered to be an *overall induced immune responder OIIR* if a positive response (see above) was noted in at least one of the assays at at least at two time points.

## 8.5 Statistical analysis

Statistical analyses were done using Statview® (SAS Institute Inc. version 5.0 USA) (Papers I, II, III, IV) as well as IBM SPSS statistics (version 19) (Papers I, III).

### Paper I

Comparison of pre-immunization anti-CEA titers of deceased and alive patients between groups were analysed by unpaired t-test. The relationship between survival and anti-CEA Ig-titers were analyzed by Cox-Regression analysis where the through mean values of IgM, IgA and IgE levels respectively for all patients at pre-immunization and at month 1, 2, 4, and 6 were calculated and listed from lowest to highest value. Patients with a through mean value below the median of all patients were labelled value 0. Patients with a mean value above the median of all patients were labelled value 1. This factor was analyzed in Wilcoxon Gehan exact test in relation to survival. As development of antibodies is “time dependant”, analysis of the prognostic impact of antibodies was performed after a fixed observation time<sup>[237]</sup>. As one patient died early, after 7 months, the 6 first months were used as observation time. The observation period and follow-up period were then held apart, avoiding “the pitfall of overlapping exposure and follow-up periods”<sup>[238]</sup>. Survival in relation to both anti-CEA IgA and anti-CEA IgG titers were also analysed. Patients were grouped according to those having both IgA and IgG anti-CEA antibody levels (through value) above the median (n=9) and the remaining patients (n=15). The relationship between anti-CEA IgA titers, GM-CSF and survival were analysed by Cox-regression analyses. The influence on survival by GM-CSF was assessed by univariate Wilcoxon Gehan exact life table test.

### Paper II

Changes in size of erythema over time were analyzed using non-parametric tests (Friedman test and repeated measure ANOVA). Continuous parameters (differences in size between Cohort A and B) were compared by the Mann-Whitney U test.

### Paper III

The non-parametric Friedman test for multiple comparisons was used to calculate relation of cell subsets at different time points. The Chi-square, Fisher test and one-way ANOVA tests were used to analyze differences in prognostic factors between groups and relation to immune responses. A p-value <0.05 was considered statistically significant. Life table landmark analysis with Wilcoxon Gehan exact test was used analyze a relationship



between OS as well as DFS to immune responses; landmark was set at 72 (part 1) and 52 (part 2/3) weeks, i.e. close to the end of immunological evaluation. The landmark method for evaluating OS and DFS by immune response selects a fixed time, chosen by the investigator prior to the start of the study. Patients still on study at the landmark time are separated into two immune response categories (non-responders and responders) depending on immune response at the landmark time point. Patients were then followed onwards in time from the landmark <sup>[239, 240]</sup>. One way ANOVA and Friedmans test (non-parametric) were used to assess PPD and PHA differences at baseline between patients and healthy donors.

#### **Paper IV**

The non-parametric Mann Whitney two-tailed rank sum test for comparison of independent variables and the two-tailed non-parametric Wilcoxon signed rank test for dependent observations were applied.

## **8.6 Ethics**

All studies were approved by the Regional Ethical Review Board in Stockholm.

## 9 RESULTS AND DISCUSSION

### 9.1 Paper I

#### **Induction of IgM, IgA and IgE Antibodies in Colorectal Cancer patients vaccinated with Recombinant CEA protein.** (*J Clin Immuno.* 32:855-856, 2012. Staff et al)

Previous clinical studies have indicated that IgG antibodies have the ability to induce apoptosis of tumor cells but IgM, IgE and IgA may also mediate tumor cell killing<sup>[241-246]</sup>. The presence of spontaneously induced IgM anti-CEA antibodies has been associated with improved survival in CRC patients<sup>[247]</sup>. However, there is scanty information on the Ig subclasses response after vaccination against cancer antigens. The clinical significance of the different Ig subclasses antibodies is not clear.

In a previously reported study, twenty-four resected CRC patients without macroscopic disease were immunized seven times with the tumor associated protein rCEA ± GM-CSF<sup>[138, 248]</sup>. Four different dose schedules were used over a 12-month period. A significant anti-rCEA-specific IgG1, IgG2 and IgG4 antibody response was shown and the induction of IgG correlated to improved survival.

Hence, a follow-up study of patients previously enrolled in this vaccination study done by our group was carried out. The aim of this study, reported in **Paper I**, was to assess the induction of IgM, IgA and IgE anti-CEA antibodies, during 36 months after vaccination with rCEA with or without GM-CSF as an adjuvant. Isolated IgA from one vaccinated patient containing high IgA anti-CEA titers was tested in ADCC and CDC, using CEA positive and negative cell lines. Patients were evaluated for DFS and OS. The findings were correlated to DFS and OS.

Most patients (83%) had a detectable IgM, a majority (54%) IgA and one patient had IgE anti-CEA antibodies before vaccination. Pre-immunization IgA, IgM and IgE anti-CEA titers did not differ significantly comparing deceased and alive patients.

GM-CSF significantly augmented the anti-CEA response for all three antibody classes ( $p < 0.05$ ). A strong IgM response was observed initially during vaccination with maximum titers at 3 months from start of vaccination. A rapid induction of IgA antibodies was noted already after one immunization, followed by gradual increase with a peak at 12 months. Repeated immunizations induced a gradual increase in the IgE anti-CEA antibody levels. Maximum IgE levels were noted at 15 months. The antibody curves over time (through values) in relation to GM-CSF were highly significant for IgM  $p=0.003$ ; IgA  $p=0.006$  and IgE  $p=0.007$ . The IgM-titers declined as expected much earlier than the IgA and IgE antibody levels. However, IgA and IgE anti-CEA levels clearly above pre-vaccination levels were still detectable 2 years after the last immunization. The dose of CEA had no significant effect on the IgM, IgA or IgE anti-CEA antibody responses.

Six patients in the CEA alone group and six patients in the GM-CSF group relapsed and died during follow-up. Two further patients in the CEA alone group died from other causes than CRC.

A significant correlation between survival and high IgA anti-CEA titers was noted ( $p=0.02$ ) irrespective of GM-CSF treatment.

Anti-CEA IgA antibodies could significantly lyse CEA positive tumor cells in ADCC ( $p<0.0001$ ) as well as in CDC ( $p<0.0001$ ).

The addition of GM-CSF was practically mandatory to elicit a vaccine induced IgM, IgA and IgE anti-CEA response as compared to patients receiving the vaccine alone. This finding corroborates our previous studies where GM-CSF was found to strongly enhance both the T cell and IgG responses [138, 248]. Fc receptors for IgA are found on monocytes/macrophages, polymorphonuclear neutrophils and NK-cells, i.e cells participating in ADCC [249]. In colon cancer patients elevated serum IgA immune complexes have been observed [241]. Moreover, IgA and IgG antitumor antibodies were shown to be equally effective in killing tumor cells in ADCC [241]. In conclusion, the observation that IgA anti-CEA antibodies were cytotoxic and associated with improved survival might indicate that also these antibodies may exert a clinical anti-tumor effect. Also, the duration of the humoral responses seen in Paper I, warrants further studies to evaluate rCEA as a therapeutic complement in patients with CRC.

## 9.2 Paper II

### **A phase I safety study of plasmid DNA immunization targeting carcinoembryonic antigen in colorectal cancer patients.** (*Vaccine*. 29:6817-6822, 2011. Staff et al)

The feasibility, safety and immunogenicity of therapeutic vaccination in CRC have been established. Different vaccination approaches targeting CEA in over 350 patients, including plasmid DNA as well as prime-boost vaccination strategies (74 patients) have only shown grade 2, and occasionally grade 3, toxicity [72, 138, 250-252].

In the present study, a plasmid DNA vaccine, encoding a truncated form of human CEA fused to a T-helper epitope (CEA66 DNA) was delivered three times i.d (2mg) or i.m (8mg), by a needle-free injector device, Biojector® to patients with colorectal cancer in the adjuvant setting. Five patients were included in each cohort (Cohorts A and B). Prior to the first vaccination, all patients received cyclophosphamide (300 mg/m<sup>2</sup>) intravenously (i.v). GM-CSF was administered s.c with each vaccination.

Safety and tolerability were evaluated in all ten patients. All patients reported local injection site reactions but the majority was mild. 89% of the total local AE in Cohort A were grade 1; the corresponding figure for cohort B was 85%. The frequency of patients with local grade 2 AEs increased during the 1<sup>st</sup>, 2<sup>nd</sup> and 3<sup>rd</sup> vaccinations and was 0%, 20% and 50% respectively in all patients. There was a significant increase in size of erythema at the injection sites from week 0 to week 6 in Cohort A, ( $p$ -value=0.015).

Systemic AEs were also mild and transient (grade 1-2). There was a tendency that fatigue, headache and myalgia were more frequent in the i.d cohort. No grade 3 or 4 adverse events possibly, probably or definitely associated with vaccination were reported. One SAE in one patient, in Cohort A, was not considered related to the study drugs. No signs of autoimmunity were seen.

During follow-up time, one patient (Cohort A) died at week 99 due to urine bladder cancer, diagnosed 72 weeks after start of treatment. One patient (Cohort B) had disease

recurrence at week 52 and was alive at data collection. Eight out of ten patients have no evidence of disease. A concern for DNA vaccination is the integration into hosts cellular DNA that could result in mutagenesis. DNA vaccines currently in test do not show of integration into the host genome <sup>[253, 254]</sup>. In light of the short time period between the initiation of immunization in this study and diagnosis of the bladder cancer in two patients, as well as the coexistence of risk factors as smoking and previous pelvic radiotherapy, integration of CEA66 DNA could not have resulted in this new malignancy <sup>[255]</sup>. No signs of autoimmunity were seen.

In conclusion, this phase I trial represents the first clinical trial on safety of a therapeutic vaccination strategy with plasmid DNA encoding human truncated CEA (CEA66 DNA), delivered by needle-free injection in combination with GM-CSF and cyclophosphamide in humans.

Skin contains higher numbers of immunoreactive cells than muscle <sup>[90]</sup>, which may explain the more frequent AE in the i.d cohort. The intradermal low dose DNA route indicated that skin may be a preferred site for immunization. Furthermore, the vaccine strategy was well tolerated, confirming the results of a preclinical model <sup>[256]</sup>.

### 9.3 Paper III

#### **Plasmid DNA immunization targeting carcinoembryonic antigen in colorectal cancer patients.** (*Manuscript, Staff et al*)

The method of DNA delivery as well as the localization of the expressed antigen can impact the outcome of DNA vaccination. The CEA66-DNA vaccine construct, evaluated in Paper II, encodes a modified, full-length, non-glycosylated form of CEA that is retained inside the cell. In preclinical studies, a T cell mediated response to CEA but weak antibody response was seen, using a needle-free pneumatic device (Biojector) <sup>[256, 257]</sup>. In contrast to the CEA66-DNA construct, DNA encoding wild type CEA (glycosylated) fused to a tetanus T-helper epitope (tetwtCEA-DNA), which is expressed as a membrane protein, induced a strong T cell as well as an antibody response <sup>[257]</sup>.

Delivery of DNA vaccine either i.m. or i.d by EP, may increase DNA uptake and antigen expression compared to needle injections <sup>[258, 259]</sup>. Short electrical pulses, at the site of vaccination, create transient pores in the cell membrane augmenting non-viral transfection of plasmid DNA. EP also causes a mild inflammation, with recruitment of APC at the site of injection without affecting persistence or integration of the plasmid <sup>[260, 261]</sup>.

In **Paper III**, we report immune data in an explorative study using CEA66-DNA and tetwt-CEADNA for immunization in combination with cyclophosphamide and GM-CSF in the adjuvant setting of CRC patients. Ten patients received i.d or i.m (cohorts I/II) CEA66-DNA by Biojector (for details see Paper II) at weeks 0, 2 and 6 (part 1). Ten patients; cohorts III/IV (part 2), received tetwt-CEADNA 400µg i.d by needle followed by electroporation at weeks 0 and 12. Cohort V (part 3) (n=6) included patients primed with CEA66-DNA and boosted with tetwt-CEADNA. GM-CSF and cyclophosphamide was included in the vaccination schedules.

Adverse events regarding part 1 has previously been published <sup>[262]</sup> (Paper II), AEs regarding part 2 and 3 are shown in Table 8. The most frequently reported AEs were local injection site reactions, fatigue and nausea, but the majority was mild. No signs of autoimmunity were seen.

**Table 8.** Frequency (%) of patients with AE treated with cyclophosphamide and vaccinated with tetwtCEA DNA without GM-CSF (part 2, cohort III; part 3, cohort V) and with GM-CSF (part 2, cohort IV) from baseline to week 24 (NCI Common toxicity criteria version 3.0 grades 1-3)<sup>a</sup>.

| Toxicity              | Cohort III (n=5) |        | Cohort IV (n=5) |        | Cohort V (n=6) |        | Total (n=16) |        |
|-----------------------|------------------|--------|-----------------|--------|----------------|--------|--------------|--------|
|                       | G* 1-2           | G 3    | G 1-2           | G 3    | G 1-2          | G 3    | G 1-2        | G 3    |
|                       | No** (%)         | No (%) | No (%)          | No (%) | No (%)         | No (%) | No (%)       | No (%) |
| <b>Local AE</b>       |                  |        |                 |        |                |        |              |        |
| Pain/discomfort       | 3 (60)           | 0 (0)  | 2 (40)          | 0 (0)  | 2 (33)         | 0 (0)  | 7 (44)       | 0 (0)  |
| Erythema              | 1 (20)           | 0 (0)  | 4 (80)          | 0 (0)  | 2 (33)         | 0 (0)  | 7 (44)       | 0 (0)  |
| Oedema                | 0 (0)            | 0 (0)  | 1 (20)          | 0 (0)  | 1 (17)         | 0 (0)  | 2 (13)       | 0 (0)  |
| Warmth                | 0 (0)            | 0 (0)  | 0 (0)           | 0 (0)  | 0 (00)         | 0 (0)  | 0 (0)        | 0 (0)  |
| Pruritus              | 2 (40)           | 0 (0)  | 1 (20)          | 0 (0)  | 5 (83)         | 0 (0)  | 8 (50)       | 0 (0)  |
| Induration            | 0 (0)            | 0 (0)  | 0 (0)           | 0 (0)  | 0 (00)         | 0 (0)  | 0 (0)        | 0 (0)  |
| <b>Systemic AE</b>    |                  |        |                 |        |                |        |              |        |
| Fatigue               | 2 (40)           | 1 (20) | 2 (40)          | 0 (0)  | 4 (67)         | 0 (0)  | 8 (50)       | 1 (6)  |
| Headache              | 1 (20)           | 0 (0)  | 1 (20)          | 0 (0)  | 0 (0)          | 0 (0)  | 2 (13)       | 0 (0)  |
| Dizziness             | 1 (20)           | 0 (0)  | 2 (40)          | 0 (0)  | 0 (0)          | 0 (0)  | 3 (19)       | 0 (0)  |
| Myalgia               | 1 (20)           | 0 (0)  | 0 (0)           | 0 (0)  | 1 (17)         | 0 (0)  | 2 (13)       | 0 (0)  |
| Nausea                | 4 (80)           | 0 (0)  | 3 (60)          | 0 (0)  | 2 (33)         | 0 (0)  | 9 (56)       | 0 (0)  |
| Anorexia              | 0 (0)            | 0 (0)  | 1 (20)          | 0 (0)  | 0 (0)          | 0 (0)  | 1 (6)        | 0 (0)  |
| Arthralgia            | 0 (0)            | 0 (0)  | 1 (20)          | 0 (0)  | 1 (17)         | 0 (0)  | 2 (13)       | 0 (0)  |
| Shivering             | 1 (20)           | 0 (0)  | 3 (60)          | 0 (0)  | 0 (0)          | 0 (0)  | 4 (25)       | 0 (0)  |
| Swelling              | 0 (0)            | 0 (0)  | 0 (0)           | 0 (0)  | 0 (0)          | 1 (17) | 0 (0)        | 1 (6)  |
| Chest tightness       | 0 (0)            | 0 (0)  | 0 (0)           | 0 (0)  | 0 (0)          | 0 (0)  | 0 (0)        | 0 (0)  |
| Diarrhoea             | 1 (20)           | 0 (0)  | 0 (0)           | 0 (0)  | 0 (0)          | 0 (0)  | 1 (6)        | 0 (0)  |
| Pruritus general      | 0 (0)            | 0 (0)  | 0 (0)           | 0 (0)  | 0 (0)          | 1 (17) | 0 (0)        | 1 (6)  |
| Rash/urticaria        | 0 (0)            | 0 (0)  | 0 (0)           | 0 (0)  | 0 (0)          | 1 (17) | 0 (0)        | 1 (6)  |
| Microscopic hematuria | 0 (0)            | 0 (0)  | 0 (0)           | 0 (0)  | 1 (17)         | 0 (0)  | 1 (6)        | 0 (0)  |
| <b>Metabolic</b>      |                  |        |                 |        |                |        |              |        |
| ALAT elevated         | 0 (0)            | 0 (0)  | 1 (20)          | 0 (0)  | 0 (0)          | 0 (0)  | 1 (6)        | 0 (0)  |
| ASAT elevated         | 0 (0)            | 0 (0)  | 1 (20)          | 0 (0)  | 0 (0)          | 0 (0)  | 1 (6)        | 0 (0)  |

a. No grade 4 AE was reported. \*G=Grade. \*\* Represents the number of subjects experiencing adverse events.

In total, 16 out of 20 (80%) patients mounted a single assay cellular response; 10/10 (100%) in part 1 and in 6/10 (60%) of the patients in part 2. The difference between the CEA vaccine naïve patients was statistically significant ( $p = 0.025$ ) in favour of part 1. OIIR was seen in 6/10 (60%) of the patients in part 1 and in 4/10 (40%) of the patients in part 2. The difference was statistically not significant.

In part 3, 5/6 (83%) of the patients mounted a single response in one assay. In part 3 OIIR was noted in 2/5 (40%) of the patients. All patients mounted a single assay cellular immune response in either part. In total, humoral responses were seen in two patients only.

There were no significant differences in either OS or DFS comparing single assay immune responders or OIIR in parts 1 and 2.

In the present study, the frequency of patients mounting a single assay cellular immune response was significantly higher following Biojector delivery (part 1) compared to vaccination with electroporation in CEA vaccine naïve patients (part 2).

An explanation might be a higher number of injections transfecting a larger number of cells and that Biojector delivery may evoke a substantial cell-trauma eliciting danger signals, enhancing an immune response induction<sup>[11, 12]</sup>. Immune responses are also higher when a higher dose of DNA plasmid is delivered<sup>[263]</sup>. Another explanation for a better immunogenicity in part 1 is the form of CEA-DNA used. The CEA-DNA construct used in part 1 generates a non-glycosylated protein that to a higher extent might be retained in the cytoplasm in comparison to the DNA construct used in part 2 yielding a glycosylated CEA protein which localizes to the plasma membrane. Thus, the CEA66-DNA might be more foreign compared to wild type CEA than the tetwtCEA-DNA used in part 2<sup>[264, 265]</sup>. Likewise, in melanoma, a more heterogenous substance compared to the wild-type were more effective in breaking immune tolerance<sup>[266]</sup>.

Mostly a Th-1 like cellular immune response was seen and only few individuals developed an antibody response, confirming the results in mice where CEA66-DNA primarily induced a cellular immune response<sup>[256]</sup>. Priming with CEA66-DNA and boosting with tetwtCEA-DNA did not augment the humoral response contradicting the results in mice by Brave et al<sup>[255]</sup>. Immune responses against CEA-derived peptides may suggest that CEA-DNA vaccination might induce both a CD4 and CD8 response. These results are supported by a preclinical study in mice vaccinated with CEA66-DNA, where a response against the B3 domain was mediated by CD4+ cells and towards the CTL-epitopes by CD8+ cells<sup>[256]</sup>.

The results in Paper III indicate that self-tolerance against the tumor-associated antigen CEA could be broken, although weak but long-lasting. Similar to CEA66-DNA, tetwtCEA-DNA, combined with cyclophosphamide and GM-CSF was safe<sup>[262]</sup>. The present study also indicates that T cell responses could best be induced against CEA-DNA delivery by Biojector, which in this respect seemed to be superior to CEA-DNA injection followed by electroporation for immune induction. Extended studies are warranted to discriminate dose and delivery devices to determine the clinical significance of CEA-DNA vaccination in patients with CRC.

## 9.4 Paper IV

### **Telomerase-peptide vaccination (GV1001) together with gemcitabine in advanced pancreatic cancer patients.** (*Manuscript, Staff et al*)

In **Paper IV**, we assess safety and immunogenicity in non-resectable pancreatic cancer patients using a 16 aa telomerase peptide (GV1001) for vaccination in combination with GM-CSF and gemcitabine as first line treatment.

Three different vaccine treatment schedules were used (A (n=6), B (n=6), C (n=5)). Groups A/B received GV1001, GM-CSF and gemcitabine concurrently. Group C received

initially GV1001 and GM-CSF while gemcitabine was added at disease progression. Group D (n=4) was treated with gemcitabine alone.

AE related to vaccination were mild (grades 1-2). Grade 3 AEs were few and transient. A higher dose of GM-CSF induced a higher frequency and severity of injection site reactions. Gemcitabine related side-effects were as expected<sup>[267]</sup> and without overlapping toxicity with the vaccine treatment. One SAE was initially suspected to be related to GV1001 or GM-CSF in one patient in group C, who developed hepatic dysfunction (grade 3) due to liver metastasis. Both median time to progression and overall survival were most poor in group C.

Four out of six (67%) patients developed a single assay immune response against telomerase and 3/6 (50%) against ras in group A. In group B, 4 out of 6 (67%) patients mounted a telomerase response and one (17%) against ras. A telomerase response in group C was noted in 2/5 (40%) patients and a ras response in 1/5 (20%) patients. In group D, a telomerase response was recorded in 2/4 (50%) patients and a ras response in 1/4 (25%) patients.

OIRR was only seen in group A patients, in one patient against telomerase and in two patients against ras. The cytokine pattern was that of a Th<sub>1</sub>-profile. A significant decrease in regulatory T cells over time was noted in patients in groups A and B ( $p < 0.05$ ).

The results might indicate that concomitant treatment with gemcitabine may not hamper the induction of an immune response but that delayed administration of gemcitabine might reduce the capacity to mount an immune response and favour tumor progression (group C).

Telomerase vaccination (GV1001) in combination with chemotherapy appeared to be safe but the immune responses were weak and transient. Based on the experience of the present study and of those of others including immune responses and clinical efficacy, measures have to be taken to augment the magnitude and duration of the immune response to GV1001. Furthermore, advanced pancreatic carcinoma patients might not be a preferred clinical setting for vaccine treatment, as is the case for other tumors and tumor cancer vaccines<sup>[268]</sup>. Maybe the GV1001 vaccine is not an optimal telomerase vaccine candidate, although it has been shown that immune responders to GV1001 vaccination survived longer than non-immune responders<sup>[165]</sup> and CLL patients exhibited spontaneous T cells recognizing GV1001, which could lyse autologous telomerase expressing leukemic cells<sup>[153]</sup>. The study might also support the notion that multiple immune read-out systems might increase the sensitivity to detect antigen specific immune responses<sup>[269, 270]</sup>.

## 10 CONCLUSIONS AND FUTURE PERSPECTIVES

Active immunotherapy is emerging as an important addition to conventional cancer treatments but many important questions remain. Effective strategies for overcoming immunosuppression need to be developed and patient selection are important. Optimal combinations of antigens, adjuvants and delivery vehicles need to be determined. Validation of surrogate endpoints and the identification of predictive biomarkers are equally important.

A concept that is gaining acceptance in the field of active immunotherapy towards cancer is that vaccine monotherapy is unlikely to succeed in generating a robust and long-lasting memory. Synergism between existing chemotherapy and immunotherapy may lead to more clinical responses <sup>[271, 272]</sup>. Gemcitabine and 5-Fluorouracil have been implicated in the suppression of MDSC activity <sup>[273, 274]</sup>. Cyclophosphamide induced leucopenia results in an expansion of immature dendritic cells favouring immune responses <sup>[275]</sup> and may suppress T<sub>regs</sub> <sup>[119, 276-279]</sup>.

The development of combination therapies is, however, associated with challenges. There is often little preclinical data on such combinations and phase I trials are required to show non-toxic combinations as well as to demonstrate immunological responses to the TCV. In Paper IV, gemcitabine combined with GV 1001 and GM-CSF did not seem to add to the toxicity but a decrease in the frequency of T<sub>regs</sub> was seen. As shown in Paper II and III, the combination of cyclophosphamide and CEA-DNA vaccination was well tolerated with mild and transient side-effects. No significant decrease in the frequency of T<sub>regs</sub> were seen, but the PHA responses increased in about 40% of the patients during vaccination, which might be due to reduced suppression induced by the cyclophosphamide treatment.

Combinations of complementary immunotherapies may further induce sustained anti-tumor responses. By combining plasmid DNA vaccination with other modalities of antigen delivery, such as recombinant proteins, in heterologous prime/boost protocols, both B and T cell immune responses can be enhanced <sup>[280]</sup>.

The results in Papers I and III combined, show a durable induction of anti-CEA antibodies following rCEA protein vaccination and long-lasting cellular responses following CEA-DNA vaccination. The induction of anti-CEA specific IgG antibodies as well as proliferative T cell responses, following rCEA protein vaccination, has previously been published <sup>[138, 248]</sup>. In a preclinical study designed as a DNA prime/protein boost setting, cellular immune responses were of higher magnitude in animals primed with CEA-DNA than in animals receiving repeated doses of rCEA protein <sup>[256]</sup>. In light of safety data presented in Paper II, a similar study would be feasible in humans.

Clinical studies have demonstrated the ability of different TCV strategies to induce antigen-specific T cells <sup>[281-284]</sup>. However, despite expanding antigen-specific T cells, in most clinical trials, tumors continue to grow <sup>[285-288]</sup>. Most clinical trials have been conducted in patients with advanced metastatic disease, and as tumors grow they evade immune destruction, for example by down-regulating HLA-expression <sup>[28, 289, 290]</sup>. In the present thesis, treatment with GV1001, GM-CSF and gemcitabine in patients with advanced



pancreatic cancer, mounted an immune response against telomerase in approximately two thirds of the patients in the best schedule. However, although the patients did not seem to be immune-hypo responsive as evaluated by PHA and PPD responses, immune response towards telomerase were weak and transient. In a number of clinical studies in which active immunotherapy failed to prolong survival, subgroup analyses suggested clinical benefit in patients with early or less aggressive disease <sup>[291-294]</sup>.

For cell surface antigens, both CTL and humoral responses may mediate antitumor activity, attesting the importance of activating both CD8+ and CD4+ T cells <sup>[72]</sup>. Previous studies on rCEA protein vaccination in CRC patients resulted in both cellular and humoral CEA-specific responses, and the IgG antibody responses to CEA correlated with increased survival <sup>[137, 138]</sup>.

Studies have suggested that IgM, IgA as well as IgE antibodies may have a role in eradicating cancer cells <sup>[241-246]</sup>. IgA antibodies have been proposed to be advantageous in tumor cell killing as compared to IgG <sup>[295]</sup>. In Paper I, rCEA vaccination combined with GM-CSF as an adjuvans, significantly induced durable antigen –specific humoral responses (IgA, IgM, IgE) against CEA and the induction of IgA was significantly correlated to survival. Furthermore, anti-CEA IgA could significantly lyse CEA positive cells. Most TCV studies focus on the induction of antigen-specific T cells. Future cancer vaccine trials might also assess the induction of the humoral response, including different isotypes.

Modifications of the antigen may alter the immune response. In Paper III, the CEA-DNA construct used in part 1 generates a non-glycosylated protein that to a higher extent might be retained in the cytoplasm in comparison to the DNA construct used in part 2 yielding a glycosylated CEA protein which localizes to the plasma membrane. Tumor associated MUC1 is frequently hypoglycosylated on myeloma cells and this is associated by a more efficient antigen processing by dendritic cells, leading to stronger T cell responses <sup>[296-298]</sup>.

DNA potency may also be improved by a) alteration to the plasmid itself to increase expression of immunogenicity (i.e. insertion of immune modulators) or b) placing DNA in/on microparticles acting as adjuvants for APCs <sup>[110]</sup>.

There is a need for development and validation of tools to identify patients who can benefit from a particular form of immunotherapy. Despite effort, we do not know which parameters of immune response, and which assays used to assess these parameters, are optimal for efficacy analysis. According to a workshop held 2011, involving FDA, National Cancer Institute and the International Society for Biological Therapy of Cancer-Society for Immunotherapy of cancer (iSBTc-SITC), several key issues need to be addressed to identify patients and patient-groups who will benefit from treatment <sup>[299]</sup>.

Recommendations included standardized handling of blood samples, use of robust and standardized assays. Furthermore, they recommend that several different assays used – to capture both the magnitude of the response, but also duration, quality and frequency.

The characterization of responders and non-responders is difficult, assays yield plenty data. Distinguishing them from assay variation and normal human variation is difficult. Tight response criteria, requiring positive responses in 2 post-therapy time points is recommended <sup>[168]</sup>. Obtaining multiple pre-therapy samples, at different time-points, can be

used to assess pre-therapy variability and published raw data enables the reader to interpret data <sup>[168]</sup>. In addition, new analytes and/or biomarkers needs to be standardized and validated by other investigators before they are correlated to clinical outcome <sup>[168]</sup>.

Looking ahead, an effective immune response that leads to an antitumor response requires not only an increase in immune activation but also reduction of suppressive immune checkpoints <sup>[278, 300]</sup>. The approval of mAb against CTLA-4 provided proof-of-principle for targeting immune suppressive checkpoints in the treatment of cancer <sup>[120]</sup> and mAb targeting PD1/PDL1 are in phase III clinical development. Synergy with TCV in mouse models suggests the potential for combining TCV with either of these mAbs <sup>[300-302]</sup>.

Moreover, the mechanisms whereby radiotherapy elicits tumor-specific immune responses have begun to emerge <sup>[303, 304]</sup>. Well designed trials are required to investigate this combatorial approach <sup>[305]</sup>.

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