



The Nature of Mutated Muc5ac, an Oncofetal Protein, Expressed in Colorectal Cancer: It's Role as a Therapeutic Immunogen

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Abstract

The present concept of effectively treating a malignant lesion, especially one in the process of metastasis, requires being able to turn on cytotoxic T cells that can recognize and destroy the tumor. All attempts at employing this system using TIL and other T cell populations have failed to date. Prehn in the 1950 s postulated that tumors act in a similar fashion to invading bacteria and viruses. Here the host can recognize the foreign invader by defining immunogenic proteins expressed on the cell membrane at the threshold level needed for recognition. He also believed that immunogenic proteins that characterize a malignancy exist, but are only present at a fraction of the level needed to induce tumor destruction. By pooling tumor membrane proteins, we were able to define 3 specific oncofetal proteins expressed only during fetal life. Later in life a viral or carcinogenic agent reactivates the needed gene to produce a post translational modification of the oncofetal protein. The more common of these tumor immunogens to be identified was a modified form of MUC5ac, Monoclonals to this target were produced GMP, tested in vitro and based on the strong ADCC noted in contrast to a CD8 response, were introduced for use in clinical trials in patients with metastatic colorectal cancer having failed all standard forms of therapy.

Introduction

Many have equated the cancer cell to a foreign host invader similar to bacteria and viruses that enter and infect the host. The cancer cell can also be defined in a similar manner, acting as a probable host "invader" [1]. The presence a cancer growth results however, in producing greater consequences. It is far more aggressive, with uncontrolled progression in growth which can lead to metastasis. Where the malignant cell is genetically programmed to invade and even destroy host cells in comparison to that of bacteria and viruses, the failure of the existing immune system to demonstrate any form of immune recognition allows for this process of continued growth of the tumor [2]. This phenomenon appears to be the result of a weakened tumor immunogenic state. As such the cancer cell is poorly recognized by host defenses and left to continue in its inherent growth pattern. In addition, as a means of self protection, the tumor cell does express inhibitory molecules on its surface which serve to help inhibit the host's immune system response. [3]. In order to have proper immunologic recognition of an invading cancer system with the ability of a patient to reject this type of invader, a threshold level of immunogenic protein that characterizes a particular malignancy must be presented to the host. In most of the tumor systems that we have evaluated, a well defined immunogenic oncofetal protein is present, but at such small levels, that the host is only capable of reaching what is termed a state of "surveillance". Here an occasional tumor cell may be destroyed but in general the developing tumor system is left to progress in its pattern of cellular proliferation. This proliferative state can only be interrupted from an immunologic standpoint, when an external source of antigen is delivered. This will then bring the threshold level of tumor immunogen to that level where recognition is accomplished. At this point, the tumor is then attacked by the host in an attempt to control if not eliminate it. The lack of tumor antigen expression was first postulated by Prehn [4]. He suggested that while the tumor cells did contain a low level of immunogen, that the use of pooled allogeneic antigen could possibly, when delivered at the threshold level, reactivates the needed immune response. He performed various experiment with mice showing that when the antigen was derived from the pooling of multiple tumor membrane proteins, that a threshold level could be achieved to prevent tumor growth. Decreasing the level of antigen resulted in regrowth of tumor. Challenging immunized mice with a different tumor system

OPEN ACCESS

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Received Date: 08 Jul 2016

Accepted Date: 25 Jul 2016

Published Date: 15 Sep 2016

Citation:

Arlen M, Arlen P, Dubeykovskiy A, Saric O, Coppa G, Conte C, et al. The Nature of Mutated Muc5ac, an Oncofetal Protein, Expressed in Colorectal Cancer: It's Role as a Therapeutic Immunogen. *Clin Oncol*. 2016; 1: 1096.

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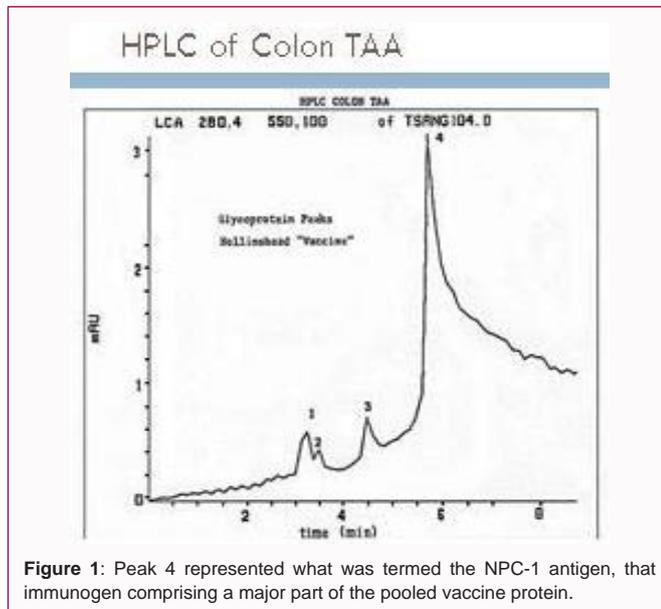


Figure 1: Peak 4 represented what was termed the NPC-1 antigen, that immunogen comprising a major part of the pooled vaccine protein.

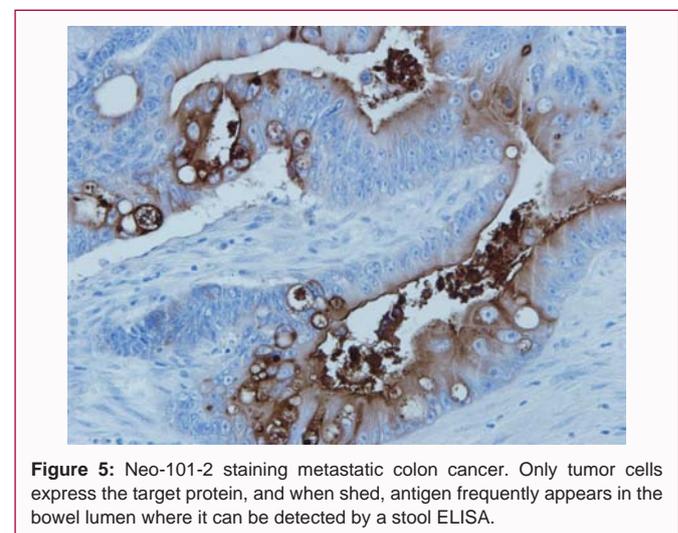
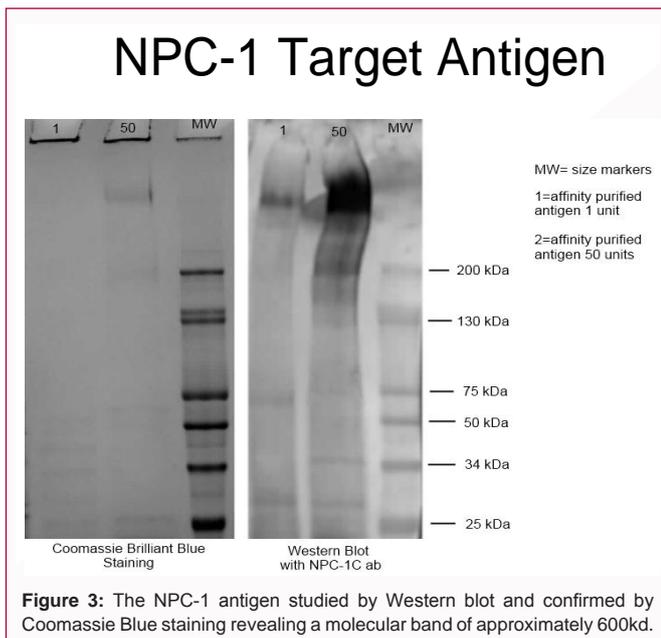
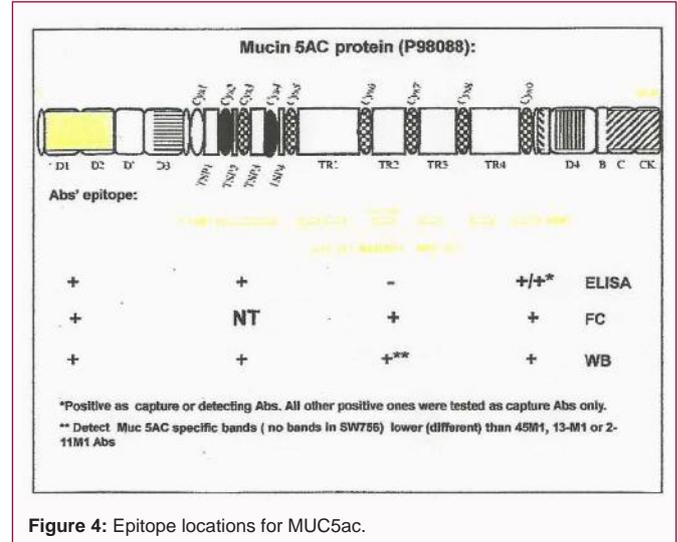
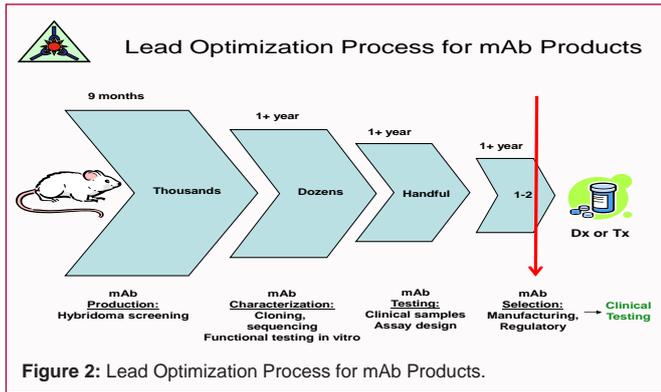
failed to prevent tumor growth. The conclusion to such experiments indicated that vaccines could be effective when delivered at threshold levels of the needed immunogen, but in addition that the vaccine also required specificity. In the 1970's Hollinshead, coming across the studies provided by Prehn [4] approached the concept of low antigen expression by tumors, she employed pooled allogeneic membrane protein to see if improving levels of tumor associated antigen (TAA) could result in enhancement of the host immune system to target residual tumor by the host (5). An initial Phase I study planned by Hollinshead et al. was performed in the 1980's. It explored the use of TAA therapy in patients with several malignancies but in a more detailed situation with those presenting with adenocarcinoma of the colon that were demonstrating metastasis or with a very high possibility of recurrence [6]. The immunogenic tumor proteins employed in each study were derived from pooled cancer membrane specimens obtained immediately after surgery [7]. Five years into the study there was evidence of marked improvement in survival. On that basis the FDA was approached to allow expansion of such studies toward developing a commercial product. At that time, the recognition of AIDS (HIV), Hepatitis C and HPV infections resulted in a change of procedure under FDA guidance. It was stipulated by FDA, that in order to advance such trials, a recombinant form of vaccine would be needed. It was suggested that the protein preparation used in the initial trials be employed to develop monoclonal antibodies for affinity purification and to eventually, by mass spectroscopy, define the structure needed to be sequenced in order to produce the required recombinant product. Most in our lab considered that contrary to our initial interpretation of the relative purity of the gel band used in preparing the final tumor immunogen as defined by patients being evaluated by delayed cutaneous sensitivity reaction (DHR), that multiple monoclonals would probably be derived from the hybridoma preparations that would be developed from the original antigen preparation. At that point any specific monoclonal antibody found to target the colon cancer protein would then be used for affinity purification and sequencing of the antigen to be isolated [8,9]. An HPLC study was performed (Figure 1) utilizing the therapeutic antigen from the initial clinical trials to develop hybridomas. At least 4 monoclonals were produced, each apparently specific to certain colorectal carcinoma cells. None reacted with normal colonic tissue.

Three of the mAbs matched the peaks seen on HPLC with that mAb defining the major peak 4 termed as NPC-1 or Neo 101. The other mabs studied were labeled as 31.1 and Neo 201. This region (peak 4), comprised the major portion of the antigenic component of the allogeneic vaccine used in the initial therapeutic trial.

Defining those oncofetal proteins expressed in the HPLC peaks

The murine monoclonal antibody termed mAb NPC-1 (Neo-101) was directed against and defined that portion of the immunogenic protein peak 4 seen on the HPLC. This particular protein component was expressed in more than 60% of the colon cancers examined by IHC. It proved to be non reactive, and as such absent in normal adjacent colon tissue [10]. Monoclonal NPC-1C (Chimeric), developed at Precision Biologics, was found by mass spectroscopy, to represent that mAb shown to target an altered form of the oncofetal form of MUC5ac. This mutated protein represented the unique Tumor Associated Antigen (TAA) comprising a major fraction of the vaccine that was previously tested in our Phase I clinical colon trial. The antibodies derived from the hybridomas produced from the original TAA, were used for further immunopurification and mass spectroscopy of the several components of the allogeneic vaccine. The resulting major antigenic component (peak 4) that was defined, was found to be homologous to the antigen MUC5ac, an oncofetal protein turned on in the latter part of fetal development [11]. The NPC-1 (Neo-101) monoclonal employed in our studies was developed at two separate intervals in our lab. The initial Neo -101 was expressed in the original bioreactor facility at 140 mg./L of bioreactor fluid and showed some minimal levels of toxicity in several patients. Utilizing the high expression vector developed by Selexis, the new mAb was now expressed at 2000 mg./L bioreactor fluid. Studies of bio similarity were required and revealed that the now termed Neo -102 was non reactive against red blood cells so that any minor degree of hemolysis was gone, there was a higher degree of ADCC and immunohistochemistry (IHC) of tumor specimens expressing the antigen appeared cleaner. Though both defined similar epitopes, they were probably glycosylated somewhat differently. This new version of the mAb targeting the altered form of MUC5ac antigen is now in a position for commercial development. The NPC-1 antibody was chimerized by genetically replacing the murine constant regions with human IgG1 constant regions, and designated as NPC-1C. The need of course was that for therapeutic purposes, the mechanism for tumor destruction was through an ADCC (Antibody Dependent Cell Cytotoxicity) response where the human Fc contained the receptors for carrying the patients NK cells to the surface of the tumor where NK destruction of tumor could be initiated. Several laboratory studies employing fresh-frozen or formalin-fixed tissues indicate that the NPC-1C antibody stains approximately 60% - 70% of colon malignancies. This supported the need for the antigenic protein to be highly expressed in the tumor cells without cross reacting with normal tissue. This potential allows the Neo-102 mAb and other similar monoclonals that we have developed to be used diagnostically, not only in terms of IHC, but in the development of an accurate serum ELISA prior to use as a therapeutic [12-14]. (Figure 2) shows the Optimization Process for use of the mAbs.

Preclinical pharmacology and toxicology studies demonstrated statistically significant tumor growth inhibition by the mAbs without significant toxicity, as judged by clinical observation, Studies of body weights, food consumption, ophthalmology and clinical pathology in test animals, supported the rationale for development of NPC-



1C antibody for clinical testing since it was the major product produced by immunization with the tumor antigen. An SDS PAGE electrophoresis and Western Blot was performed using the NPC-1 (Neo 102) chimeric of affinity purified NPC-1 antigen (50ugm/well, lanes 1 and 2) from LS174T lysate (Figure 3). Note that the high molecular wt. band (approx 600 KD) reacted with Neo-102C antibody in Western blot. The Coomassie blue band was later excised and proceeded to MALDI-TOF analysis which revealed its identity as a MUC5ac modified protein of approximately 600kd.

Background

The origin of the colon oncofetal protein

When the fetus matures in the final term of pregnancy, the gene needed for producing the oncofetal (immunogenic) protein required for organ function is demethylase. This allows activation of that protein or molecule necessary for the fetus to progress to term. In the case of the colon, the MUC5ac gene is activated to produce the mucin needed for the bowel and lung to function properly. Just prior to completion of pregnancy, this gene is re: methylated. It is however re expressed in some patients later in life where a modification of the methylated gene by a virus or carcinogen takes place. The oncofetal protein expressed by this newly modified “malignant cell” has little resemblance to the original protein functioning in the fetus. During this post translational state of expression, the modified protein

becomes immunogenic, thus characterizing the nature and function of the tumor. Such post translational modifications of the oncofetal protein are not found in normal cells and do not react with those monoclonals directed against the original fetal (antigen) protein. The commercial monoclonals (mAbs) to MUC5ac used to identify cystic fibrosis products do not recognize the mutated form of the oncofetal protein and the monoclonals (Neo -101 and 102) developed against the post translation molecule do not see the cystic fibrosis molecule. By Immunohistochemistry (IHC), the mAbs that we have developed for the modified MUC5ac protein only target the tumor protein in the cancer cell and not in the adjacent normal cell [15,16]. As such when the mAbs are employed therapeutically, there is a requirement that the tumor specimen be evaluated by IHC to assure that there is adequate recognition of the target protein by the monoclonal antibody being employed for therapy. (Figure 5) illustrated the specific staining needed to identify the potential of the tumor to respond to treatment.

Discussion

Our understanding of the nature of tumor induction took many years to understand. It was recognized early on, that certain viruses could induce malignant transformation. This was first noted with the Rous sarcoma virus transfecting the chicken fibroblast and later with the Rauscher virus inducing malignancy. It took Varmus, during his

Normal appearing colonocytes at the margin of a colon cancer resection

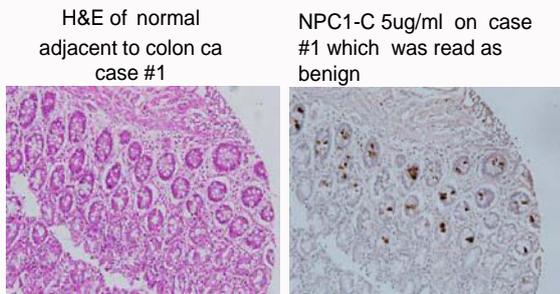


Figure 6: Normal appearing colonocytes that have undergone malignant transformation and are seen expressing tumor protein.

time at NCI, to define the mechanism by which the transformation process first took place. That is that the virus would enter the DNA of the nucleus, and in terms of the Rous virus, a defined site on the chromosome termed the SRC gene would come into play and function in the mechanism of malignancy [17]. Next it was noted that when the SRC gene was activated that it induced a neo-protein that when present would present features of the sarcoma cell that was nonexistent in the normal fibroblast. This mechanism of induction was considered to be a prominent feature in most forms of malignant transformation. Why the tumor continued to progress in patterns of growth without interruption by the host immune system was again a matter of speculation. It was not until Prehn had demonstrated in in-vivo models that the tumor in actuality, behaved as a foreign cell, not completely different from that of invasion by bacteria and viruses, that a potential approach toward utilizing the host immune system for tumor therapy became a reality. The major issue was in defining the nature of the oncofetal protein that represented the specific malignancy and making sure that it could be delivered to the host wherein a proper level of an immune response could be initiated.

Prior to the realization that specific active immunity utilizing tumor immunogens and their corresponding monoclonals had the potential to bring many malignancies under control, many therapeutic products for advanced and recurrent colon cancer were and still are widely employed. . But despite the development of several new treatment regimens for this particular form of malignancy (colorectal cancer) little if any benefit has been appreciated at the clinical level. A directional change from combination chemotherapy to immuno-chemotherapy is presently in a transition stage of clinical development and appears to definitely suggest an improvement in therapeutic value. . Some immunotherapeutic in 1998, Trastuzumab (Herceptin) in 1999, Alemtuzumab (Campath) in 2001, Bevacizumab (Avastin) in 2004, Cetuximab (Erbix) in 2004, and Panitumumab (Vectibix) in 2006. More recently, the number of new monoclonal antibodies on the market has increased with Trametinib, Alectinib, and Elotuzumab. Many other monoclonal antibodies are currently in clinical trials as monotherapy.

The ideal monoclonal to employ is the one that can characterize the tumor via its oncofetal protein. As such the mAb becomes both diagnostic and therapeutic as seen with the Neo-101 and 102. Such mabs demonstrating high levels of ADCC can be used alone, but probably become more effective when used in combination

with other therapies. This monoclonal antibody, when used in the treatment of solid tumor malignancies should of course target an immunogenic protein, probably an oncofetal protein characterizing the tumor rather than a growth factor expressed on the surface of the tumor but seen as well in many normal tissues. Such oncofetal proteins do not appear to mutate as we find with the epidermal and vascular growth factor targets. This latter group can mutate every 4-6 months requiring that the target be reevaluated due to loss of recognition by the therapeutic antibody Those antibodies used for targeting an immunogen such as MUC5ac in colon tumors cover a broad spectrum of transformation processes and therefore can also be used for defining the premalignant as well as metastatic lesion.

Summary and Conclusion

The concept of the need for immunotherapy as part of any clinical study to control advanced colorectal cancer arose from the initial trials of Hollinshead in the 1980's. In these studies, pooled allogeneic colon cancer membrane protein was employed. After 5 years, significant improvement in survival was noted, but FDA would not allow further use of such preparations. They realized the possibility that any such vaccine that could be developed by pooling of tumor specimens could possibly be contaminated by viruses such as HPV and HIV. In an attempt to produce a recombinant vaccine, monoclonals were developed from the original vaccine for immune purification and sequencing. It became apparent following the development of the needed monoclonals, that reevaluation of the response to vaccine data needed to be reviewed.

We quickly recognized at this time, that the expected cell mediated response in tumor destruction, played only a small role. It appeared that the major mechanism for tumor destruction was via ADCC and not via a cytotoxic T cell response As such those mAbs that we had developed for other needed uses, were now evaluated for their therapeutic potential.

The first in human investigational study of NPC-1C (Study # NEO-0901) using NEO-101, was initiated in patients with advanced colorectal cancer. Pre-clinical testing showed minimal toxicities. In 26 subjects treated in this study, side effects possibly attributable to NPC-1C (Neo 101) included hives, chills, flushing, shortness of breath, lower back pain, hematuria, elevated creatinine, elevated LDH, decreased haptoglobin, and anemia. This was virtually eliminated when monoclonal Neo-102 was used to replace the original version.

After administration of NPC-1C (NEO-101) to 26 subjects in the NEO-0901 study, the glycoengineered form of NPC-1C, called NEO-102, was manufactured for improved stability and decreased RBC agglutination.

The clinical trial, NEO-0901, dose escalation phase was re-initiated with the new formulation, and the first combination clinical trial using NPC-1C and gemcitabine (Study # PB-1201) is underway. NEO-102 has been administered as monotherapy in 100 patients with colorectal cancer with a very favorable toxicity profile. The trials employing the high production system mAb Neo 102, were established and have now completed the Phase II studies and as reported have shown indications of a significant benefit to the host [18]. Since the MUC5ac related antigen is the most common immunogen found among the tumor specimens examined, we are specifically targeting that protein, a post translational modification of MUC5ac with its corresponding mAb Neo 102.

The work performed as part of the clinical trial defined the sequence of the mutated MUC5ac protein and provided the peptidomimetics of an NPC-1 epitope derived from MUC5ac. This included composition comprising the amino acid sequence of the epitope binding site defined by Phage display. That is the polypeptide: F(PHE) P(PRO) E(GLU) D(ASP) Y(TYR) F(PHE) R(ARG) Y(TYR) T(THN(ASN) Q((GLN) K(GLY)). This sequence proved to be that fragment of the 600kd MUC5ac molecule responsible for antibody production and thus could potentially be utilized for eventual peptide vaccine therapy as a preventative post surgical resection in high risk patients.

At the 2015 ASCO GI Symposium [19] results of the Phase 1 study of Ensituximab (Neo-102) in chemotherapy refractory metastatic colorectal Cancer Study were presented in a poster session. The study revealed a maximal tolerated dose of 3.0 mg/kg IV every 2 weeks. The overall survival observed in this study demonstrated 10.4 months comparing favorably to the historical control for a similar population of patients with advanced colorectal Ca (5 months). This led to a larger Phase 2 multi-center colorectal cancer study using Ensituximab in the same patient population. This study is finalizing with the results employed in planning for the Phase 3 study. We are planning to introduce the immunotherapy at the time of initial recurrence where the immunotherapy will be given in conjunction with chemotherapy. The latter will be given in doses that do not suppress the immunocytes but will function to minimize to shedding of inhibitory molecules produced by the tumor that might interfere with optimum tumor control.

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