Designing vaccines to prevent breast cancer recurrence or invasive disease

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Keywords: antigen • breast cancer prevention • MHC Class II • Th1 • vaccine

Introduction

With the emerging importance of the tumor immune environment in both invasive and preinvasive breast cancer, vaccine therapy may provide a well-tolerated, durable therapy for preinvasive breast cancer by preventing both recurrent preinvasive disease and progression to invasive disease. Preinvasive breast cancer is ideal for vaccines because the disease is slow growing and the patients are not immunosuppressed. However, design of the vaccines are critical including consideration of appropriate antigens, selecting Th1 stimulating epitopes, and using multiantigen vaccines to address tumor heterogeneity.

The importance of the immune system in preinvasive breast cancer

Although breast cancer has not historically been believed to be an immunogenic tumor, the importance of the role of the tumor immune environment is emerging for both in invasive and preinvasive breast cancer. In invasive breast cancer (IBC), increased cytotoxic CD8⁺ T-cell infiltrate shows improved breast cancer specific survival [1] whereas increased FOXP3⁺ T-cell immune suppressive infiltrate is associated with inferior survival [2]. On examining specific breast cancer subtypes, a subset of triple negative breast cancer defined by robust immune infiltrate

shows similar prognosis to luminal A tumors [3]. Preinvasive tumors and high-risk breast lesions also have increased immune infiltrate seen as early as benign proliferative breast disease, for example, CD3⁺ T-cell infiltration increases from the mean in normal breast of 2.8 ± 1.0 to 81.5 ± 14.0 in benign proliferative breast disease, to 84.0 ± 14.9 in ductal carcinoma *in situ* (DCIS) and to 103.7 ± 3.9 in IBC [4]. Similar to invasive breast cancer, the populations of immune cells present in preinvasive disease are important for prognosis. A study of 62 DCIS tumors showed that increased FOXP3⁺ infiltrate conferred a shorter relapse-free survival (HR: 2.81; 95% CI: 0.99–7.99; $p = 0.05$) [5]. Furthermore, gene expression profiling studies demonstrate that a chronic inflammatory signature is present in both DCIS and IBC [6], and a T helper 1 (Th1) proinflammatory signature demonstrated improved outcome as compared with a T helper 2 (Th2) immune suppressive signature ($p = 0.0394$) [7]. These data suggest that as early as preinvasive disease, the immune system mounts an adaptive immune response to the tumor; therefore, if the immune system can be stimulated to mount the appropriate antitumor Th1 immune response to these preinvasive tumors it may be possible to prevent recurrence of the preinvasive lesions or progres-

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sion to invasive cancer. Vaccine therapy should work well for patients with preinvasive breast cancer because these patients have slow growing disease, a low tumor burden and do not have a suppressed immune system. However for vaccine therapy to be effective, the vaccine has to be carefully designed to identify antigens that are present in preinvasive disease, the vaccine has to be designed to trigger a Th1 immune response and the heterogeneity of preinvasive breast cancer has to be addressed by targeting multiple immunogenic proteins present in the lesion.

Identifying appropriate preinvasive vaccine antigens

Antigen targets for preinvasive breast cancer should be immunogenic proteins that are overexpressed in high-risk DCIS or other preinvasive breast lesions and are associated with an increased risk of developing invasive breast cancer [8]. Overexpressed proteins are believed to be immunogenic by revealing cryptic epitopes and by increasing the number of peptides available for recognition by the immune system [9]. Three examples of possible preinvasive antigen targets include IGF1R, IGFBP2 and HER2. IGF1R is overexpressed in both preinvasive and invasive breast cancer and has increased autoantibodies in breast cancer patients as compared with normal donors ($p < 0.04$) [10]. Increased autoantibodies to IGFBP2 have been seen in breast cancer patients over controls ($p = 0.0008$) and IGFBP2 has been shown to be important in pathogenesis of both preinvasive and invasive breast cancer [11]. IgG HER2 specific antibodies are found to be increased in patients with the HER2⁺ breast cancer subtype, and patients with high levels of IFN- γ secreting HER2-specific T cells are significantly more likely to have 3+ overexpressing tumors ($p = 0.001$) [12]. Furthermore, HER2 is also found overexpressed in DCIS and confers a higher risk of progressing to invasive disease [13]. Importantly, all three of these preinvasive antigens are also overexpressed in invasive breast cancer; therefore, a vaccine against these targets should not only prevent recurrence of preinvasive disease but also prevent against developing invasive disease.

Designing vaccines to produce a Th1 immune response

The proinflammatory Th1 immune environment promotes the best antitumor response; however, most tumors stimulate a Th2 immunosuppressive environment [14]. For example, PBMC from breast cancer patients had an increased Th2 response to MHC class II predicted IGF1R epitopes as compared to matched normal controls [10]. Th1 immune response both destroys the current disease and provides immunologic memory

to protect against recurrence or progression because of IFN- γ associated inflammatory cytokines which are necessary for enhanced function of antigen presenting cells, epitope spreading, activation of the CD8⁺ T cells and immunologic memory [15,16]. Using MHC class II vaccines which stimulate Th1 CD4⁺ T cells can both increase recruitment of CD8⁺ T cells to the tumor and allow them to be active by promoting a Th1 immune environment [17]. In the preclinical transgenic mouse mammary tumor model TgMMTV-neu, an MHC class II vaccine against IGFBP2 could inhibit tumor growth and this antitumor function was CD4⁺ T-cell dependent [11]. Furthermore, MHC class II epitopes from the same protein can either stimulate a predominantly IFN- γ Th1 or IL-10 Th2 response and inclusion of any Th2 epitopes in a vaccine prevents the antitumor immune response. For example, in the protein IGFBP2, the N terminus stimulated predominantly a Th1 IFN- γ immune response and as a vaccine could inhibit tumor growth in the transgenic TgMMTV-neu mice. However, when the vaccine included both the N terminus and the C terminus (which stimulated predominantly a Th2 response) there was no tumor inhibition [18]. For effective prevention vaccines, epitopes need to be selected that primarily promote a Th1 response to avoid the immunosuppressive Th2 epitopes that prevent an antitumor immune response.

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Multiantigen vaccines to address tumor heterogeneity

Preinvasive breast tumors are heterogeneous, therefore, there is differential expression of target proteins even in the same premalignant lesion [19]. Since HER2 had been shown to be both overexpressed and immunogenic in preinvasive breast cancer, a dendritic cell vaccine directed against HER2 was tested in patients with HER2⁺ DCIS [13]. The HER2 vaccine was immunogenic and 91% of the subjects had greater than fivefold response to HER2 peptides by IFN- γ ELISPOT after vaccination. Furthermore, there were increased HER2 peptide responsive CD8⁺ T cells in DCIS tissue by FACS in 88% of patients after vaccination. However, 84.6% of the patients had residual DCIS, although 55% of patients had decrease in size of DCIS, and the DCIS remaining had lost HER2 expression [13]. Using a single antigen vaccine allowed for selection of clones that do not overexpress the particular immunologic

target. Furthermore, HER2 is only expressed in a subset (~20%) of DCIS; therefore, single antigen HER2 vaccines only address a subset of DCIS.

A preclinical multivalent vaccine developed by our group has demonstrated that vaccination with a vaccine containing HER2, IGF-IR, and IGFBP2 epitopes can prevent both DCIS and progression from DCIS to invasive breast cancer in the TgMMTV-neu transgenic mouse mammary tumor model which develop tumors similar to luminal B breast cancer [20]. When mice were immunized at approximately 18 weeks old, the multiantigen vaccine was able to block development of palpable lesions in 65% of the mice as compared with adjuvant alone ($p < 0.0001$) and improve overall survival ($p < 0.0002$) in TgMMTV-neu mice. Furthermore, while animals vaccinated with the single antigens had similar disease-free survival ($p = 0.965$), the disease-free survival of the animals vaccinated with the multiantigen vaccine was significantly improved over any of the individual antigens ($p = 0.018$). The tumors that did develop in the multiantigen-vaccinated mice were slower growing and demonstrated increased CD8⁺ T-cell tumor infiltration as compared with control-vaccinated mice ($p = 0.003$). Similar prevention of invasive disease was found when vaccinating the very aggressive C3(1) Tag transgenic mice which develop tumors that are genetically similar to triple negative breast cancer ($p < 0.0001$). The protection was CD4⁺ mediated and depletion of the CD4⁺ T cells caused loss of the tumor inhibitory effect ($p = 0.013$) [20]. These studies give preclinical evidence that a multiantigen vaccine using Th1-specific MHC class II epitopes

can be effective at preventing development of both preinvasive and invasive breast cancer. The immunogenicity of this vaccine in women will soon be tested in a clinical trial.

Conclusion

The emerging role of the immune system in breast cancer has shown that the quantity and composition of the immune response to the tumor affect prognosis even in preinvasive lesions. Using intelligent vaccine design to ensure that the vaccines can stimulate a Th1 immune environment and target multiple antigens has been shown to both prevent preinvasive and invasive disease in preclinical transgenic mouse models where without treatment all mice develop mammary tumors. Preinvasive breast cancer would be an ideal disease for vaccine therapy and vaccines would address an unmet need in these patients for a durable, safe, well-tolerated therapy that has the potential to prevent both recurrence and progression.

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