Salivary Protein Antigens for Breast Cancer Biomarkers

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Abstract

Breast cancer remains the world leading cancer type in women. Its management requires a plethora of techniques from imaging to serum analysis; the latter using the presence of mostly protein-based biomarkers for the detection, classification and prediction of breast cancer’s behavior. Saliva is the latest medium in which detection of these biomarkers is being carried out. In this review, we would like to discuss salivary biomarkers for the detection of breast cancer.

Introduction

Breast Cancer is the most regularly diagnosed type of cancer in women in the world, making up on its own 25% of all cases, or nearly 2 million new cases in 2018, and 15% of all cancer related deaths, or around 626,700 deaths for that same year. [1].

The management of breast cancer patients is usually done through biomarker detection, their role is particularly important in deciding the type of systemic therapy for administration [2].

A biomarker is a molecule (protein, nucleic acids, etc) found in either a bodily tissue or fluid that changes its concentration by the presence of one or more types of cancer. It is the product of either the tumor itself or the immune response to a tumor. The ideal biomarker has the potential to detect small tumors with great specificity and selectivity in order to either inform for an early diagnosis or help in the tumor’s screening [3].

Biomarker detection technology have also progressed to the point where it plays a key role in improving conventional medical assessments to serious systemic diseases. Hence, the development of a salivary sensitive assay for the accurate, rapid, and non-invasive identification of breast cancer biomarkers would greatly benefit breast cancer’s detection and screening.

When compared to blood, saliva holds several advantages as a diagnostic medium; it is non-invasive and readily available in large quantities, allowing for repeated collection without patient’s discomfort [4].

In addition, salivary biomarkers are a filtrated fraction of their blood counterparts, as such, their analysis offers an advantage through their reflection of the physiological conditions of the body. Analysis of salivary samples can be used to monitor and predict clinical status of systemic diseases [5].

Because of all these advantages, saliva-based detection has been garnering increasing interest and technology based on their detection is offering a promising new clinical strategy for breast cancer [4].

In this mini review, we would like to expand on the salivary protein biomarkers used in breast cancer diagnosis, their biology and their clinical use.
Saliva

Saliva is a biological fluid found in the mouth; it derives from the secretion of the salivary glands in the buccal cavity.

The composition of whole saliva varies depending on a large number of factors and is in itself highly diverse with both body secretions, molecules, and the buccal cavity’s microflora, hence, in recent years, interest has been increasing to develop technologies capable of using saliva as a diagnostic medium [6].

Saliva for diagnosis

The main reason why saliva is garnering that much attention is because of its property of being a filtrated fraction of blood, as such its composition reflects the body’s physiological state. In addition, saliva collection is safer, simpler, and is non-invasive when compared to that of blood. For the diseases where saliva is used for detection, the medium showed great promise and high clinical appeal [5].

Despite that, saliva analysis is considered less reliable than blood or urine analysis; the reason for that is the high difficulty in managing the salivary matrix due to the influence of salivary flow rate on its pH and composition; while also containing lower concentration of said analytes to begin with [7]. In addition, collection methods affect salivary composition greatly, the use of cotton swabs and/or chewing gum to stimulate salivary secretion for example have an effect on decreasing protein concentration while increasing salivary flow rate; while passive drool have the opposite effect that can be attributed to which salivary gland is activated during salivary collection [8].

Saliva and breast cancer

Not many articles have explored the potential of using saliva to detect breast cancer protein biomarkers [1]; the most prominent article regarding this subject was written by Streckfus et al. [9], nearly 20 years ago, it highlighted the correlation of six breast cancer biomarkers found in saliva and their relation to the progression of the disease, these biomarkers will be explained further bellow. However, the small number of participants of this research cast doubt. Since then however, many researchers have focused their attention on Mucin 1 and on Human Epithelial Receptor 2 (HER2) [10-14], while others have focused on other non-protein biomarkers such as microRNA, and metabolomics [4], however, discussing non-protein antigens is beyond the scope of this review.

Breast Cancer Biomarkers

Epidermal growth factor receptors family

The epidermal growth factor receptor family is frequently over-expressed in human cancers. They are one of the 20 families of human receptor tyrosine kinases (RTK). They are 185 kDa transmembrane glycoproteins that get activated by binding with epidermal growth factors (EGF), EGF-like, or neuregulin ligands. The binding of these proteins form homodimers and heterodimers, which activates their intracellular tyrosine kinase domain leading to the activation of several important oncopogenes downstream in the signaling cascade [15].

The epidermal growth factor receptor (EGFR) also known as HER1 and ERBB1 has an important role in starting the signaling that directs epithelial cell behavior (an example of which can be found in Figure 1) and subsequently, tumor cells of epithelial origins [16]. EGFR overexpression often results in a poor prognosis and lower survival rates [17]. It has been established for a while now that EGFR presence is inversely correlated with Estrogen Receptors (ER) [18,19], despite the fact that extensive aspect of its function remains unexplored [20], it has proven to be a useful marker for the few subsets of breast cancer in which it has been fully explored (for example triple negative breast cancer [21] and inflammatory breast cancer [22]). HER2, also known as ERBB2, is unique in that it lacks a ligand; its signaling depends upon heterodimer formation [15].

HER2 is over-expressed or its gene is amplified in approximately 20% of breast cancer patients, and is associated with having a bad prognosis, thus requiring anti-HER2 therapy (trastuzumab or lapatinib) to enhance the odds and length of survival [23].

Mucin 1

Many different names have been used for Mucin 1 (MUC1) over the last 25 years, the most commonly used name however is the protein’s serum markers for breast cancer antigen 15-3 (CA 15-3) and cancer antigen 27.29 (CA 27.29). MUC1 is a large cell surface glycoprotein with a molecular weight of around 120 to 225 kDa, this variation is due to the large number of O-linked glycosylation sites on its extracellular domain.

MUC1 go through auto-cleavage resulting in 2 different subunits: the amino terminal subunit (MUC1-N) and the carboxyterminal subunit (MUC1-C). The cleavage occurs at a serine residue near the N-terminal in a GSVVV motif which is located in the extracellular SEA domain. A

**Figure 1:** Structure and function of the human epithelial receptor proteins [24].

**Figure 2:** Schematic representation of MUC1 protein [25].
picture of MUC1 structure can be found in Figure 2 [25].

MUC1’s function is to provide protection against fluctuation of pH and osmolarity, lubrication, and hydration, to the apical surface facing the lumen of ducts by trapping cellular debris on the cell surface. MUC1 can also be found in milk where it acts as a distraction for different types of infective pathogens [26]; its function as a regulatory protein for T-cells is newly discovered and its potential as an immune system regulator is still being discovered [27].

MUC1 is altered in a number of diseases including but not limited to Crohn’s disease, Inflammatory bowel disease, interstitial pneumonitis, and many others. While in cancer, it is overexpressed in several types of cancer from breast and ovarian to prostate, lung and pancreatic cancer. Because of that, it is considered as a universal marker for human cancer.

Because of its universality, many types of serum tests were developed for its detection, chiefly among them are CA 15-3 which is considered a prognostic biomarker for breast cancer worldwide and CA 27.29 that is used to detect early recurrence of breast and ovarian cancer [26,28]. Its mechanism with breast cancer relates to its ability to aid in immune cell evasion due to MUC1 interactions with siglets or Sialic acid based lectins found on a number of immune cells [29].

In terms of subtypes, there are no evidence found for a link between preoperative CA15-3 levels and different breast cancer subtypes [30].

**Tumor protein 53**

Tumor Protein 53 (p53) is a remarkable multifunctional protein that plays a role in cell cycle and differentiation, inflammation and immune response, transcription and DNA repair, epigenome, and lastly senescence and autophagy.

In normal cell function, p53 is maintained at low levels, however, being an influential inductor of apoptosis, p53 level increases in response to diverse stress conditions through different posttranslational modifications, causing p53 stabilization and activation leading to apoptosis [31].

In breast cancer, p53 has been in case studies from its prognostic and predictive behaviors [32] toward development of several therapeutic strategies relating to it [33].

In regard to different breast cancer types, p53 is mostly overexpressed in triple negative breast cancer while it is less frequently overexpressed in hormone receptor positive breast cancers [34].

**Vascular endothelial growth factor**

Vascular endothelial growth factor (VEGF) is a cellular signal protein whose function in the body involves the formation of new blood vessels during embryonic development or after injury or to bypass blockage in a vessel, also it aids in muscle growth following exercise [35].

VEGF’s role in cancer is the promotion of vascular leakage, allowing more blood vessels to reach the tumor than normal cells [36].

In Breast Cancer, elevated levels of VEGF leads to worse clinical outcome [37,38]. However, VEGF is a key player in triple negative breast cancer, where it helps the cancer to vascularize and proliferate, as such it is an important biomarker for prognosis and predictive behavior for this particular type of cancer [39,40].

**Carcinoembryonic antigen**

Carcinoembryonic antigen (CEA) is the first ever cancer biomarker discovered in 1965 [41], since then it has fallen out as a cancer biomarker due to its poor specificity in the case of breast cancer [42]. It also stopped being used in metastatic breast cancer due to the large number of false positives it gives [43]. This is despite the fact that recent meta-analysis have shown particular benefit for CEA as a good prognostic biomarker for poorer disease free and overall survival [44], as well as the possibility of CEA to be a good predictive for metastatic behavior of early breast cancer regardless of subtype [45]. Also, recent findings shows that CEA is higher in patients with a history of breast cancer in the family than in history without breast cancer [46].

**Conclusion**

Salivary protein breast cancer biomarker has benefitted greatly from the advancement of detection techniques. However, much remains to be tested; the biggest issues remain is the lack of large scale research to grant legitimacy for the use of saliva as a medium for detection of breast cancer, and a standardization of saliva collection in order to minimize the matrix effect of the salivary matrix. Despite the lack of data regarding salivary protein biomarker and breast cancer, what we understand of these proteins now is sufficient to form the basis of future research and further development of the techniques for their detection.


