Breast cancer tumor markers

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Abstract
Breast cancer is one of the most common cancer in women, presented with a lump and skin colour change, is one of the commonest cancer in women. A wide variety of tumor markers viz. uPA, BRCA1 & 2, Ki76, ER, PR, HER etc. are used for establishing carcinoma of breast but none being a proper diagnostic tool for early detection, though each one has its own prognostic value. Though proposed but only few are used in clinical practice, research studies are still ongoing to identify and establish new biochemical parameters/markers that can be of used not only in advanced disease, but also in early stages of the diagnostic workup of breast cancer.

Key words
Biomarkers, Carcinoma, Tumor

Introduction
Tumor biomarkers are substances which show up or are elevated in blood, urine or tumor. These substances can be hormone, proteins, peptides etc. Tumor markers can be specific or non-specific, making it useful in detection, diagnosis and prognosis of cancer. Breast cancer, a heterogeneous disease, is one of the most common cancers in women, after skin cancer, representing about 16% of total female cancers. Breast cancer is less evident in women under 40 years. Breast cancer also known as malignant breast neoplasm is tumor originating from the inner lining of milk ducts or the lobules that supply the ducts with milk. Depending on which it is known as ductal carcinomas and lobular carcinoma respectively. Breast cancer is generally presented with lump in breast, nipple change or discharge and skin contour changes. Though a wide range of tumor markers have been identified for breast cancer but lack of sensitivity and specificity for early diagnosis is main disadvantage. Consequently, the available markers are of no value in either screening or diagnosing early breast cancer. Sometimes, these tumor markers are helpful but have to be used together with other tests such as biopsy, mammograms, and ultrasound and breast MRI’s. These tests are powerful, highly sensitive and specific to detect breast cancer in early stages even before symptoms are manifested [1]. It is specifically to be noted that tumor marker tests alone cannot provide enough or rather no information to screen/diagnose breast cancer. The main aim of this review is to have a brief insight of all the existing conventional and emerging biomarkers, and presenting the most effective biomarkers for breast cancer, as treatment of breast cancer is possible only if it is detected in early stages.

Markers
There is wide range of tumor markers but none being effective.
Conventional tumor markers: These include: uPA, PAI-1, CA 15-3, CA 27.29, CEA, ER, PR, HER2, BRCA1, BRCA2 and Ki67.

Emerging tumor markers: Includes: p53 protein, bcl-2, ARF, TBX2/3, Cyclin D, Cyclin E, VEGF, hTERT DNA, Glycan biomarkers, Stem cell markers, topoisomerase IIa, Serum autoantibodies, PPIA, PPRDX2, FKBP52 and Micro RNA’s.

**Urokinase plasminogen activator (upa); plasminogen activator inhibitor (pai-1)**

uPA, produced as proprotein i.e. pro uPA, which is cleaved to form a 53 Kda serine protease. Major function of this protease is to cleave plasminogen to form active protease plasmin and is involved in tissue remodelling, inflammation, fertilization, and embryogenesis and tumor invasion. PAI-1 is a glycoprotein, a serine protease inhibitor- binds to tPA and uPA, prevents cleavage of plasminogen \(^2\). PAI-1 has a role in cancer migration, invasion, inflammation, neutrophil recruitment, proliferation of smooth muscle cells and obesity \(^3\). uPAR is a cell surface receptor for uPA and is required for endocytosis of uPA-PAI-1 complexes and uPA activation. In breast cancer over expression of uPAR, PAI-1 and uPA attributes to decreased survival. UPA and PAI-1 has poor prognostic indicators for member of cancers, as they have been shown to be elevated in cancers other than that of breast \(^4\).

**CA 15-3; CA 27.29 and CEA**

*Increased CA 15-3 in serum is reported* in about 10% of patients in early breast cancer and is subsequently elevated as the disease progresses. An increase of CA 15-3 five to ten times above normal upper limit can predicts breast cancer; however a low value cannot exclude metastasis \(^5, 6\), making this more of prognostic marker rather than diagnostic marker.

**Ki67**

ki67 antigen is used to evaluate the proliferative activity of breast cancer, though its role in prognosis is still unclear it can be used as a marker for this. \(^7\) ki67 reacts with nuclear non histone protein in all active phases of cell cycle. The presence of ki67 in all proliferating cells makes it a weak marker for diagnosis \(^7, 8\).

**ER, PR, HER**

ER is probably the most powerful diagnostic and prognostic marker for breast cancer. About 60% of women below 50 years has been reported with ER positive and is increased up to 80% in women’s above 50 years. Oestrogen hormone is an established risk factor for breast cancer, which exerts its effect via oestrogen receptors (ER) - a nuclear protein. Two types of ER is seen ERa and ERb; ERa is expressed in upto 70% of all breast cancer. ER is an important mediator of carcinogenesis, inhibition of this can lead to the breast cancer therapy \(^9, 10\). PR is controlled by oestrogen therefore it shows the indication of proper functioning of ER pathway and thus PR assay can be used for prognostic purposes. \(^11\) HER2- Human epidermal growth factor 2 oncogene encodes epidermal growth factor receptor (EGFR). HER2 is important in cell differentiation, adhesion and motility. In nearly 20% of breast cancer, overexpression of this gene leads to abnormal high level of glycoprotein. HER2 positivity is associated with number of tumors and cell growth owing to its low diagnostic and prognostic value \(^11\).

p53 tumor suppressor gene products are one the major mechanism to control cancer. Inactivation of this gene leads to overexpression of p53 protein. Though overexpression is seen highly in breast cancer but is also associated with other tumors making it a poor diagnostic and prognostic marker \(^12\).

**bcl2**

Cancer cell does not undergo apoptosis. Bcl2 protein regulates the apoptosis in normal cellular pathway i.e. antiapoptotic pathway. Increase in bcl2 leads to cell survival. Bcl2 overexpression is reported in many tumors and lymphomas. In breast cancer it is increased due to oestrogen, and is less evident in other tumors \(^13\).
ARF
p14 ARF protein blocks cell cycle at G1 and G2. It acts with correlation with p53, and inhibits growth of cancerous cells by activating p53 \[^9\].

TBX2/3
TBOX protein 2 and 3 affects dimerization and DNA binding, both generally affect tumor development through down regulation of ARF p53. TBX2/3 has been overexpressed in breast cancer \[^9\].

Cyclin D1 and Cyclin E
Cyclin D1 is associated with G1 phase of cell cycle and is synthesised in response to growth factors. Cyclin D1 is overexpressed in many tumors and is detected in half of the breast cancer.

Cyclin E is the limiting factor for G1 phase progression and S phase entry. Overexpression is seen in about 40% of breast cancer \[^9\].

BRCA1 and BRCA2
Mutation in genetic level of BRCA1 and BRCA2 genes are strong evidence of breast cancer with nearly 40-80% chance of developing cancer. Genetic testing of BRCA mutation is one of the powerful tools for predicting Breast cancer \[^14\].

VEGF
VEGF is associated with angiogenesis in breast cancer and is required for tumor growth and metastasis. VEGF is reported to be overexpressed in breast cancer \[^9\].

hTERT DNA
Human telomerase reverse transcriptase (hTERT) underlies cancer cell immortalization, and the expression of hTERT is regulated strictly at the gene transcription \[^13\]. Circulating human telomerase reverse transcriptase hTRET DNA is one of the better markers for diagnosis of breast cancer \[^15\].

Glycan biomarkers
It’s still under research; certain links with cancer and altered protein glycosylation have been reported. Once established this can be a highly sensitive method for detecting early breast cancer \[^16\].

CD44 and CD24
Stem cell CD44 and CD24 evaluation in colon, breast and prostate cancer can be a candidate for breast cancer diagnosis and prognosis \[^17, 18\].

Micro RNA
Recently the miRNA has been shown to be deregulated in human breast cancer leading to genomic changes. Evaluation of this can be a better candidate for breast cancer marker \[^10\].

Conclusion
Though a clear biomarker for carcinoma of breast has not yet found but assessing certain parameters like uPA, BRCA1 &2, Ki76 and ER, PR, HER can give an indication of breast cancer, though a negative value or normal values doesn’t rule out the possibilities for the breast cancer, so as for detecting early breast cancer, some unconventional markers needed to be assayed such as micro RNAs. hTERT gene, and glycan biomarkers. These though still under research can give a proper
indication of breast cancer. However research is still needed to establish a proper protocol for detection of early breast cancer.

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