



ASSOCIATION OF VASCULAR ENDOTHELIAL GROWTH FACTORS WITH BREAST CANCER:



A REVIEW

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Abstract:

Breast Cancer is the leading cancer diagnosed in women aged 20 - 45 with risk factors including breast density, age and family history. Vascular endothelial growth factors is usually referred as VEGF family, which comprises of VEGFA, VEGFB, VEGFC, VEGFD and placental growth factor. The human VEGF gene is localized in chromosome 6p21.3 and it is organized as eight exons separated by seven introns. The VEGF and VEGFR family system is responsible for regulation of both the vascular and lymphatic endothelium and also plays a key role in tumor formation through promotion of both angiogenesis and lymphangiogenesis. A humble attempt is made to find the association between vascular endothelial growth factors and breast cancer through this review. A literature search was conducted in Pubmed for articles published in the English language using combinations of the following terms: Vascular endothelial growth factors, breast cancer and tumor.

Keywords: Vascular endothelial growth factors, breast cancer, tumor.

Introduction:

Vascular endothelial growth factors (VEGF) is usually referred as VEGF family, which comprises of VEGFA, VEGFB, VEGFC, VEGFD and placental growth factor (PGF). The human VEGF gene is localized in chromosome 6p21.3 and it is organized as eight exons separated by seven introns [1-3]. Alternative exon splicing was initially shown to result in the generation of four different isoforms (VEGFA121, VEGFA165, VEGFA189 and VEGFA206), having 121, 165, 189 and 206 amino acids respectively, after signal sequence cleavage [2, 3] the predominant isoform, lacks the residues encoded by exon 6, whereas VEGFA121 lacks the residues encoded by exons 6 and 7. VEGF121 lacks heparin binding activity because of its acidic property and is free to diffuse into the extracellular space. Due to its bioactivity and biological potency, VEGF165 is the predominant isoform of VEGF [4]. Although VEGF165 is also secreted, a significant fraction remains bound to the cell surface and extracellular matrix (ECM). VEGF189 and VEGF206 are mostly sequestered in the ECM because of their high binding affinity for heparin [5, 6]. Less frequent splice variants also have been reported, such as VEGFA145 and VEGFA183 [7]. The VEGF and VEGFR family system is responsible for regulation of both the vascular and lymphatic endothelium and also plays a key role in tumor formation through promotion of both angiogenesis and lymphangiogenesis.

Role of VEGF in Breast Cancer:

Several lines of evidence implicate the importance of VEGFA in Breast Cancer (BC) [8]. Patients with loco regional ductal cancers have elevated serum VEGFA concentrations in comparison with women with benign breast tumors. The highest concentrations of serum VEGFA were found in metastatic BC, in particular among patients who did not receive cancer therapy for metastatic disease [9]. A significant correlation between VEGFA concentration and microvessel density (MVD) has been reported [10]. A study in 574 node-negative BC patients showed that high VEGFA levels in tumor tissue were associated with larger tumor size, older age and negative progesterone receptor (PR). Tissue microarray of 642 BCs demonstrated that high levels of VEGFA and its receptors- VEGFR-1, VEGFR-2 and NRP-1 were significantly associated with poor survival. This study also demonstrated the correlation between the expression of VEGFA and its receptors within tumor cells, supporting an autocrine and paracrine function of VEGFA [11]. In a study of 50 invasive ductal carcinomas, expressions of VEGFA protein and mRNA were correlated with tumor size, lymph node metastasis and TNM staging. The MVD counts were correlated with the expression of VEGFA and axillary lymph node metastasis [12]. Cox analysis revealed that intratumoral VEGFA was an independent prognostic factor for node-negative BC [13]. High VEGFA and low soluble VEGFR-1 levels (an intrinsic negative counterpart of VEGFA) were significantly associated with poor prognosis [14, 15]. Randomized study of tamoxifen adjuvant treatment in postmenopausal patients revealed that the patients with positive VEGFA had no response to tamoxifen [16]. In advanced BC patients receiving bevacizumab and vinorelbine who had lower baseline plasma VEGFA before treatment had longer time to progress (9.3 months vs. 3.7 months) than those with higher plasma VEGFA [17].

Fibrocystic lesions with the highest vascular density are associated with a greater risk of BC [18]. Histopathologically aggressive ductal carcinoma-in-situ lesions demonstrate higher (MVD) and increased VEGF expression [19, 20]. Numerous studies have indicated that intratumoral VEGF expression is significantly correlated with MVD and poor prognosis in a variety of human solid cancers including BC, brain tumors, head and neck cancer and gastrointestinal cancer [21]. The prognostic value of VEGF has been confirmed not only in immune histochemical studies but also in other studies using enzyme-linked immunosorbent assay (ELISA) and Northern blotting. In most clinical studies that examined the prognostic value of VEGF in primary BC, intratumoral VEGF expression was a significant marker of poor prognosis in both node-negative and node-positive subgroups [22]. Carriers of the C936T allele in the VEGF gene were more frequent among controls (29.4%) than among a cohort of BC patients (17.6%), implying a protective effect for carriers of the variant polymorphism [23]. A recent report also demonstrated that this polymorphism is associated with an improved metastasis-free survival time in patients that have low-grade BC [24]. Increased VEGF expression is associated with impaired response to tamoxifen or chemotherapy in patients with advanced BC [25]. Some studies have shown that VEGF status is a significant predictor of relapse in primary BC patients receiving adjuvant endocrine treatment with tamoxifen [26, 27]. These clinical data appear compatible with the results of

experimental studies in which transfection of hormone-sensitive tumors with the VEGF121 gene rendered them hormone refractory, and antiangiogenesis therapy achieved a combined effect with hormone therapy in animal experiments [28]. A microenvironment rich in VEGF might facilitate up-regulation of growth factor production and intracrine hormone synthesis, although little is known about the mechanism involved. Similarly, VEGF expression level in the tumor tissue was a significant predictor of relapse-free survival and overall survival in node-negative BC patients treated with locoregional radiotherapy [29]. VEGFR-1 antagonists and a tyrosine kinase inhibitor of VEGFR-2 reverted radiation refractory tumor models to a radiation-sensitive phenotype. These findings suggest that the high VEGF expression might define a radio-resistant phenotype [30].

The above review supports the crucial role of VEGF in the transition from benign to malignant breast disease and BC aggressiveness and it should be further investigated in the future prospective studies in different ethnic groups to get more accurate results.

Abbreviations:

VEGF	:	Vascular endothelial growth factors
PGF	:	Placental growth factor
ECM	:	Extracellular matrix
BC	:	Breast Cancer
PR	:	Progesterone receptor
MVD	:	Micro vessel density

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