

Visfatin and its Role in Breast Cancer

Robab Sheikhpour

Hematology and Oncology Research Center, Shahid Sadoughi University of Medical Sciences, Yazd, Iran

Abstract

Breast cancer, the most common cancer in women, is a leading cause of cancer death among women worldwide. Obesity is associated with increased risk of breast cancer. Discovery of biomarkers that can be used for diagnosis, prognosis, and predictors of breast cancer is very important. Visfatin, a new adipokine found in visceral fat, plays a main role in metabolic and stress responses. There is an association between elevated expression of visfatin with malignant behavior and adverse prognosis in breast cancer. Visfatin promotes malignancy via signaling pathways that include Ras, Raf, MEK1/2, ERK, phosphoinositide 3-kinase (PI3K), Akt, and NF- κ B. Visfatin up-regulates G1-S phase cell cycle progression through up-regulation of mRNA levels of cyclin D1 and CDK2. Visfatin plays a main role in metastasis and synthesis of genes that play a significant role in tumor-related angiogenesis such as vascular endothelial growth factor, progression and tumor invasion such as matrix metalloproteinases in cancer. Visfatin may increase breast cancer cell growth and metastasis capability through c-Abl and STAT3 activation (two oncoproteins). According to the results of these studies, visfatin expression appears to be associated with virulent behavior and its inhibition may be an effective treatment for breast cancer patients.

Keywords: Breast cancer, Visfatin, NF- κ B

Introduction

Cancer is one of the most fatal diseases in humans with an annual mortality rate of 30000 individuals in Iran.¹ Breast cancer (BC) is the most common cancer in women and a leading cause of cancer deaths among women worldwide.^{1,2} Of note, the incidence of BC is increasing.³ Several factors that include cytokines, metabolic parameters, environmental factors, reproductive history, and

heredity are important in BC development.⁴⁻⁶ Currently, there are no adequate serum biomarkers for early BC detection. Therefore, the discovery of biomarkers useful for diagnosis, prognosis, predictive, and therapeutic targets for BC is very important.⁷ Visfatin, as a pre-B cell enhancing factor (PBEF),⁸ is a new adipokine found in the visceral fat that plays a main role in stress responses and cellular energy

♦Corresponding Author:

Robab Sheikhpour, PhD
Hematology and Oncology
Research Center, Shahid
Sadoughi University of Medical
Sciences, Yazd, Iran
Tel: +98 9131522462
Email: r.sheikhpour@yahoo.com

Table 1. Relation between visfatin and obesity.

Study	Results	Reference No.
Berndt et al. (2005)	Visfatin plasma concentration had a positive correlation with visceral visfatin mRNA expression and body mass index (BMI).	33
Pagano et al. (2006)	Visfatin was associated with BMI in the investigated disease.	28
Zahorska-Markiewicz et al. (2007)	Serum concentration of visfatin was significantly higher in obese women when compared to controls.	34
Shea et al. (2007)	An association existed between visfatin and the investigated disease with obesity.	35
Kovacikova et al. (2008)	Visfatin was associated with BMI.	36
Malavazos et al. (2008)	BMI >30 in obese patients was associated with disease.	37
Nakajima et al. (2010)	Visfatin levels increased with obesity.	38
Kim et al. (2010)	The level of visfatin increased with obesity.	8
Dalamaga et al. (2012)	One parameter that determined serum visfatin level in PBC cases was BMI.	7
Romacho et al. (2013)	Increased circulating levels of visfatin were reported in obese patients.	12
Assiri et al. (2015)	A positive correlation existed between visfatin and BMI.	9
El-Benhawy et al. (2015)	A high level of serum visfatin was independent of BMI.	39
Nourbakhsh et al. (2015)	Visfatin levels increased in obese people and were associated with insulin resistance (IR).	40
Hung et al. (2016)	High serum visfatin levels had a significant association with BMI.	13
Liang et al. (2016)	Elevated visfatin serum levels increased weight.	41

metabolism as nicotinamide phosphoribosyl transferase (Nampt).⁷ There is an association between elevated expression of visfatin with malignant behavior and adverse prognosis in BC.⁹ Few studies have evaluated the relationship between visfatin levels and BC, and its molecular basis remains poorly understood. Hence, this review aims to evaluate the role of visfatin in BC patients.

Visfatin

The visfatin gene has been placed on the long arm of chromosome 7 (7q22.2).¹⁰ Visfatin was discovered in adipocytes by Chang et al. in 2005.¹¹ Visfatin, a multifaceted molecule, is known as a 52 kDa adipocytokine.^{12,13} It also acts as a pro-inflammatory cytokine¹⁴ and inflammation increases visfatin synthesis.¹² Visfatin is ubiquitously generated in many tissues and several cells such as synovial fibroblasts,¹⁵ articular chondrocytes,¹⁶ or monocytes.¹⁷ Pro-inflammatory cytokines that include tumor necrosis factor alpha (TNF α) or interleukin (IL)-1 β , by lipopolysaccharide (LPS) and dexamethasone regulate expression of visfatin.¹⁸⁻²⁰ It exists in both intracellular and

extracellular states.¹³ The action of intracellular visfatin as a rate-limiting enzyme is the biosynthesis of nicotinamide adenine dinucleotide (NAD). NAD⁺ is a ubiquitous coenzyme that performs an important role in redox reactions and carries electrons from one reaction to another.²¹ It receives electrons from other molecules and becomes NADH. Therefore, it seems that Nampt/visfatin as a regulator of NAD⁺ metabolism can play a main role in the control of basic processes.²¹ Visfatin release outside of cells shows dual roles in enzyme-like activity on extracellular NAD formation and cytokine-like activity through a putative receptor-mediated pathway.⁴ In both the intracellular and extracellular states it can be involved in tumor development and cancer progression.¹³

Contradictions exist regarding the insulin mimetic effect and protective role of visfatin against development of insulin resistance (IR).²² Some studies report that visfatin can bind to insulin receptors in a place separate from insulin and act as an insulin mimetic, increasing glucose uptake and metabolism in myocytes and adipocytes, and block the gluconeogenesis

pathway. These functions of visfatin cause hypoglycemia. Studies report an association between visfatin and insulin levels,^{23,24} however others report no such association.²⁵⁻²⁸

Relation between obesity and visfatin in breast cancer (BC)

Visfatin may be a promising predictor for obesity, IR, and metabolic syndrome.²⁹⁻³² Table 1 shows the relation between obesity and visfatin.

Obesity and related metabolic alterations are associated with increased risk for cancer, particularly BC.⁸ Obese postmenopausal women compared to non-obese women had a higher overall risk of BC.⁴² Although the relation between obesity and BC is complex and not understood,⁴³ a number of parameters can contribute to this process.

Role of visfatin in breast cancer (BC)

Visfatin is over-expressed in many inflammatory diseases that include atherosclerosis, rheumatoid arthritis, osteoarthritis (OA), metabolic syndrome, and sepsis.^{15-17,44,45} Visfatin also promotes malignancy and is related to a worse clinical prognosis.⁴⁶ Elevated expression of visfatin is associated with adverse prognosis of BC.⁴⁷ Therefore, visfatin may show a new link with BC pathophysiology.^{47,48} Table 2 lists studies that have reported the role of visfatin in BC.

Signaling pathways of visfatin

Although studies have shown the role of serum visfatin as a biomarker¹³ in BC, the relation between visfatin levels and BC risk is unclear. The molecular basis for this link is not understood. Visfatin up-regulates G1-S phase cell cycle progression through up-regulation of mRNA levels of cyclin D1 and CDK2.⁸ Visfatin has been shown to increase expressions of matrix metalloproteinases MMP2 and MMP9 mRNA, and vascular endothelial growth factor (VEGF).⁵⁴ Therefore, it seems that visfatin can play a main role in metastasis, as well as synthesis of genes that play a significant role in tumor-related angiogenesis (VEGF as a pro-angiogenic protein),⁷ and

progression and tumor invasion (MMP in BC).⁵²

Park et al. have reported that visfatin increased NF- κ B p65 and Notch1 levels. NF- κ B p65 is a positive regulator of Notch1 stimulated by visfatin in BC cells.⁵² Notch1 was inhibited via NF- κ B signaling pathway inhibition. It has not been determined if NF- κ B directly or indirectly mediates the stimulation of Notch1 expression.⁵² Visfatin, via Notch1 signaling activation, promoted BC. Reduction of visfatin and Notch1 suppressed cell proliferation and stimulated apoptosis.⁵³ However, over-expression of active Notch1 increased tumor formation in murine mammary tissue.⁵³ Therefore, increased expression of Notch1 has been shown to correlate with poor overall survival in BC patients. It seems that Notch1 can be as a prognostic biomarker for the pathogenesis of BC and a new therapeutic target.⁵³

Hung et al. reported that extracellular visfatin increased BC cell growth and metastasis capability through c-Abl and STAT3 activation (two important oncoproteins).¹³ They reported that imatinib, a tyrosine kinase inhibitor, and Stattic as STAT3 inhibitor might be useful to treat BC patients who have elevated levels of visfatin.¹³

It has been shown that numerous signaling pathways such as PI3K,⁵⁴ p38 MAPK,⁵⁵

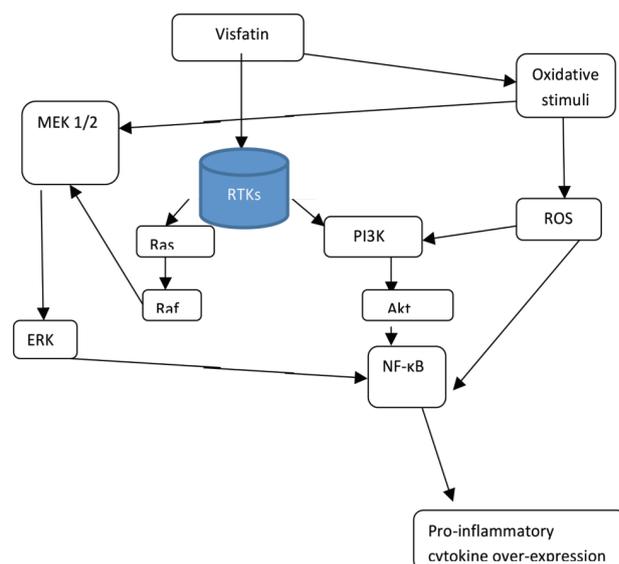


Figure 1. Visfatin signaling pathway which leads to pro-inflammatory cytokine over-expression by oxidative stress and receptor tyrosine kinases (RTKs).⁵⁹

Table 2. Different studies that discussed the role of visfatin in breast cancer (BC).

Study	Results	Reference No.
Folgueira et al. (2005)	Visfatin was expressed in human BC tissue.	49
Yonezawa et al. (2006)	Visfatin was expressed in MCF-7 BC cells.	50
Kim e al. (2010)	Visfatin motivated proliferation of MCF-7 human BC cells.	8
Moschen et al. (2010)	The level of visfatin as a pro-inflammatory adipocytokine increased in various inflammatory conditions and was correlated with the degree of inflammation.	51
Dalamaga et al. (2011)	A significantly greater mean level of visfatin existed in BC patients compared to the control group.	4
Lee et al. (2011)	Visfatin highly expressed in human BC cells both <i>in vitro</i> and <i>in vivo</i> .	47
Kim et al. (2012)	Visfatin increased the invasion of MDA-MB-231 human BC cells.	52
Dalamaga et al. (2012)	Higher visfatin expression in BC tissue correlated with increased malignant cancer behavior.	7
Park et al. (2014)	Visfatin stimulated BC proliferation and invasion.	53
Li et al. (2014)	Women with high serum levels of visfatin had significantly shorter overall survival compared to those with low serum levels of visfatin.	3
Assiri et al. (2015)	Visfatin increased lymph node metastasis in postmenopausal BC.	9
El-Benhawy et al. (2015)	High levels of serum visfatin were seen in BC patients compared to controls.	39
Hung et al. (2016)	High serum visfatin levels were associated with advanced tumor stage, increased tumor size, lymph node metastasis, and poor survival in BC patients.	13

ERK1/2,⁵⁶ JNK,⁵⁷ STAT3,⁵⁸ AKT,⁵⁹ NF- κ B⁵² can contribute to activation of downstream target gene transcription induced by visfatin. Some studies reported that STAT3 as a downstream target gene transcription for visfatin does not contribute to cancer progression.⁵⁹ Figure 1 shows visfatin signaling pathway which leads to pro-inflammatory cytokine over-expression by oxidative stress and receptor tyrosine kinases (RTKs).⁵⁹

Visfatin and oxidative stress

Several metabolic pathways such as aerobic metabolism in the mitochondrial respiratory chain can generate reactive oxygen species (ROS) and subsequently initiate and progress several types of cancers.⁶⁰ Reactive oxygen species can affect many signals such as growth factors and mitogenic pathways.⁶⁰ The role of visfatin in inflammation, progression, and oxidative stress is emerging.⁵⁹ Visfatin is a strong marker for inflammation and dysfunction.²¹

Visfatin can increase expressions of inflammatory adhesion molecules including

vascular cell adhesion molecule (VCAM-1) and intercellular adhesion molecule 1(ICAM-1) in vascular endothelial cells in a ROS-dependent manner via NF- κ B signaling.⁶¹ Therefore, these studies have shown that visfatin increased oxidative stress and inflammation through NF- κ B activation and other stress-related signaling pathways in a cell-type dependent manner.⁶¹ Visfatin, by stimulation of PI3K, caused activation of NF- κ B and pro-inflammatory answer.^{59,62} Other studies have reported that visfatin activates members of the MAPK signaling pathway involved in stress responses, including ERK⁶² and p38.^{61,62} Marseglia et al. have reported that nicotinamide phosphoribosyltransferase (NAMPT) / visfatin can act as a cytokine, including granulocyte-macrophage colony-stimulating factor (GM-CSF), IL-2, IL-1 β , IL-6, and IL-13 which up-regulate in several acute and chronic inflammatory neonatal diseases.²¹ However Buldac reported that visfatin increased cytokine production via increased intracellular ROS levels.⁶⁰

Lipid peroxidation increased in the presence of visfatin.⁶⁰ Visfatin could interact with protein

mediated oxidative stress and inflammation including ND1, ferritin light chain, and interferon-induced transmembrane protein 3 (IFITM3) which led to increased levels of ROS.⁶³

Conclusion

According to the result of these studies, it seems that visfatin expression is associated with virulent behavior. Its inhibition may be effective therapy for breast cancer patients.

Conflict of Interest

No conflict of interest is declared.

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