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Immunohistochemical Study of the Angiogenetic Network of VEGF, HIF1 α , VEGFR-2 and Endothelial Nitric Oxide Synthase (eNOS) in Human Breast Cancer

Maria Kafousi • Thomas Vrekoussis • Eleftheria Tsentelierou • Kitty Pavlakis • Iordanis Navrozoglou • Vassilios Dousias • Elias Sanidas • Dimitrios Tsiftsis • Vassilios Georgoulias • Efstathios N. Stathopoulos

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Abstract

Background The role of Nitric Oxide (NO) in angiogenesis has not been fully clarified yet. A dual role for NO, either inductive or inhibitory, has been proposed on the basis of different effects that high or low concentrations of NO may exert on the angiogenic process. Additionally, it has been referred that NO may induce VEGF production, while VEGF may induce NO production via up-regulation of the endothelial nitric oxide synthase

Vassilios Georgoulias and Efstathios N. Stathopoulos share senior authorship

M. Kafousi · T. Vrekoussis · E. Tsentelierou · E. N. Stathopoulos (⊠) Department of Pathology, Medical School of the University of Crete, Voutes, 70003 Heraklion, Crete, Greece e-mail: stath@med.uoc.gr

T. Vrekoussis · I. Navrozoglou · V. Dousias Breast Unit, Department of Obstetrics and Gynecology, Medical School, University of Ioannina, Ioannina, Epirus, Greece

K. Pavlakis Department of Pathology, Medical School, University of Athens, Athens, Greece

E. Sanidas · D. Tsiftsis Department of Surgical Oncology, Medical School, University of Crete, Heraklion, Crete, Greece

V. Georgoulias Department of Medical Oncology, Medical School, University of Crete, Heraklion, Crete, Greece (eNOS), the two pathways being reverse. The aim of the current study was to investigate the expression of key molecules involved in these opposite pathways in primary breast cancer.

Methods Representative tumor samples from 242 patients with early-stage breast cancer (invasive ductal breast carcinomas) were investigated for the expression of VEGF, VEGFR-2, HIF1 α , iNOS, and eNOS using immunohistochemistry.

Results Endothelial NOS was found in 159 cases, VEGF in 131 cases, HIF-1 α in 139 cases, VEGFR2 in 185 cases and inducible NOS (iNOS) in 22 cases. There was a significant correlation between the expression of VEGF and VEGFR-2, eNOS and VEGF, eNOS and VEGFR-2, eNOS and HIF1 α . No statistically significant correlation was found between iNOS and the rest of the studied molecules.

Conclusions In breast cancer cases, the major molecules regulating NO and VEGF production can be co-expressed in the individual carcinomas implying a possibility for the relevant pathways to be active; however appropriate functional experiments remain to be conducted to prove such a hypothesis

Keywords Angiogenesis · Breast Cancer · eNOS · VEGF

Background

Angiogenesis is essential for a tumor to grow and disseminate [1]. Several factors have been identified as inducers or inhibitors of angiogenesis. Among them, nitric oxide (NO) seems to be enrolled in modulating either positively or negatively the angiogenic process [2].

NO is produced by nitric oxide synthases (NOS), enzymes that catalyze the transformation of L-arginine to L-citruline. To date two categories of NOS have been recognized: a) the calcium-dependent constitutive isoforms that are found in neural tissue (neuronal NOS or nNOS) and in endothelial cells (endothelial NOS or eNOS) and b) the calcium-independent inducible NOS or iNOS, found primarily in mesenchymal or parenchymal tissues of several organs [2].

The role of NO in tumor angiogenesis has not been entirely clarified yet. Low NO concentrations seem to induce tumor angiogenesis, while high NO concentrations are inhibitory. Several studies so far have revealed that NO may induce VEGF production via the PI3K/Akt/PKB-HIF-1 α pathway [3–6]; however reports claim that NO might also act as an inhibitor of VEGF expression [7]; this contradiction has not been cleared so far. This dual and bi-directional angiogenesis regulatory mechanism becomes more complex by the interference of hypoxia. Under hypoxic conditions, there is an activation of the c-Src pathway leading to HIF-1 α up-regulation and finally to increased VEGF production [8, 9], which induces NO production via VEGFR-2 and its downstream effector c-Src [10-14]. Additionally, hypoxia seems to induce NO production [15]. On the contrary, high NO concentrations inhibit c-Src leading to HIF-1 α down-regulation, thus decreasing VEGF production [2].

The current knowledge regarding NO/NOS and their relationship with angiogenesis in breast cancer is based primarily on in vitro experiments, since there are only a few studies reporting on the relationship of NO/NOS expression in the prognosis of human tumors [16–19]. To our own best of knowledge there is no study in the literature investigating the relationship between nitric oxide synthases and other angiogenic factors or receptors on clinical specimens.

The aim of the present study was to investigate whether molecules of the above interrelating pathways are expressed in human breast cancers.

Patients, Materials and Methods

Patients and Molecular Subtyping Two hundred and 42 cases of operated early breast cancer (invasive ductal breast carcinomas) were randomly retrieved from the archives of the Pathology Department of the University General Hospital of Heraklion, Crete, Greece. All patients had undergone primary surgical treatment (breast surgery \pm axillary sentinel lymph node dissection \pm complete axillary lymph node dissection). No patient had received preoperative chemotherapy. Archival data regarding estrogen receptors (ER), progesterone receptors (PR) and Her-2 status on each patient was retrieved as well. All cases were subgrouped according to their molecular profile as previously described [20]: a) Luminal A [ER (+) or PR (+), Her-2 (-)], b) Luminal B [ER (+) or PR (+), Her-2 (+)], c) Triple negative [ER (-), PR (-), Her-2 (-)] and d) Her-2 positive [ER (-), PR (-), Her-2 (+)].

Immunohistochemistry for Detection of VEGF, VEGFR-2, eNOS, iNOS, HIF-1 α Expression Immunodetection protocol details are presented in Table 1. Briefly, formalin-fixed and paraffin-embedded tissue sections (3 µm thick) were deparaffinized in xylene and rehydrated through graded concentrations of ethanol. Antigen retrieval via microwave treatment (3 cycles of 5 min. each) followed. After incubation with the primary antibodies, positivity was revealed by using a commercially available immunodetection system (UltraVision LP, TL-125-AL, Lab Vision, CA, USA with fast red as chromogen) followed by Mayer's hematoxylin counterstaining for 3 min. Slides were then

Table 1 Basic features of the immunohistochemistry detection performed in the study

Antibody/Origin/Source	Antigen retrieval	Dilution in TBS	Incubation	Positive Control	Detection System
Anti-VEGF, mouse, MS-1467-P, Neomarkers, CA, USA	MW^1	1/50	1 h	Angiosarcoma	UltraVision LP/AP with Fast-Red (Lab Vision, CA, USA)
Anti-HIF1a, mouse, MS 1164, Thermo scientific, CA, USA	MW^1	1/20	overnight	Oral squamous cell carcinoma	
<i>Anti-VEGFR-2, mouse,</i> SC-6251, SantaCruz, CA, USA	MW^1	1/40	2 h	Human colon carcinoma	
Anti- eNOS, rabbit, RB-1711-P, Neomarkers, CA, USA	MW^2	1/100	overnight	Capillary endothelium	
Anti-iNOS, rabbit, RB-9242, Thermo scientific, CA, USA	MW^1	1/300	1 h	Lung macrophages	

¹MW (350 W, three times for 5 min. each) in 0.01 M Citrate Buffer, pH 6

 2 MW (350 W, three times for 5 min. each) in 0.001 M EDTA, pH 8

rinsed with ammonia water, washed in tap water and covered using Glycergel (Dako, CA, USA). Positive controls (according to the manufacturer's protocol) were included in the study. Sections stained with the same protocol by omitting the primary antibodies were used as negative controls.

Scoring of Immunoreactivity

The evaluation of all immunostainings was performed by two independent observers (MK and ENS) working blindly to the clinical and histopathological data, and the mean of the two recorded observations was considered as the final value. In cases with greater than 10% discordance in reading results, the final conclusion was reached in consensus [21]. Immunostaining results were evaluated in a Nikon, Eclipse, E-400 photomicroscope equipped, among others, with a $40X/0.65-\infty/0.17$ WD 0.65 objective lens.

Nuclear reactivity for HIF-1a was scored as previously reported [22], based on reaction intensity and percentage of positive cells as follows: a) staining not detected: 0, b) positive cells less than 1%: 1+, c) 1–10% cells with slight to moderate intensity or 11–50% slightly positive cells: 2+, d) 11–50% positive cells with moderate to marked staining: 3+ and e) more than 50% positive cells with moderate to marked staining: 4+.

The VEGFR2 expression was evaluated as previously described [23]: a) carcinomas with less than 10% positive cells/10 hpf (x400): 0, b) 11%–30% positive cells/10 hpf: 1, c) 31%–50% positive cells/10 hpf: 2, and d) more than 50% positive cells/10 hpf: 3 [23].

The eNOS immunostaining was evaluated as previously reported [16]. Intensity of cytoplasmic immunostaining was scored from 0 to 4+ (no staining, weak staining, moderate staining, strong staining, very strong staining, respectively). There was no need for positive tumor cells counting, since in any single case all cells were equally either positive or negative.

Regarding iNOS, there was no need for positive tumor cells counting and grading, since in any individual case all cancer cells were mostly equally either positive, with a granular cytoplasmic positivity pattern without differences in intensity from cell to cell, or negative.

VEGF staining was evaluated as previously described [24]. In brief, intensity of cytoplasmic immunostaining was scored from 0 to 3+ (no staining, weak staining, moderate staining, strong staining, respectively).

Each section was also characterised as negative (IHC score=0) or positive (IHC score>1+) according to previously mentioned reports [16, 25–27].

Statistical Analysis Potential correlations between expressions of molecules involved in angiogenesis were assessed either by evaluating the Gamma correlation co-efficient or by applying the chi-square test (when appropriate). Every association presented with p < 0.05 was considered statistically significant.

Results

Patient Clinicopathological Features All cases enrolled in this study were invasive ductal carcinoma. Among these, 146 (60.33%) were ER positive, 135 (55.79%) were PR positive and 36 (14.88%) were Her-2 positive. According to ER/PR/Her-2 expression status, cases were grouped as Luminal A (n=156, 64.46%), Luminal B (n=16, 6.61%), Her-2 expressing (n=19, 7.85%) and triple negative (n=51, 21.07%). Axillary lymph node dissection was performed on 227 cases; 138 (60.79%) were found to have one or more positive lymph nodes (Table 2).

The selected tissue sections of 200 and 42 cases of early primary invasive ductal breast carcinomas examined immunohistochemically for eNOS, VEGF, VEGFR2, HIF1 α and iNOS expression were representative of the individual tumors architectural and cytological pattern. Representative figures of immunostained slides are presented in Fig. 1.

Distribution of IHC Positivity Endothelial NOS (eNOS) positivity was cytoplasmic, mainly on tumor cells; 36 (14.88%), 80 (33.06%), 29 (11.98%) and 14 (5.79%) were scored 1+, 2+, 3+ and 4+ respectively (Table 2). VEGF was expressed in the cytoplasm of tumor cells, and 81 (33.5%), 44 (18.2%) and 6 (2.4%) were scored as weak, moderate and strong positive, respectively (Table 2). HIF1 α was found to be expressed in the nucleus of tumor cells; 82 cases (33.88%) presented as 2+, 18 (7.44%) as 3+ and 39 (16.12%) as 4+ (Table 2). VEGFR2 was expressed on tumor cells as well as on endothelial cell membranes and was found in 11 (4.55%) cases as weakly positive, in 6 cases (2.48%) as moderately positive and in 168 (69.42) cases as strongly positive (Table 2). Finally, iNOS was detected in 22 cases (9.09%) in a cytoplasmic pattern (Table 2).

Sixty five cases (26.85%) were positive for all the four molecules included in the VEGF-eNOS pathway (eNOS, VEGF, VEGFR2, HIF1 α) (Fig. 1b). Thirty nine cases were of luminal A, 7 were of luminal B, 5 were of Her-2 expressing type and 14 were triple negative cases. This distribution however was not significant (p>0.05).

Correlation Analysis eNOS, VEGF, HIF-1a, VEGFR2 and iNOS Endothelial NOS expression was detected in 104 out of 131 (79.39%) of the VEGF positive cases and in 55 out

molecules of the current study (%) n Histology 100 Invasive Ductal Carcinoma 242 Grade G1 38 15.70 G2 46.69 113 G3 91 37.60 Nodal Status (n=227) N₀ 89 39.21 N^+ 138 60.79 **Estrogen Receptors (ER)** Positive 146 60.33 Negative 96 39.67 **Progesterone Receptors (PR)** Positive 135 55.79 Negative 107 44.21 Her-2 Positive 36 14.88 Negative 206 85.12 Molecular subtypes Luminal A 156 64.46 Luminal B 16 6.61 Her-2 19 7.85 Triple Negative 51 21.07 Immunohistochemistry scores

Table 2 Major clinicopathological features and IHC scores of the

eNOS	n	(%)	HIF-1a	n	(%)
0	83	34.30	0	103	42.56
1	36	14.88	1	0	0
2	80	33.06	2	82	33.88
3	29	11.98	3	18	7.44
4	14	5.79	4	39	16.12
VEGF	n	(%)	VEGFR2	n	(%)
0	111	45.87	0	57	23.55
1	81	33.47	1	11	4.55
2	44	18.18	2	6	2.48
3	6	2.48	3	168	69.42
iNOS		n		(%)	
Positive		22		9.09	
Negative		220		90.91	

of 111 (49.55%) of the VEGF negative cases. Moreover, 104 out of 159 (65.40%) of the eNOS positive cases were found to express VEGF compared to 27 out of 83 (32.53%) of the eNOS negative cases (Table 3). The correlation between eNOS and VEGF was significant (gamma co-efficient: 0.377, $p < 10^{-6}$) (Table 4). Concurrently eNOS was detected in 133 out of 185 (71.89%) of



Fig. 1 Representative sections of breast cancer specimens, immunostained for the molecules studied (the scale bar represents 10 μ m) and schematic presentation of the sample regarding eNOS (**a**), VEGF (**b**), HIF-1a (**c**) and VEGFR2 (**d**). In sixty five cases all the molecules coexist (*marked with a dash*). Orange: Negative IHC staining, *Red*: Positive IHC staining

Table 3	Cross-tabulation	of data	according to	the	expression of	of the	molecules	involved	in the	e current study	V
			0							-	

	VEGF		HIF 1a		VEGFR2	
	Negative (n=111)	Positive (n=131)	Negative (n=103)	Positive (n=139)	Negative $(n=57)$	Positive (n=185)
eNOS negative (n=83)	56	27	44	39	31	52
eNOS positive (n=159)	55	104	59	100	26	133
iNOS negative (n=220)	104	116	95	125	55	165
iNOS positive (n=22)	7	15	8	14	2	20
	VEGF negative (n=	-111)	60	51	40	71
	VEGF positive (n=	131)	43	88	17	114
			HIF 1a negative $(n=103)$		29	74
			HIF 1a positive $(n=139)$		28	111

the VEGFR2 positive cases and in 26 out of 57 (45.61%) of the VEGFR2 negative cases, while 133 out of 159 (83.64%) of the eNOS positive cases compared to 52 out of 83 (62.65%) of the eNOS negative cases were found to express VEGFR2 (Table 3). The correlation between eNOS and VEGFR2 was significant (gamma co-efficient: 0.244, p=0.000848), as well (Table 4). Finally, eNOS was detected in 100 out of 139 (71.94%) of the HIF-1 α positive cases and in 59/103 (57.28%) of the HIF-1 α negative cases, whereas 100/159 (62.89%) of the eNOS positive cases were found to express HIF-1 α compared to 39/83 (46.99%) of the eNOS negative cases (Table 3). The correlation between eNOS and HIF1 α was also significant (gamma co-efficient: 0.130, p=0.031229) (Table 4).

Regarding VEGFR-2, its expression was found to be significantly correlated to the expression of VEGF (gamma co-efficient: 0.385, $p=10^{-6}$) (Table 4). VEGFR2 was

Table 4 Pair-wise correlations of the molecules under study

Pairwise correlations								
	Gamma correlation	Gamma correlation						
	Gamma	р						
eNOS & VEGF	0.377	$< 10^{-6*}$						
eNOS & HIF 1	0.130	0.031229*						
eNOS & VEGFR	0.244	0.000848^{*}						
VEGF & HIF 1	0.307	0.000001^{*}						
VEGF & VEGFR	0.385	0.000001^{*}						
HIF 1 & VEGFR	0.131	0.094057						
	Chi square test (χ^2), df=3	р						
iNOS & VEGF	6.135	0.10519						
iNOS & HIF 1	3.086	0.37847						
iNOS & VEGFR	2.925	0.40327						

Significant observations are marked with an asterisk

detected in 114 out of 131 (87.02%) of the VEGF positive cases and in 71 out of 111 (63.96%) of the VEGF negative cases. One hundred and 14 (61.6%) out of 185 VEGFR-2 positive cases were found expressing VEGF, compared to 17 out of 57 (29.82%) of the VEGFR-2 negative cases (Table 3). Finally a significant correlation was found between VEGF and HIF-1 α expression (Table 4).

The correlations between VEGFR2-HIF-1 α , iNOS-VEGF, iNOS-VEGFR2, iNOS-HIF1 α were not statistically significant (Table 4).

Correlation Analysis Between Clinicopathological Features and the Molecules of the Study All the clinicopathological features were examined for possible correlations with the molecules of the study. Estrogen receptor status (Table 5) and progesterone receptors status (Table 6a) was significantly correlated to eNOS expression. Indeed 72.6% and 71.1% of the ER/PR positive cases respectively, express eNOS, while only 55.2% and 58.8% of the ER/PR negative cases respectively are eNOS reactive (Table 6a).

Additionally a significant correlation was revealed between PR and HIF-1a expression (Table 5), since 49.6% of the PR positive and 67.28% of the PR negative cases express HIF-1a (Table 6b). Analysis between HIF-1a groups and different IHC scores (Table 6b) also revealed that intense HIF-1a IHC score (i.e. 4+) is significantly more common (p=0.028) in ER negative (23.9%) than in ER positive (10.9%) cases and in HER-2 positive (25%) than in HER-2 negative (14.5%) cases (p=0.004).

Taking into account the molecular subtyping, it is shown that HIF-1a expressing cases are significantly more common (p=0.036) in Luminal B (75%), HER-2 (68.42%) and triple negative (68.62%) cases, than in Luminal A (50.64%) cases (Table 7). HIF-1a, VEGF and VEGFR2 expressions were positively correlated to the histological grade (Table 5). The rest of the investigated correlations were not significant (Table 5).

Table 5 Correlations between salient clinicopathologic features and the molecules of the study. Significant results are underlined. Analysis on ER, PR, Her-2 and lymph node status has been performed by

applying chi square. Grade and tumor diameter analysis has been performed by applying gamma correlation statistic

	ER (p)	PR (p)	HER-2 (p)	Grade (p)	Tumor Diameter (p)	Lymph node status (p)
eNOS (df=4)	0.042	0.078	0.483	0.837	0.986	0.729
$VEGF_{(df=3)}$	0.362	0.842	0.559	0.001	0.942	0.763
HIF-1a _(df=3)	0.028	0.027	0.004	$\leq 10^{-3}$	0.305	0.588
$VEGFR2_{(df=3)}$	0.520	0.626	0.978	0.028	0.986	0.756
$iNOS_{(df=1)}$	0.212	0.219	0.153	0.396	0.766	0.863

Discussion

Angiogenesis is an essential process for tumor growth and dissemination. The key-role in angiogenesis is held by VEGF [28], being regulated among others by hypoxia and NO. Hypoxia and NO production are expected to be strongly associated, since it has already been shown that HIF1 α is enrolled in the NO-induced VEGF production [5, 29].

Although efforts have been made to identify and study VEGF, HIF1 α , eNOS, iNOS and VEGFR-2 and the in between them relation using in vitro models, little progress has been made in studying these molecules on human breast cancer specimens. Most of the studies so far (frequently reporting contradictory results) have focused either on NO production or on hypoxia, in the aim of

identifying significant correlations regarding clinical features such as overall or disease-free survival [18, 30–32].

To the best of our knowledge, there are no studies in humans investigating the major components of the two pathways involving VEGF-induced NO production, and vice versa, as an intergraded system of molecules, interacting or opposing or even being complementary to each other. Our work, using a statistically adequate sample (n=242 breast cancer cases), is the first one clearly demonstrating that the above molecules can co-exist in human breast cancers, since 26.85% of the cases in our sample are positive for all the relevant molecules.

We have shown statistically significant co-expression between eNOS and VEGF on the one hand, and eNOS and HIF1 α on the other; this observation indicates an important role in angiogenesis for all the above three molecules,

Table 6 A & B. Selective detailed presentation of the significant correlations. Further analysis (by a 2×2 contingency table) of eNOS expression has been performed by grouping all positive IHC scores together, as previously performed. (df. degrees of freedom)

A.							
	eNOS						
	0	1+	2+	3+	+4	p (df=4)	p (df=1)
ER							
Positive	40	21	54	22	9	0.042	0.005
Negative	43	15	26	7	5		
PR							
Positive	39	17	50	21	8	0.078	0.046
Negative	44	19	30	8	6		
B.							
		HIF-1	ı				
		0	2+	3+	4+	p (df=3)	
Her-2							
Positive		10	10	7	9	0.004	
Negative		93	72	11	30		
ER							
Positive		66	55	9	16	0.028	
Negative		37	27	9	23		
PR							
Positive		68	43	8	16	0.027	
Negative		35	39	10	23		

Table 7 Distribution of the molecules of the study among themolecular subtypes

	Luminal A	Luminal B	Her-2	Triple Negative	р
eNOS					
Positive Negative	108 48	13 3	12 7	26 25	0.056
VEGF					
Positive Negative	84 72	13 3	9 10	25 26	0.131
HIF-1a					
Positive Negative	79 77	12 4	13 6	35 16	0.036
VRGFR2					
Positive Negative	115 41	11 5	15 4	44 7	0.264
iNOS					
Positive Negative	17 139	1 15	0 19	4 47	0.428

which have already been demonstrated to participate in NO-induced VEGF production [2]. Co-expression of VEGF and HIF1 α has been previously reported [8] and it is in consistence with our results (Table 4). Additionally, VEGF-mediated NO production has been reported via eNOS upregulation by VEGFR-2 signaling [13]. The above finding is in agreement with the significant co-expression of VEGFR-2 and eNOS that was found in our series. Finally, the lack of statistical significance found in our results (Table 4), when a tentative HIF1 α and VEGFR-2 co-expression was evaluated, seems to indicate that both in human breast cancers and in vitro [2], VEGF-induced NO-production may not be -at least directly- HIF-1a-dependent.

Another interesting finding of our study is that eNOS, but not iNOS, is strongly correlated with VEGF, VEGFR-2, and HIF-1 α . iNOS is considered essential for tumor angiogenesis, since it is responsible for NO production in the stroma [33], and has been reported as a predictor of survival in axilla negative breast cancer cases [34]. Nevertheless, as our results indicate, it seems that in breast cancer, its contribution to the mechanism suggested and discussed herein is not significant.

The association of eNOS and estrogen receptors shown in our series (Tables 5 and 6a) is in concordance with previous knowledge. In the past, few reports demonstrated the current ER-eNOS association in breast cancer patients [16, 35]. Since then, it has been reported that ER induces eNOS expression [36], explaining why estrogens induce eNOS but not iNOS production in the MCF-7 breast cancer cell line [37]. Recently an association between ER α and eNOS expression has been found in healthy women as well [38]. Our findings verify previous results and thus We have, also, shown a marginally significant association of progesterone receptors and eNOS expression, being in accordance with a previous report. The effect however of progesterone upon eNOS expression is not clear yet. Progesterone has been described to oppose to estrogen by down-regulating eNOS expression [39, 40]. Such an observation might seem as discordance to our results. However the evidence regarding the progesterone effect in breast cancer is weak so far. It could be hypothesized though that as long as eNOS is expressed, the significant co-existence of progesterone receptors could be a part of a negative feedback mechanism.

In cancer, the major role of hypoxia has been established by an expression profiling study [41], in which, genes affected by hypoxia were clustered to form the "hypoxia signature". This hypoxia signature was proved to be of clinical importance, since it could significantly predict survival rates [41, 42]. The key role in this hypoxia signature is held by the HIF family molecules via which other genes and proteins are regulated [42]. In our study, we analysed the effect of HIF-1a as part of the hypoxia signature and we demonstrated that in addition to the above discussed correlations, HIF-1a is inversely related to ER and PR expression and positively related to Her-2 expression. The inverse association of HIF-1a with the hormone receptors can be justified since it has been reported that hypoxia downregulates ER and PR expression in breast cancer cell lines [43]. However to our knowledge our report is one of the few verifying that HIF-1a (demonstrated by IHC and not by expression profiling) is clinically associated with reduced ER and PR expression in breast cancer cases [44]. Few reports correlate HER-2 with HIF-1a expression as well [45, 46]. Our results are in agreement with these reports.

Finally, HIF-1a was found to be more abundant in the more aggressive breast cancer subtypes (luminal B, Her-2 expressing and triple negative). Such an observation is in concordance with a recent report associating HIF-1a expression with aggressive breast cancer and worse disease free and overall survival [44].

All the above taken together produce evidence that in human breast cancer there is, at least, a proportion of cases that can express the key-molecules needed for the regulation of VEGF and NO production. Patients within this group are likely to benefit from targeted treatments, including VEGF inhibition, such as the commercially available anti-VEGF antibody Bevacizumab [47, 48]. Additionally, HIF1 α and thus VEGF inhibition is nowadays an appealing solution. Several HIF1 α inhibitors (microtubule modifiers, HSP90 inhibitors, topoisomerase inhibitors and other molecules) have been identified; however most of them are not as specific as required in order to be clinically exploited [49, 50].

However, interactions between NO, hypoxia factors and VEGF might be more complex than hypothesized in the present work, since a) HIF-1 α is regulated by oncogenic signaling pathways which are cell-type specific [49] and b) hypoxia elements for HIF-2 have already been identified on the eNOS promoter region [51]. Consequently, combined functional, molecular and immunohistochemical studies are necessary to further clarify the relationship between hypoxia, NO pathways and angiogenesis.

Conclusions

In conclusion, we have shown that in breast cancer cases, the major molecules regulating NO and VEGF production can co-exist, implying-but not proving-functional activity of the relevant pathways. However appropriate experiments, such as functional assays, remain to be conducted to strengthen such a hypothesis. This finding could be of clinical importance for designing future anti-angiogenic interventions. More extended studies are certainly required in order for the above concept to be applied to breast cancer therapy.

Conflict of Interest The authors declare that they have no conflict of interest.

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