

# Candidate pathways linking inducible nitric oxide synthase to a basal-like transcription pattern and tumor progression in human breast cancer

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**I**nducible nitric oxide synthase (NOS2) is an inflammation responsive enzyme (EC 1.14.13.39) that is induced during acute and chronic inflammation and tissue injury as part of the host defense and wound healing process. NOS2 upregulation leads to increased nitric oxide (NO) production, the means by which this enzyme can initiate NO-dependent signal transduction, influence the redox state of cells and induce modifications of proteins, lipids and DNA. Aberrant expression of NOS2 has been observed in many types of human tumors. In breast cancer, increased NOS2 is associated with markers of poor outcome and decreased survival. Growth factor and cytokine signaling, tissue remodeling, NFκB activation and hypoxia are candidate mechanisms that induce NOS2 in tumor epithelial and tumor-infiltrating cells. NOS2 induction will trigger the release of variable amounts of NO into the tumor microenvironment and can activate oncogenic pathways, including the Akt, epidermal growth factor receptor and c-Myc signaling pathways, and stimulate tumor microvascularization. Constitutively increased NO levels may also select for mutant p53 cells to overcome the tumor suppressor function of NO-activated wild-type p53. More recent findings suggest that NO induces stem cell-like tumor characteristics in breast cancer. In this review, we will discuss the effects of NO in tumor biology and disease progression with an emphasis on breast cancer, and will examine the mechanisms that link increased NO

to a basal-like transcription pattern in human breast tumors and poor disease outcome.

## Introduction

Stromal gene signatures predict resistance to therapy and clinical outcome in breast cancer.<sup>1,2</sup> Other studies revealed that transcriptional signatures reminiscent of the host wound healing response are induced in breast tumors and other epithelial cancers, and are associated with cancer progression and poor outcome.<sup>3,4</sup> Together, these findings highlight the importance of stromal biology and the process of tumor-associated wound healing in cancer biology, disease aggressiveness and therapy response. One of the genes with a key function in wound healing is NOS2.<sup>5</sup> In tumor biology, aberrant NOS2 induction may occur to facilitate tissue remodeling and to stimulate neovascularization.<sup>6-9</sup> In this function, NOS2 may synergize with the endothelial isoform, eNOS.<sup>10,11</sup> NO regulates metalloproteinase activity,<sup>9,12</sup> crucial players in tissue remodeling and oncogenic pathways,<sup>13-15</sup> and induces angiogenesis by mechanisms that involve activation of vascular endothelial growth factor (VEGF),<sup>7,8,16</sup> and inhibition of thrombospondin-1.<sup>17</sup> Moreover, recent findings indicate that NO induces stem cell-like tumor characteristics, including upregulation of c-Myc and CD44, and a basal-like phenotype in breast cancer<sup>15</sup> which are unfavorable prognostic markers for the disease.<sup>18-21</sup> Together, these functions of NO suggest that NOS2 expression

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in cancer biology could be disease promoting and upregulation of the enzyme in breast cancer and other cancer types could cause a poor outcome phenotype.

### NOS2 in Human Tumors

Aberrant expression of NOS2 has been commonly observed in human tumors including human breast,<sup>22,23</sup> colon,<sup>24</sup> stomach,<sup>25</sup> lung,<sup>26</sup> head and neck cancer,<sup>27</sup> among others. In the two breast cancer studies, increased NOS2 correlated with dedifferentiation as indicated by a high tumor grade<sup>22,23</sup> and increased tumor angiogenesis.<sup>23</sup> The correlation between increased NOS2 and a high tumor grade was also observed in two other studies<sup>15,28</sup> and one of them corroborated the finding that NOS2 upregulation is associated with an increased tumor vascularization.<sup>15</sup> In most breast cancer studies, NOS2 was found to be expressed by the cancer cells themselves. Several reports examined the relationship between NOS2 and patient outcome in breast cancer.<sup>15,28,29</sup> The two earlier studies detected NOS2 expression in a majority of the analyzed tumors (61 and 78%) and observed borderline associations between increased NOS2 and inferior breast cancer survival in two patient groups that consisted mainly of estrogen receptor (ER)-positive patients.<sup>28,29</sup> The third and largest study detected a moderate to strong NOS2 expression in 70% of the analyzed tumors.<sup>15</sup> This study found a significant association of NOS2 with poor survival only in the ER-negative patients. From these results it appears that tumor NOS2 expression may have a limited prognostic value in ER-positive breast cancer but is a predictor of outcome in the ER-negative disease.

### Induction of NOS2

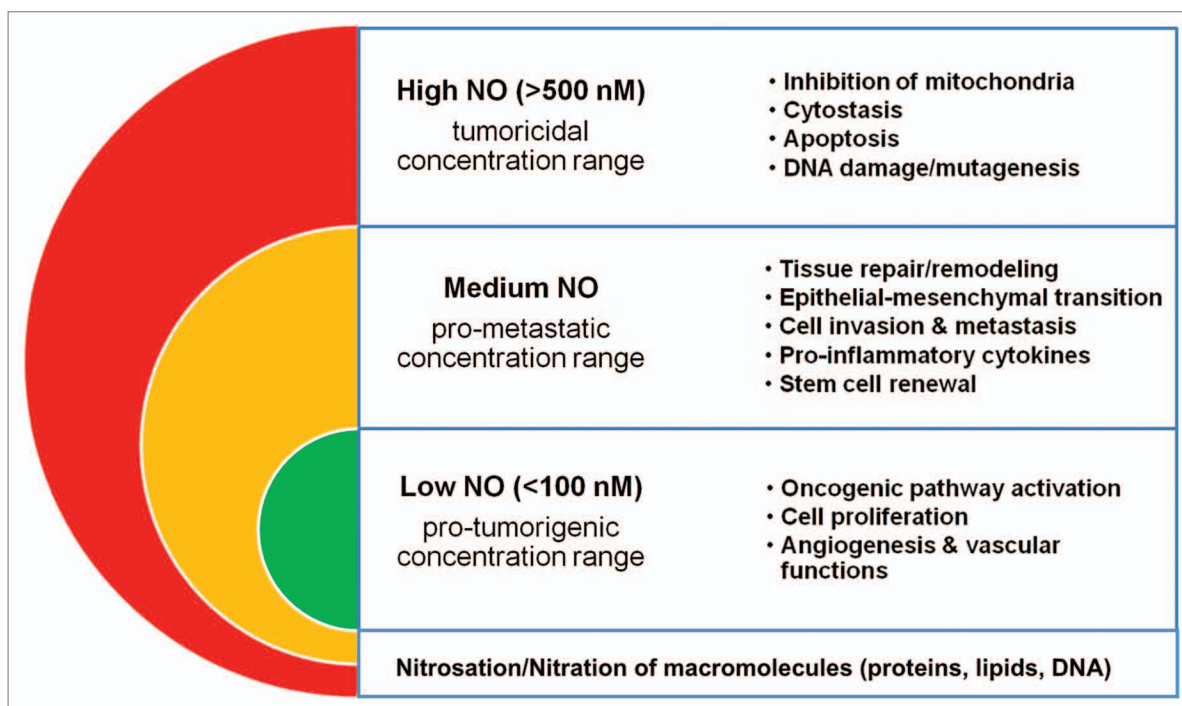
Constitutive expression of NOS2 is rarely observed and most cells do not express the enzyme without a stimulus. NOS2 expression occurs in normal breast tissue where weaning induces this enzyme in the lactating gland.<sup>30</sup> Estrogen has been found to alter NOS2 expression in murine macrophages, splenocytes and vascular smooth muscle cells by mechanisms involving ER,<sup>31,32</sup> but because of

the known differences in promoter regulation of murine and human NOS2,<sup>33,34</sup> we cannot be sure that estrogen would have similar effects on human NOS2. NOS2 is an inflammation responsive enzyme and bacterial lipopolysaccharide and cytokines have been shown to induce NOS2 in a variety of cell types through mechanisms that involve NFκB.<sup>33-35</sup> Hormones may interact with cytokines in NOS2 induction and progesterone has been shown to enhance cytokine-stimulated NOS2 expression in MCF7 human breast cancer cells.<sup>36</sup> However, the level of NOS2 induction can be different between murine and human cell lines, with human cell lines commonly showing a lower expression of NOS2 following cytokine exposure. This observation is explained, at least partly, by promoter differences that regulate NOS2 expression in response to inflammatory stimuli. These findings suggest that the induction of NOS2 in murine cells can lead to a phenotype that is somewhat different to that observed in their human counterparts because the effects of NO are strictly concentration-dependent with high concentrations causing cytostasis and apoptosis while lower concentrations may activate oncogenic signaling pathways, promote proliferation and invasion and inhibit apoptosis.<sup>14,17,37-39</sup> Thus, NOS2 effects in murine models may not apply to humans, and the use of the recently established human NOS2 transgenic mouse with a more human-specific release of NO may prove advantageous in experimental settings studying NOS2 effects in mouse models of cancer.<sup>40</sup> Other cancer-related signaling pathways that stimulate NOS2 expression include the Wnt adenomatous polyposis coli pathway, hypoxia, colony-stimulating factor-1 and epidermal growth factor (EGF) signaling. In human colon and liver cancer cell lines, Wnt signaling induces NOS2 in a β-catenin-dependent manner.<sup>41</sup> Hypoxia synergizes with interferon, resulting in a more than additive induction of NOS2 in murine macrophages.<sup>42</sup> This mechanism could be important for NOS2 induction associated with tumor hypoxia and the tumor-associated wound healing response. Colony-stimulating factor-1 released by human breast carcinoma cells was found to induce NOS2 in macrophages, leading

to increased NO-induced invasiveness of the breast cancer cells.<sup>43</sup> Lastly, EGF was shown to induce NOS2 in MDA-MB-468 human breast cancer cells. This cell line has basal-like breast cancer characteristics and EGF enhanced both NO production and vascular endothelial growth factor expression in these cells by a mechanism that involves the nuclear interaction of the EGF receptor and STAT3.<sup>44</sup>

### Effects of NO in Tumor Biology

NO can influence tumor biology in various and sometimes dichotomous ways (Fig. 1). Genetic ablation of NOS2 increases mammary tumor latency and inhibits lung tumor development in mouse models of cancer.<sup>45,46</sup> These genetic findings demonstrate that NOS2 can have tumor promoting activities. However, NOS2 expression or other forms of NO exposure are not always tumor promoting, and there are many examples where exposure to NO either delayed or inhibited tumor growth and metastasis.<sup>47-49</sup> In some studies, tumor cell NO inhibited primary tumor growth but stimulated experimental metastasis,<sup>50</sup> while others found that metastasis was suppressed by NO.<sup>49</sup> Because the effects of NO are strictly concentration-dependent with high concentrations causing cytostasis and apoptosis, at least some of the observed differences are likely explained by a difference in levels of NO exposure. If a NOS2 transgene is used to experimentally study the effect of NO on tumor growth, the use of a weak promoter, when compared to the CMV promoter, may reveal cancer promoting activities of the enzyme,<sup>6,8</sup> while CMV-driven expression of the transgene may yield high NOS2 expression and cytotoxic effects of NO.<sup>48,49</sup> Another factor that influences the effect of NO is the tumor microenvironment. For example, it was found that the ablation of NOS2 in cancer-prone p53 knockout mice can either suppress or enhance cancer development depending on the presence of chronic inflammation.<sup>51,52</sup> This dichotomy of NO function is well documented, and NO can cause both DNA damage and protect from cytotoxicity, and either inhibit or stimulate cell proliferation and migration, and apoptosis.<sup>14,37,39,53-56</sup> The NO effect will also



**Figure 1.** Concentration-dependent effects of NO in tumor biology.

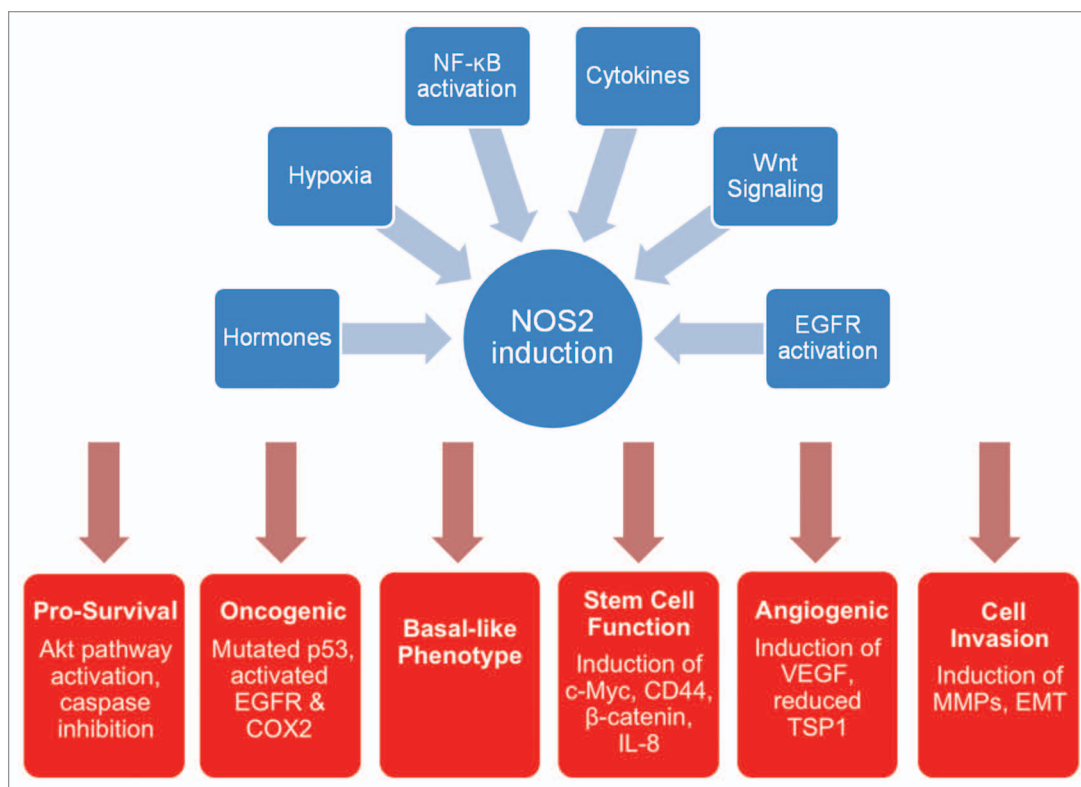
dependent on tissue oxygen tension and local superoxide concentrations,<sup>55</sup> and are therefore difficult to predict in human tumors. Thus, NO effects in human cancer are perhaps best estimated from the expression analysis of nitric oxide synthases in the tumors, adjacent non-tumor tissues, and in the involved lymph nodes, and examining their association with tumor markers and survival.

Cancer cell lines can adapt to high NO concentrations and thereby change their phenotype.<sup>57</sup> NO targets DNA repair processes and can induce genetic instability and cell transformation.<sup>58-61</sup> DNA damage can trigger activation of the tumor suppressor p53. We and others have reported the existence of a negative feedback loop in which NO-induced p53 activation results in p53-mediated growth arrest and apoptosis, and in the trans-repression of NOS2.<sup>8,62-64</sup> NO may promote carcinogenesis through the inactivation of wild-type p53 function, either by causing loss of DNA-binding activity<sup>65</sup> and/or selecting for mutant p53. The latter hypothesis is supported by our observations that the p53 mutation frequency in colon and breast tumors is increased in tumors with high NOS2.<sup>15,66</sup> Other candidate oncogenic functions of NO include

the activation of the cyclooxygenase-2 (COX2) pathway by NO and the induction of the chemokine receptor, CXCR4, which has a critical function in the homing of cancer cells to specific metastatic sites.<sup>67-69</sup> NOS2 and COX2 are both inflammation responsive enzymes that are commonly expressed in breast tumors and their co-expression could have more deleterious effects than the expression of only one of these two markers. The activation of COX2 by NO may also enhance the anti-apoptotic and pro-angiogenic effects of NO that are observed when local NO concentrations remain below a 300 to 500 nmol/l threshold. Here, NO can suppress apoptosis through the inhibition of caspases by S-nitrosylation.<sup>39</sup>

Other studies discovered that NOS2 expression correlates with increased Akt phosphorylation in breast tumors, and that NO induces Akt phosphorylation and activation of the oncogenic Akt pathway in breast cancer cells in culture.<sup>13,14</sup> Akt activation is a pro-survival signal that suppresses the activation of key factors like BAD and caspase-9 in the apoptosis pathway, but also activates eNOS and eNOS-mediated tumor maintenance.<sup>11,70</sup> We found that NOS2 expression was associated with Akt activation in both

ER-positive and ER-negative tumors.<sup>13</sup> However, NOS2 expression was a predictor of survival only in the ER-negative disease.<sup>15</sup> It is likely that both the difference in ER expression and also intrinsic differences between ER-negative and ER-positive breast tumors independent of the tumor ER status contribute to the increased responsiveness to NO in ER-negative breast cancer. We observed that interleukin-8 (IL-8) is significantly upregulated in ER-negative breast tumors, but not ER-positive tumors, with high NOS2. IL-8 is induced by NO.<sup>15,71-73</sup> This chemokine is preferentially expressed by ER-negative breast tumors and cancer cell lines and its expression is low or absent in most ER-positive breast cancer cells.<sup>74</sup> IL-8 is of particular importance in human breast cancer biology and is associated with increased cancer cell invasion and microvessel density.<sup>75</sup> IL-8 mediates metastasis in breast cancer and other cancers and its secretion correlates with early disease dissemination and poor survival.<sup>76</sup> We hypothesize that NOS2 may lead to poor survival among ER-negative patients partly because it induces IL-8 selectively in tumors of these patients, leading to increased angiogenesis and early metastasis. ER-negative tumors also tend to



**Figure 2.** Candidate pro-tumorigenic properties of NOS2 in ER-negative breast tumors. EMT = epithelial to mesenchymal transition.

have more tumor-associated macrophages than ER-positive tumors,<sup>77</sup> and many pro-inflammatory cytokines are more highly expressed in ER-negative tumors than ER-positive tumors.<sup>78</sup> Both macrophages and cytokines will alter the tumor microenvironment and may lead to a pro-inflammatory state and increased oxygen radical formation. As already mentioned at beginning of this section, the presence of chronic inflammation can lead to an opposite effect of NO in a mouse model of cancer.<sup>51,52</sup> Thus, NO biochemistry and signaling, which is greatly influenced by reactive oxygen species availability, may be different in ER-negative and ER-positive breast cancer.

### NOS2 and Basal-Like Breast Cancer

Recent results from large-scale gene expression profiling studies showed that ER-negative and ER-positive breast tumors should be further subdivided based on their gene expression profiles.<sup>79</sup> Distinct molecular signatures characterize three luminal subtypes among

the ER-positive tumors and basal-like, HER2-positive and normal-like subtypes among the ER-negative tumors. Among all subtypes, basal-like and HER2-positive, ER-negative tumors have been recognized as the most aggressive subtypes.<sup>21,80</sup> They are not treatable by endocrine-targeted therapy such as tamoxifen and aromatase inhibitors and have a worse prognosis than ER-positive breast tumors independent of therapy, particularly in the first five years following diagnosis.<sup>81,82</sup> There is an urgent need to identify novel targets for the treatment of ER-negative breast cancer in general, and even more for the basal-like subtype and the so called triple-negative tumors (ER/PR/HER2-negative) which mostly represent basal-like tumors but also some other ones.

We think that NOS2 and downstream targets of NOS2 are candidate new targets for therapeutic intervention in both the basal-like and triple-negative disease (Fig. 2). Recent research by our laboratory led to the novel and clinically significant observation that NOS2 expression is associated with a prognostic basal-like transcription pattern and is an

independent predictor of poor survival in women with ER-negative breast tumors.<sup>15</sup> NOS2 remained a predictor of poor survival when the analysis was restricted to only those patients with basal-like breast tumors. This result suggests that in addition to inducing a basal-like signature in ER-negative tumors, NOS2 may further enhance disease aggressiveness in the presence of this signature. Increased NOS2 in breast tumors also correlated with other poor outcome markers, such as an increased tumor vascularization and p53 mutation frequency, and activated EGFR. Both p53 mutations and EGFR overexpression occur more commonly in basal-like breast tumors than other breast cancer subtypes.<sup>21,79</sup> Additional work showed that NO activates EGFR, consistent with previous findings in lung cancer,<sup>83</sup> and induces proteins, such as CD44 and c-Myc, in ER-negative human breast cancer cells. These markers have been linked to an embryonic stem cell-like phenotype in breast cancer and disease outcome.<sup>19,84,85</sup> Recently, an increased expression of stem cell markers in basal-like tumors has been reported.<sup>86,87</sup> Among

them was CD44, which is a receptor for hyaluronan and osteopontin.<sup>88</sup> CD44 is a poor outcome marker in breast cancer<sup>18</sup> and CD44-positive breast cancer cells have an increased invasive activity,<sup>89</sup> increased resistance to radiation therapy and chemotherapeutics.<sup>90,91</sup> Together, these novel observations link NOS2 to the development of a poorly differentiated breast cancer phenotype with stem cell-like characteristics. NO may induce this phenotype by activation of c-Myc or by inducing the release of stem cell renewal factors like IL-8, a property that NO has, as we have shown.

### Concluding Remarks

In conclusion, NOS2 was found to be a predictor of survival and determinant of disease aggressiveness associated with ER-negative breast cancer. The underlying mechanisms that lead to an NO-induced poor outcome phenotype in breast cancer may include a combination of events, such as the induction of a basal-like phenotype, activation of the EGFR pathway, increased IL-8 and tumor angiogenesis and selection for mutant p53 cells. We propose that NOS2 and downstream targets of NOS2 signaling are novel therapeutic targets for ER-negative breast cancer in general and more specifically for basal-like breast cancer and the triple-negative disease.

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