



Article Differential Expression of Nitric Oxide Synthase Isoforms nNOS and iNOS in Patients with Non-Segmental Generalized Vitiligo

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Abstract: Nitric oxide (NO) is involved in several biological processes, but its role in human melanogenesis is still not well understood. Exposure to UVA and UVB induces nitric oxide production in keratinocytes and melanocytes through the activation of constitutive nitric oxide synthase, increasing tyrosinase activity and melanin synthesis, whereas inducible nitric oxide synthase over expression might be involved in hypopigmentary disorders. The aim of this study was to evaluate whether inducible nitric oxide synthase and neuronal nitric oxide synthase expression were modified in vitiligo skin compared to healthy controls. Skin biopsies were obtained from inflammatory/lesional and white/lesional skin in 12 patients with active, non-segmental vitiligo; site-matched biopsies of normal skin from eight patients were used as controls. Nitric oxide synthase isoforms expression was evaluated by confocal laser scanning microscopy and Western Blot analysis. Inducible nitric oxide synthase expression was significantly increased in inflammatory/lesional skin compared to healthy skin; melanocytes showed a moderate neuronal nitric oxide synthase expression in white/lesional skin, demonstrating that metabolic function still goes on. The obtained data demonstrated that vitiligo lesions were characterized by modifications of nitric oxide synthase isoforms, thus confirming the hypothesis that nitric oxide imbalance is involved in vitiligo and supporting the idea that nitric oxide synthase inhibitors might be used as a possible therapeutic approach for the management of vitiligo.

Keywords: vitiligo; nitric oxide; nitric oxide synthase isoforms; melanocytes; cytokines

1. Introduction

Vitiligo is a common progressive depigmentation of epidermis caused by the destruction of melanocytes and it is considered one of the most common pigmentary disorders occurring in around 0.1–2% of the worldwide population [1]. The pathogenesis of vitiligo is still not fully understood but several studies have suggested the possible involvement and interaction of genetic, biochemical, environmental, and immunological factors [2,3]. This theory is supported by robust evidence, such as the association of vitiligo with other autoimmune disorders of the endocrine glands [4–6] and the presence of a lymphocytic infiltrate observed in perilesional skin of actively spreading vitiligo [7].

Several cells are involved in vitiligo such as lymphocytes, keratinocytes, and endothelial cells, thus collaborating to melanocyte migration, proliferation, and differentiation [8–10].

To date, there is accumulating evidence that vitiligo may be a disorder of the entire epidermal melanin unit [11]. In particular, keratinocytes contribute to melanocyte homeostasis and the keratinocyte alteration may effect melanocyte dysfunction in vitiligo epidermis. Melanocytes adhere to the surrounding keratinocytes to the basement membrane of the epidermis with laminin-binding integrins and via E-cadherin-dependent adherence junctions. Through the interaction with the integrins, a whole series of messages concerning adhesion, differentiation, proliferation, and migration processes is transmitted from the extracellular matrix to melanocytes [12,13].

Vitiligo lesions are characterized by melanocyte death/disfunction caused by autoimmune, autocytotoxic, and neural mechanisms [2]. Vitiligo is also characterized by a blunt increase of reactive oxygen species (ROS) and inflammatory mediators, such as cytokines and nitric oxide (NO), thus contributing to melanocyte dysfunction and/or destruction [14,15]. NO is a highly reactive free radical with a short half-life, involved in several biological processes such as vascular homeostasis, neurotransmission, immunomodulation, and inflammation [16,17]. Although the well-known involvement in several physiological and pathophysiological conditions, its role in human melanogenesis is still under investigation [18–22].

NO is endogenously produced by three NO synthase enzymes (NOS): neuronal NOS (nNOS), constitutive endothelial NOS (eNOS) and inducible NOS (iNOS). All these isoforms have been isolated in human skin [16,23].

Exposure to UVA and UVB may induce the production of nitric oxide, in particular in keratinocytes and melanocytes, through the activation of constitutive NOS, leading to an increase in tyrosinase activity and melanin synthesis (paracrine and autocrine mediation of UV-induced melanogenesis) [24,25].

Alterations of NOS activity could influence melanocytes in many ways: (a) reduction of melanogenesis and accumulation of toxic intermediates of melanin synthesis; (b) increase of reactive oxygen species; and (c) decrease of cell ability to compensate oxidative stress. The possible consequences of abnormal production of nitric oxide on a photoinduced tissue reaction, like excessive entity and duration of the normal inflammatory response and facilitation of the induction of autoimmune phenomena, also have to be mentioned: they can be, in fact, causal or concausal factors of melanocyte death [14,26,27].

It has been demonstrated that an association between promotor polymorphisms within iNOS gene and NO overproduction, thus inducing melanocyte destruction and death [28,29]. Alterations of NOS activity influence melanocytes reducing the melanogenesis process, accumulating toxic intermediates of melanin synthesis, and increasing ROS production.

Therefore, the aim of this study was to evaluate the distribution and the expression of iNOS and nNOS in skin biopsies of patients with non-segmental generalized vitiligo.

2. Results

2.1. Immunofluorescence for nNOS and iNOS

The nitric oxide synthase isoform distribution in normal skin was used as control (Figure 1A–D). In particular nNOS was significantly expressed in keratinocytes of both basal and suprabasal layer. Enzymes expression was also variably detected in the cell body and dendrites of epidermis melanocytes. iNOS expression was not detected in keratinocytes and in melanocytes.

Vitiligo skin samples showed significant alterations in protein expression and distribution. In all inflammatory/lesional skin samples, iNOS expression was higher than control (Figure 2A–D). nNOS isoform was expressed in the stratum basal of epidermis, which is composed by proliferative keratinocytes and active melanocytes. Protein expressions were low in differentiating keratinocytes of superficial epidermal layers and were also variably detected in the cell body of epidermis melanocytes

in the suprabasal layer (Figure 3A–D). Furthermore melanocytes showed a moderate activation of nNOS in white/lesional skin (Figure 4A–D).



Figure 1. Normal skin: immunofluorescence staining of nNOS and iNOS (**A**,**B**). Confocal images obtained in "overlay" mode showed nNOS expression (arrows) in the sections (**A**,**B**); iNOS expression was not detected (**C**,**D**). The confocal image was "colored" and superimposed on a transmission image of the same skin area to show the exact localization of the reaction. Argon laser (568 nm) lens 3, zoom 1, original magnification $40 \times .$ (**A**,**B**) bar = 25 μ m; (**C**) bar = 250 μ m; (**D**) bar = 100 μ m.



Figure 2. Inflammatory/lesional vitiligo skin: immunofluorescence staining of iNOS (**A**,**B**). Confocal image obtained in "overlay" mode showed a significant iNOS expression both in the basal and in suprabasal layer. iNOS expression was not observed in epidermal melanocytes. Argon laser (568 nm) lens 3, zoom 1, original magnification $40 \times$. bar = 20 µm; (**C**,**D**) three-dimensional reconstruction obtained in depth coding mode. The colors reveal the spatial distribution (arrangement) of iNOS throughout the thickness of the section. (**A**,**B**) Argon laser (568 nm) lens 3, zoom 2, original magnification $40 \times$. bar = 25 µm.



Figure 3. Inflammatory/lesional vitiligo skin: immunofluorescence staining of nNOS (**A–D**). Confocal image obtained in "overlay" mode showed nNOS expression in in the stratum basal of epidermis. nNOS was also detected in the cell body of melanocytes in the suprabasal layers; (**B**) Three-dimensional reconstruction obtained in depth coding mode. The colors reveal the spatial distribution (arrangement) of nNOS throughout the thickness of the section; (**A**,**B**) Argon laser (568 nm) lens 3, zoom 2, original magnification $40 \times .$ bar = 25 µm; (**C**,**D**) Argon laser (568 nm) lens 3, zoom 1,1, original magnification $40 \times .$ bar = 25 µm.



Figure 4. White/lesional vitiligo skin: immunofluorescence staining of nNOS (**A–C**); Confocal image obtained in "overlay" mode showed a moderate nNOS expression in white/lesional skin (**B–D**). Three-dimensional reconstruction of the same confocal image obtained in "depth coding" mode. The colors reveal the spatial distribution (arrangement) of nNOS throughout the thickness of the section. (**A**,**B**) Argon laser (568 nm) lens 3, zoom 1,2, original magnification $40 \times$. bar = 25 µm; (**C**,**D**) Argon laser (568 nm) lens 3, zoom 1, original magnification $40 \times$ bar = 20 µm.

These results demonstrated that iNOS expression was significantly increased in inflammatory/ lesional skin compared to healthy skin; melanocytes showed a moderate expression of nNOS both in inflammatory/lesional and white/lesional skin, suggesting that melanocytes express nNOS and still carry out some metabolic function.

2.2. Western Blot Analysis for nNOS, iNOS, pFAK, and Determination of NO₂⁻/NO₃⁻ Levels

nNOS, iNOS, and pFAK expression was also evaluated by Western Blot analysis (Figure 5A–C). Vitiligo patients showed an increased expression of both iNOS and pFAK, whereas a reduced expression of nNOS in inflammatory/lesional skin was observed compared to control group (p < 0.01 vs. CTRL) (Figure 5A,B). Moreover, vitiligo skin samples showed an increased NO₂⁻/NO₃⁻ levels compared to controls (Figure 5D).



Figure 5. Western Blot Analysis of nNOS (**A**); iNOS (**B**); and pFAK (**C**) in inflammatory/lesional skin. Lesional skin showed a reduced expression of nNOS and an increased expression of iNOS and pFAK. Samples were coded and the experiments were repeated three times; NO_2^-/NO_3^- levels (**D**) in vitiligo and control samples. Vitiligo skin samples showed increased production of NO_2^-/NO_3^- levels. * p < 0.01 compared to control group.

2.3. Statistical Analysis

Vitiligo patients showed an increased expression of iNOS and a reduced expression of nNOS in vitiligo skin compared to control group (p < 0.01 vs. CTRL).

3. Discussion

Nitric oxide is an important mediator in many physiological and pathological processes such as microcirculation, melanogenesis, keratinocyte response to UV radiation, cell growth and differentiation. A potential role for NO in hair follicle biology has been recently advanced however, little is known about the role of NO in the human hair follicle and in the hair cycling processes [30,31]. Several

evidences have suggested its critical role in many inflammatory, hyperproliferative, and autoimmune diseases [20–22,32].

NO is a highly reactive messenger: it has no electric charge and easily passes through membranes. It is produced by the conversion of L-arginin into L-citrullin through a reaction catalyzed by enzymes belonging to the family of nitric oxide synthase (NOS). Two major isoforms of NOS are known, "constitutive", further divided into neuronal (nNOS) and endothelial (eNOS) types, and "inducible" (iNOS), involved in cell homeostasis and in modulation of immune and cytotoxic response [16,17,22]. Both eNOS and nNOS were found in several tissues, and they are synthesized by keratinocytes, melanocytes, and fibroblasts in skin. iNOS plays an important role in inflammatory and immune processes, both physiological and pathological ones. Several cytokines (IL-1, IL-6, IL-23, IL-17, IL-33, IFN- γ , TNF- α) stimulate iNOS expression [9,22,33] and play an important role also in establishing depigmentation. For example, TNF- α may increase iNOS release through the production of tetrahydrobiopterin, a potent inhibitor of melanin biosynthesis [34] and an essential cofactor of iNOS [35,36].

Increased serum levels of IL-23 and IL-33 have been shown in patients affected by vitiligo, which are responsible for the exacerbation of the disease [37,38].

Both NO and IL-23 may induce an autocrine loop within the innate immune system, supporting the hypothesis of an existing association between vitiligo, autoimmune disorders and T helper-17 axis [39,40].

Also IL-33 stimulates NO release by iNOS activation [41] and promotes TNF- α and IL-6 release in keratinocytes [42].

Among oxidative processes activated by cytokines accumulation, non-enzymatic glycation and oxidation of macromolecules, corresponding respectively to AGEs and AOPPs, can be closely associated with autoimmune diseases involving skin, included vitiligo. In fact, AOPPs and AGEs levels are significantly higher in serum of patients affected by vitiligo than healthy controls [43]. Both RAGE (receptor for advanced glycationed products) and ST2 are also up-regulated in vitiligo and the IL-33/ST2 signaling may be considered as a pathway that plays a pivotal role in immune response and tissue injury/repair, causing in turn the secretion of proinflammatory factors [38,44]. Moreover, exogenous AGE administration may result in a more pronounced IL-23/IL-17 immune response, thus increasing NO secretion [45].

The production of large amounts of NO related to the increased activity of iNOS leads to self-destruction of melanocytes [18]. De novo attachment of melanocytes to the extracellular matrix is consequently reduced [46] and skin depigmentation is promoted [47]. This mechanism could be important in vitiligo, where an initial imbalance of epidermal cytokines at sites of lesions could cause tetrahydrobiopterin overexpression and iNOS activation, with consequent nitric oxide overproduction leading to loss and self-destruction of melanocytes [34,36].

Alterations on skin cells, such as melanocytes and keratinocytes may be the result of deficits in redox equilibrium, with an engagement of mitochondria [48,49]. One of the main molecules involved in mitochondrial control is PGC1 α . When damage occurs, it may stimulate FAK phosphorylation, thus promoting PGC1 α [50]. Impaired mitochondrial activity raised mitochondrial mass and increased PGC1 α and FAK expression [51]. In the present study, vitiligo skin samples showed phospo-FAK expression stimulation compared to controls. This result let us hypothesize that FAK overexpression might also be responsible for PGC1 α activation in these group of patients affected by vitiligo, but further experiments will be necessary.

Moreover, here we provide evidence that iNOS expression and NO_2^-/NO_3^- levels release were increased in vitiligo lesions compared to healthy controls. In particular, iNOS expression was augmented in all patients and in all layers of epidermis. The increased activity of iNOS may be responsible for the increased production of NO in vitiligo, thus contributing to reduction/death of melanocytes up to depigmentation. However, we have demonstrated that nNOS is expressed in vitiligo lesions, thus confirming that melanocytes are not totally missing and some metabolic function is still going on [11,52].

Vitiligo is the result of several factors, among them increased cytokines expression and iNOS activation play a key role in the depigmentation process even if NOS overexpression may not be considered the only important factor involved in vitiligo. Previously, some studies demonstrated that modification of biosynthesis of tetrahydrobiopterin (especially, recycling of BH4), as well as imbalance of anti-oxidant system in vitiligo [34,53], could deteriorate the balance of BH4/BH2 levels, with following dysfunction of NOS activity [54].

As the current study is largely descriptive, further functional studies on altered biological pathways are necessary: knowing exactly the role that nitric oxide plays is important not only to clarify the pathogenesis of this disease, but also because in the future it could be an important therapeutic target.

4. Materials and Methods

4.1. Patients

The study was carried out in 12 patients, four males and eight females, aged 52–76 years affected by active non-segmental vitiligo. The duration of vitiligo ranged from 1 to 18 years, mean extent was $51 \pm 24.3\%$ of involved BSA (range 35–90%). Vitiligo was considered "active" whether BSA was increased in the last six months.

Three patients were affected by autoimmune thyroiditis, two patients by autoimmune diabetes mellitus, two patients by psoriasis, one by alopecia areata, and four patients by psychological stress related to their lifestyle. Two patients had family history of vitiligo; six patients were used as control, matched for age and gender.

All patients and controls gave their written informed consent before the enrollment. The protocol was approved by the Ethics Committee of the University Hospital of Messina, Italy (prot. 15/17, 02/05/2017). All procedures performed in the study involving human participants were in accordance with the ethical standards of the institutional and/or national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards.

4.2. Skin Biopsies

Biopsies of inflammatory/lesional and white/lesional skin were collected from 12 patients with active, non-segmental vitiligo, whereas site-matched biopsies of normal skin were obtained from three healthy males and three females with no history of neither vitiligo nor psoriasis.

4.3. Immunofluorescence

Skin biopsies were fixed in 3% paraformaldehyde in 0.2 mol·L⁻¹ of phosphate buffer (pH 7.4). Skin samples were washed in 0.2 mol·L⁻¹ phosphate buffer and in phosphate-buffered saline, infused with 12% and 18% of saccharose, frozen in liquid nitrogen and then sectioned with a cryostat.

The obtained sections were placed on gelatin-coated slides and incubated with 0.6 mL of sheep serum, 1 mL of 45 mmol·L⁻¹ NaCl and 1 mL of 20 mmol·L⁻¹ phosphate buffer for 15 min to avoid non-specific background staining.

Sections were then incubated with primary anti-nNOS and anti-iNOS antibodies (1:200; BD Transduction Laboratories, Lexington, KY, USA). Biotinylated anti-immunoglobulin (1:500; Amersham, UK) was used as secondary antibody and Texas red–streptavidin (1:100; Amersham, UK) was used as fluorochrome.

4.4. Confocal Microscopy

Zeiss LSM 410 inverted confocal microscope with an argon laser (wavelength 488 nm, 100 mW) and two He–Ne lasers (wavelength 543 and 633 nm, 5 mW) was used to study all sections. Collected images were digitized at a resolution of 8 bits into an array of 512×512 pixels.

Fluorescent optical sections were obtained using an He–Ne laser (wavelength 543 nm and a TRIC set of filters) at a 1-s scanning speed; an average of up to 16 sections was obtained.

Images were collected using different software functions: (i) overlay (or projection on *y*-axis) to obtain images representing samples throughout their thickness; and (ii) depth coding (or color overlay), attributing a color. A color scale represented different levels in the *z*-axis.

4.5. Western Blot Analysis for nNOS, iNOS, and pFAK

Skin samples were homogenized in a lysis buffer (1% Triton; 20 mM Tris/HCl, pH 8.0; 137 mM NaCl; 10% glycerol; 5 mM EDTA 1 mM phenylmethylsulfonyl fluoride; 1% aprotinin; 15 μ g/mL leupeptin). Protein samples (30 μ g) were denatured in a reducing buffer (62 mM Tris pH 6.8, 10% glycerol, 2% sodium dodecyl sulfate, 5% mercaptoethanol, 0.003% bromophenol blue) and separated by electrophoresis on a sodium dodecyl sulphatepolyacrylamide gel (7.5%).

The separated proteins were then transferred into PVDF membranes using a transfer buffer (39 mM glycine, 48 mm Tris pH 8.3, 20% methanol) at 200 mA for about 60 min. After incubation with 5% non-fat dry milk in TBS 0.1% and three washes with Tween TBS 0.15% Tween, membranes were coated with primary nNOS, iNOS and pFAK (rabbit polyclonal to FAK phospho Y397; ab194935, Abcam) antibodies diluted in TBS 0.1%, overnight at 4 °C.

The day after, membranes were washed three times with TBS 0.15% Tween and then incubated with a secondary antibody peroxidase-conjugated goat anti-rabbit immunoglobulin G (Pierce, UK) for 1 h at room temperature. The membranes were washed and analyzed by the enhanced chemiluminescence system according to the manufacture's protocol (Amersham Bioscience, Amersham, UK). The protein signals were quantified by scanning densitometry using a bioimage analysis system (BioProfil, Celbio, Milan, Italy). The results from each experimental group were expressed as relative integrated intensity compared with control skin. Equal loading of protein was assessed on stripped blots by immunodetection of β -actin with a rabbit monoclonal antibody (Cell Signaling, Danvers, MA, USA) and peroxidase-conjugated goat antirabbit immunoglobulin G (Pierce, UK).

4.6. NO₂⁻/NO₃⁻ Determination

 NO_2^-/NO_3^- levels were determined in skin biopsies using the Griess reaction, as previously described [55,56]. Both samples and standards were assayed in duplicate. Data are expressed as means \pm SD

4.7. Assessment of the Distribution of nNOS and iNOS and Statistical Analysis

The epidermis and dermis were studied. Three fields (magnification $\times 20$, area approximately 800 μ m²) for each specimen were considered to assess the distribution of the proteins. Comparison was made by three independent observers in a blind manner.

The results were analyzed by analysis of variance. The results are expressed as mean \pm SD p < 0.01 was considered as statistically significant.

Author Contributions: Mario Vaccaro and Natasha Irrera conceived and designed the study and revised the paper; Giuseppina Cutroneo, Giuseppina Rizzo, Federico Vaccaro, and Francesco Borgia performed the data; Giuseppe P. Anastasi and Serafinella P. Cannavò analyzed and interpreted data. Francesco Squadrito and Domenica Altavilla led the design and drafted the manuscript.

Conflicts of Interest: The authors declare no conflict of interest.

Abbreviations

NO	Nitric oxide
NOS	Nitric oxide synthase
iNOS	Inducible nitric oxide synthase
nNOS	Neuronal nitric oxide synthase
eNOS	Endothelial nitric oxide synthase

BH4 Tetrahydrobiopterin

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