Vitamins C and E downregulate vascular VEGF and VEGFR-2 expression in apolipoprotein-E-deficient mice

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Abstract

Anti-angiogenic therapy reduces both plaque growth and intimal neovascularization in apolipoprotein-E-deficient mice (apoE−/−). Vascular endothelial growth factor (VEGF) has been suggested as playing a role in the development of atherosclerosis. We examined the hypothesis that VEGF and VEGF receptor-2 (VEGFR-2) expression is upregulated in apoE−/− and, since it could be driven by oxidative stress, tested whether dietary supplementation with vitamins C and E could downregulate it.

Two-month-old apoE−/− received vitamin C combined with α- or β-tocopherol for 4 weeks. Aortic VEGF and VEGFR-2 expression were measured by RT-qPCR and western blot.

ApoE−/− showed significantly higher expression of aortic VEGF and VEGFR-2 mRNA (P < 0.001) and protein (P < 0.001) than wild-type mice, as well as increased plasma VEGF (P < 0.001). Vitamin C and α-tocopherol significantly reduced aortic VEGF and VEGFR-2 expression in apoE−/− (P < 0.001), circulating VEGF (P < 0.01) and plasma lipid peroxidation (P < 0.01). apoE−/− receiving vitamin C and β-tocopherol showed diminished lipid peroxidation and VEGFR-2, but only partial reduction of VEGF expression.

These data demonstrate that augmented VEGF and VEGFR-2 expression in apoE−/− vasculature can be downregulated by vitamins C and E, at least partially through oxidative stress reduction. This novel mechanism could contribute to explaining the beneficial effects of antioxidant vitamins in experimental atherosclerosis.

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1. Introduction

Vascular endothelial growth factor (VEGF), the main agent responsible for angiogenesis and vascular permeability, is a 45kDa glycoprotein secreted in the vascular wall by endothelial and smooth muscle cells. It has also been implicated in the maintenance of endothelial integrity, partly through upregulation of endothelial nitric oxide synthase expression [1]. VEGF expression is regulated mainly by hypoxia [2], oxidative stress [3] and nitric oxide [4,5]. The biological effects of VEGF are mediated mainly by two tyrosin kinase receptors: VEGFR-1 (Flt-1) and VEGFR-2 (Flk-1/KDR). VEGFR-2 is essential for vasculogenesis, angiogenesis and hematopoiesis [6], and mediates VEGF antiapoptotic effects in endothelium [7].

Despite these functions, VEGF could also be considered a pro-atherogenic factor, because its administration increases the rate and degree of atherosclerotic plaque formation in apo-E/apo-B100 deficient mice and in hypercholesterolemic rabbits [8]. Moreover, VEGF has been found to be upregulated in human coronary atherosclerotic lesions but not in normal coronary segments [9]. Although the mechanism by which VEGF induces plaque formation has not been established, it has been suggested...
that new vessel formation causes plaque progression [8]. The administration of the anti-angiogenic drug endo-
statin to apolipoprotein-E-deficient mice (apoE/−/−), a well-characterized model of atherosclerosis, reduces both plaque growth and intimal neovascularization, providing evidence that neovascularization is required for atheroscle-
rosis plaque development [10].

Interestingly, the apoE−/− murine model of hypercholes-
terolemia is characterized by increased oxidative stress [11] and reduced nitric oxide bioavailability [12], two mecha-
nisms known to contribute to atherosclerotic plaque develop-
ment and also involved in the regulation of VEGF expression
[3,4]. Increased VEGF levels have been reported in plasma from hypercholesterolemic individuals [13] and in coronary arteries from hypercholesterolemic pigs [14]. Moreover, it has been shown that oxidized LDL, a key molecule in the atherosclerotic process, can upregulate VEGF expression in macrophages and endothelial cells, at least partially through the activation of PKA [15]. In this context, assuming that VEGF acting through VEGFR-2 can be at least one of the factors responsible for the angiogenic processes required for atherosclerosis plaque development, we hypothesized that either of them could be upregulated in the vascular wall of apoE−/− mice.

A potential role of antioxidant vitamins in cardiovascular disease prevention has been proposed, based on the essen-
tial role of lipoproteins oxidation in the initiation and pro-
gression of atherosclerotic lesions. To date, there is no clear evidence from clinical trials that antioxidants provided by dietary intake or in the form of vitamin supplements can re-
duce the risk of atherosclerosis [16,17]. However, a number of studies in different animal models have considered the potential for different dietary antioxidants to help prevent development and progression of atherosclerosis, and most of the experimental evidence helps substantiate this hypoth-
esia [18,19]. Particularly, α-tocopherol has been shown to decrease lipid peroxidation and atherosclerosis in apoE−/− [11].

Since VEGF and/or VEGFR-2 could be upregulated in apoE−/− vasculature as a consequence of increased oxida-
tive stress, we investigated whether dietary supplementation with vitamin C and α-tocopherol could downregulate VEGF and VEGFR-2 expression in this animal model. The com-
bination of vitamin C and β-tocopherol was also studied to elucidate whether antioxidant mechanisms can fully account for the effects of vitamin supplementation.

2. Methods

2.1. Mice

C57BL/6 wild type (WT) mice and apoE−/− were pro-
cured from the colony at CIFA (GLP accredited center at the University of Navarra) animal facilities, originat-
ing from breeders obtained from Center for Transgene Technology and Gene Therapy (Flinders Interuniversity

Institute for Biotechnology, Leuven). The investigation was performed in accordance with the European Com-
munity guidelines for animal ethical care and use of lab-
oratory animals (Directive 86/609), and was approved by the University of Navarra Animal Research Review Committee.

Two-month-old male wild type and apoE−/− mice fed a standard rodent diet (Harlan Teklad 9605) were di-
vided into four groups (n = 11): non-treated wild type and apoE−/−, and vitamins C and E supplemented mice (WT-CE, apoE−/−/CE). Ascorbic acid (120 mg kg−1 per
day) and α-tocopherol (210 mg kg−1 per day) were given in drinking water. An additional group of apoE−/− mice re-
ceived 120 mg kg−1 per day ascorbic acid and 210 mg kg−1 per day β-tocopherol (n = 5, apoE−/−/−CE). Two hun-
dred micromolar stock solutions of α-tocopherol and β-tocopherol in ethanol were prepared to facilitate final dilutions in water. β-tocopherol was generously provided by Dr. Christine Gartner (Cognis, Germany). Daily liquid intake, measured by weighing water bottles, was similar in mice from each experimental group (5 ml per animal per
day). After 4 weeks of treatment, animals were sacrificed and blood was drawn by cardiac puncture.

Plasma total cholesterol and triglycerides were deter-
mined by colorometric assay (Sigma). Plasma concentration of thiobarbituric acid-reactive substances (TBARS) was measured, as indicator of lipid peroxidation [20]. Aorta was re-
moved, dissected free of interstitial tissue with the help of a surgical microscope (Zeiss OPMI 6CR/5S), immediately frozen and stored at −80°C. Tocopherol concentration in plasma was measured by high performance liquid chro-
matography [19].

2.2. RNA isolation and real-time quantitative RT-PCR

Total RNA was extracted from each aorta with TRIzol (Invitrogen) and quantified with ultraviolet spectroscopy. Total RNA (5 μg) was reverse-transcribed with oligo (dT) primers with M-MLV reverse transcriptase (Invitrogen), in the presence of SuperRNase In (Ambion).

Three hundred nanograms of reverse-transcribed RNA was primed with oligonucleotides specific for VEGF (5′-GAGACCTTGTTGGAATC-3′ and 5′-TTTTAGGGT- TTCATCC-3′), VEGFR-2 (5′-GACCCGGGGAAGTGGTTTGTGA-3′ and 5′-GTCGAGGGGAGGAGTGGTGAGT- GAAAT-3′), and β-actin (5′-GCTTCTTCTGTTGTATG-3′ and 5′-ACCGAGCTGACTAAACTCA-3′). PCR was per-
formed as previously described [95°C for 10 min and run for 40 cycles at 95°C for 15 s and 60°C for 1 min] with SYBR Green PCR Master Mix (Applied Biosystems) on the ABI PRISM 5700 Detection System (Applied Biosystems) [21]. Potential genomic DNA contamination was excluded by using intron-encompassing primers. All samples were assayed in triplicate and normalized on the basis of their β-actin content.

2.3. RNA isolation and real-time quantitative RT-PCR

In the present study, we have investigated the

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2.3. Protein extraction, electrophoresis and western blotting

Following TRizol manufacturer’s instructions, protein was extracted from aortic homogenates after RNA isolation. Protein concentration was determined according to Bradford [22]. VEGF was quantified in murine plasma and aorta by western blotting: plasma samples (diluted 1:20 in saline) and aortic homogenates (10 μg total protein) were subjected to 12% SDS-PAGE, electrophoresed onto nitrocellulose membranes and probed with anti-VEGF (1:1000, sc6251, Santa Cruz Biotechnologies), followed by a peroxidase-linked sheep anti-mouse IgG (1:5000, Amersham Biosciences). Blots were developed by enhanced chemiluminescence according to the manufacturer’s instructions (ECL, Amersham Biosciences). Aortic expression of VEGFR-2 was assessed similarly, subjected to 6% SDS-PAGE, probed with anti-VEGFR-2 (1:1000, sc6251, Santa Cruz Biotechnologies) and developed by enhanced chemiluminescence (ECL Plus, Amersham Biosciences). Even loading was verified by Ponceau red staining.

2.4. Statistical methods

Data are presented as mean ± standard deviation. Analysis of variance (ANOVA) or Kruskal–Wallis test was applied to assess differences between treatment groups. After a significant ANOVA or Kruskal–Wallis, comparisons were made with Bonferroni posthoc or Mann–Whitney test, respectively. A value of P < 0.05 was accepted as statistically significant.

3. Results

3.1. Cholesterol, triglyceride and lipid peroxidation in murine plasma

Table 1 shows the general characteristics of the different animal groups. Plasma total cholesterol and triglycerides were significantly higher in apoE−/-- than in WT mice (P < 0.001 in both cases). Both types of mice presented similar plasma tocopherol concentration, although tocopherol/cholesterol ratio was markedly reduced in apoE−/-- (P < 0.001), in agreement with the increased TBARS (P < 0.01), in comparison with WT mice.

3.2. Vascular VEGF and VEGFR-2 expression in apoE−/-- mice

Vascular expression of VEGF mRNA in apoE−/-- was nearly five times higher than in WT group (P < 0.001; Fig. 1A). As shown in Fig. 1B, the observed mRNA increase was accompanied by a higher VEGF protein content in the vessel wall (P < 0.001).

Since VEGFR-2 plays a leading role in VEGF-induced angiogenesis, we analyzed vascular VEGFR-2 expression in the same in vivo model. As expected, a threefold increase in VEGFR-2 mRNA was observed in apoE−/-- mice as compared with WT (P < 0.001; Fig. 2A), and a higher VEGFR-2 protein content in the vessel wall could also be demonstrated (P < 0.001; Fig. 2B).

Relative VEGF levels in plasma were analyzed by western blotting to determine if they would reflect the observed changes in the vascular wall. VEGF plasma levels were significantly higher in apoE−/-- rather than WT mice (P < 0.001; Fig. 3).

3.3. Effect of vitamin C and α-tocopherol on VEGF and VEGFR-2

Compliance with the vitamins supplementation was evident, as plasma tocopherol levels in the groups receiving vitamin E increased to above the levels in non-supplemented mice (P < 0.001), and significantly reduced apoE−/-- plasma TBARS (P < 0.01) down to WT levels, even though tocopherol/cholesterol ratio was still lower than WT (P < 0.01; Table 1). While plasma total cholesterol and triglycerides remained unchanged after treatment in WT mice, a marginal reduction in total cholesterol levels was observed in treated apoE−/-- (P < 0.01), the levels still remaining eightfold higher than in WT.

Table 1

<table>
<thead>
<tr>
<th></th>
<th>Total cholesterol (mmol/l)</th>
<th>Triglycerides (mmol/l)</th>
<th>Tocopherol (μmol/l)</th>
<th>Tocopherol/cholesterol</th>
<th>TBARS (mmol MDA/l)</th>
</tr>
</thead>
<tbody>
<tr>
<td>WT (n = 11)</td>
<td>1.71 ± 0.34</td>
<td>0.60 ± 0.26</td>
<td>35.3 ± 13.7</td>
<td>18.1 ± 7.1</td>
<td>1.18 ± 0.36</td>
</tr>
<tr>
<td>WT-CE (n = 11)</td>
<td>1.78 ± 0.31</td>
<td>0.53 ± 0.12</td>
<td>94.9 ± 28.3**</td>
<td>50.2 ± 13.2**</td>
<td>0.92 ± 0.52</td>
</tr>
<tr>
<td>apoE−/-- (n = 11)</td>
<td>18.78 ± 4.04***</td>
<td>1.16 ± 0.34***</td>
<td>36.8 ± 12.5</td>
<td>1.9 ± 0.7***</td>
<td>1.76 ± 0.45***</td>
</tr>
<tr>
<td>apoE−/-- CE (n = 11)</td>
<td>14.74 ± 3.76***</td>
<td>1.04 ± 0.15***</td>
<td>103.4 ± 34.5††</td>
<td>7.4 ± 2.7††</td>
<td>1.25 ± 0.39††</td>
</tr>
<tr>
<td>apoE−/-- CE (n = 5)</td>
<td>18.83 ± 2.38***</td>
<td>0.88 ± 0.10***</td>
<td>100.7 ± 46.7†††</td>
<td>7.5 ± 3.5†††</td>
<td>1.44 ± 0.16†††</td>
</tr>
</tbody>
</table>

Plasma total cholesterol (mmol/l), triglycerides (mmol/l), tocopherol (μmol/l) and TBARS (mmol MDA/l) are shown. Statistically significant differences from WT are marked as **(P < 0.01), *** (P < 0.001), and from apoE−/-- as †† (P < 0.01), ††† (P < 0.001).

WT: wild type; WT-CE: supplemented with vitamins C and E; apoE−/--: apoE−/-- deficient; apoE−/-- CE: apoE−/-- supplemented with vitamin C and apoE−/-- CE: α-tocopherol or β-tocopherol.
Fig. 1. Changes in vascular VEGF expression (mRNA and protein), reported as percentage increase with respect to untreated wild type (WT) mice. (A) VEGF mRNA, assessed by RT-qPCR and corrected by β-actin mRNA content, is increased in apoE−/− mice. Vitamin C + α-tocopherol treatment (apoE−/−CE) reduces it significantly, but vitamin C + β-tocopherol (apoE−/−CIE) only achieves a partial reduction. (B) Augmented VEGF protein in apoE−/−, assessed by western blot (densitometric analysis and representative blot showing the 25 kDa VEGF monomer), is significantly reduced in apoE−/−CE and apoE−/−CIE. Bars represent the mean ± S.D. for all the animals in the experimental group (size indicated below x-axis). Statistically significant differences from WT are marked as ***(P < 0.001), and from apoE−/− as †††(P < 0.001).

Interestingly, the combination of vitamin C and α-tocopherol significantly decreased aortic VEGF and VEGFR-2 expression in apoE−/−, both at mRNA and protein level (P < 0.001 in all cases; Figs. 1 and 2), but it did not substantially modify their expression in WT. It only showed a marginal reduction in VEGF and VEGFR-2 mRNA expression remaining unchanged.

As found in the vessel wall, vitamin C and α-tocopherol also markedly reduced VEGF plasma levels in apoE−/− mice (Fig. 3, P < 0.01), as compared with WT.

3.4. Effect of vitamins C and α-tocopherol on VEGF and VEGFR-2

Dietary supplementation of apoE−/− mice with ascorbic acid and β-tocopherol was equally efficient to increase plasma tocopherol concentration (P < 0.01) and to reduce plasma TBARS (P < 0.01), the tocopherol/cholesterol ratio being still lower than WT (P < 0.01; Table 1). No modifications were observed in total cholesterol or triglyceride concentrations.

In contrast to mice receiving vitamin C and α-tocopherol, treatment of apoE−/− with vitamin C and β-tocopherol only achieved a partial, although significant, reduction (20%) in vascular VEGF mRNA and protein expression (P < 0.05, Fig. 1). However, no differences between the effect of α and β-tocopherol were observed in plasma VEGF (Fig. 3) or in vascular VEGFR-2 mRNA and protein levels (Fig. 2).

No deleterious effects were observed in any of the animals receiving the different vitamin treatments.

4. Discussion

We report herein that vitamins C and E markedly reduce the expression of VEGF and its endothelial receptor
differences from WT are marked as ††. Statistically significant differences from WT are marked as **P < 0.01, and from apoE−/− as †† P < 0.01.

VEGFR-2 in a murine model of atherosclerosis, representing a novel effect of these compounds. In this study, we demonstrate a significant increase of VEGF and VEGFR-2 in the aorta of apoE−/− mice, as compared to WT. Furthermore, to our knowledge, this is the first report showing that dietary supplementation with vitamins C and E significantly reduces vascular expression of VEGF and VEGFR-2 in this experimental model.

VEGF has a dual role in vascular homeostasis: it is not only the principal agent responsible for angiogenesis, vascular permeability and maintenance of endothelial integrity, but it also favors the progression and growth of the atherosclerotic plaque [8]. VEGF and VEGFR-2 expression were measured in apoE−/− mice, an accepted model of atherosclerosis showing high cholesterol concentration, increased lipid peroxidation [11] and low nitric oxide bioavailability [12]. We found higher levels of aorta and plasma VEGF in apoE−/− than in WT, in agreement with previous observations in the coronary arteries of hypercholesterolemic pigs [14], as well as in the plasma of hypercholesterolemic patients [13]. Moreover, increased vascular VEGF was accompanied by a marked upregulation in VEGFR-2 expression, a novel finding not described previously in hypercholesterolemia.

Upregulation of VEGF and VEGFR-2 in the vessel wall could account for the increased vascular permeability in atherosclerotic plasma [23] and enhanced vasa vasorum neovascularization observed in experimental models of hypercholesterolemia [24]. Although the mechanism by which hypercholesterolemia may induce VEGF upregulation in the vessel wall remains unclear, possible pathophysiological stimuli include hypoxia [14], oxidative stress [3] and low NO bioavailability [4].

Moreover, VEGF upregulation has been proposed to be a non-specific stress-induced vascular response in the adult organism [25], or to constitute a vascular homeostatic mechanism for compensating endothelial dysfunction [1]. Independently of the mechanism responsible, sustained vascular VEGF overexpression could contribute to the atherosclerotic process by favoring lipoprotein infiltration through increased vascular permeability [23], inflammation [26] and angiogenesis [8]. The anti-atherosclerotic effect of the angiogenesis inhibitors endostatin and TNP-470 in the apoE−/− mice [10] would support this hypothesis.

Dietary supplementation of apoE−/− mice with ascorbic acid and α-tocopherol significantly decreased both VEGF and VEGFR-2 expression, as well as plasma TBARS, while it augmented the tocopherol/cholesterol ratio. It can be speculated that the observed downregulation could be due to their antioxidant activity and/or the protein kinase C (PKC) inhibitory activity of α-tocopherol. A similar vitamin E dose, higher than that used therapeutically in humans, has been proved to reduce oxidative stress and atherosclerosis in apoE−/− mice [11], and vitamin C was administered to ensure α-tocopherol regeneration from α-tocopherohydroxyl radical, thereby preventing vitamin E pro-oxidant activity and tocopherol-mediated peroxidation [27]. In spite of the high dose of vitamin E administered, the threefold increase observed in plasma tocopherol is not far away from the twofold increase reported in several human trials studying vitamin E supplementation.

The combination of ascorbic acid and β-tocopherol, an equally strong antioxidant in comparison to α-tocopherol but lacking other biological properties present in α-tocopherol (PKC inhibitory and antiproliferative activities) [28] partially decreased VEGF expression, suggesting that the reduction of oxidative stress is not the only mechanism involved. Since augmented PKC activity has been shown in apoE−/− [29] and PKC activation can upregulate VEGF expression [4], it is conceivable that the observed in vivo effect of α-tocopherol on VEGF expression is due to both antioxidant and PKC inhibitory effects. It could be also argued that differences in the intestinal absorption, translated into different bioavailabilities for α and β-tocopherol, could account for the observed effects on VEGF expression. However, no differences were observed in plasma tocopherol between vitamin-treated apoE−/− mice, and the fact that both treatments similarly reduced vascular VEGFR-2 expression suggests that antioxidant mechanisms are fully responsible for VEGFR-2 downregulation while only accounting for partial VEGF reduction. This point would require further confirmation by measuring local production of reactive oxygen species and/or lipid peroxidation markers more specific and sensitive than TBARS (i.e. isoprostanes).

Interestingly, both treatments also achieved a similar significant reduction in circulating VEGF levels. Although the pathophysiologival meaning of the observed changes is not clear yet [13], given that systemic VEGF levels may not strictly reflect changes occurring in the vessel wall, its determination could become a valuable tool to investigate the...
relationship between hypercholesterolemia, atherosclerosis and angiogenesis, together with its pharmacological modulation.

Since VEGF seems to be necessary in the development of atherosclerosis [26] and treatment with angiogenesis inhibitors reduces both plaque growth and intimal neovascularization [19,30], the observed downregulation of VEGF and VEGFR-2 expression by the combination of vitamins C and E would be an additional mechanism by which these agents could prevent and limit atherosclerotic plaque progression [18,30]. Further studies should be conducted in a well-defined human hypercholesterolemic population to assess whether the effects of vitamins C and E supplementation observed in mice can be reproduced also in humans.

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