Repeated and long-term treatment with physiological concentrations of resveratrol promotes NO production in vascular endothelial cells

Satoru Takahashi1,2* and Yukiko Nakashima2

1Department of Immunobiology, School of Pharmacy and Pharmaceutical Sciences, Mukogawa Women’s University, 11-68 Kyuban-cho, Koshien, Nishinomiya, Hyogo 663-8179, Japan
2First Department of Biochemistry, School of Pharmaceutical Sciences, Kyushu University of Health and Welfare, 1714-1 Yoshino, Nobeoka, Miyazaki 882-0072, Japan

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Abstract

In the present study, we examined the effect of repeated and long-term treatment with resveratrol on NO production in endothelial cells as a model of routine wine consumption. Repeated treatment with resveratrol for 5 d resulted in an increase in endothelial NO synthase (eNOS) protein content and NO production in human umbilical vein endothelial cell (HUVEC) in a concentration-dependent manner. A significant increase in functional eNOS protein content was observed with resveratrol, even at 50 nM. In contrast, eNOS phosphorylation was not stimulated and inducible NO synthase (iNOS) was not detected after resveratrol treatment. Both eNOS protein and mRNA expression were promoted by 50 nM-resveratrol in a time-dependent manner. Increased eNOS mRNA expression in response to resveratrol was not decreased by an oestrogen receptor (ER) antagonist ICI182780, a PPARα inhibitor MK886 or a sirtuin inhibitor Salermide. However, a combination of ICI182780 and MK886 significantly inhibited resveratrol-induced eNOS mRNA expression. Salermide had no effect even in the presence of ICI182780 or MK886. These results demonstrate that resveratrol within the physiological range increases eNOS mRNA and protein expression through ER and PPARα activation, thereby promoting NO production in endothelial cells. eNOS induction might result from the accumulative effect of nanomolar concentrations of resveratrol. The present study results can account in part for the observation that cardiovascular benefits of red wine are experienced with routine consumption, but not with acute consumption.

Key words: Resveratrol: Endothelial cells: NO synthase: Gene expression

Epidemiological studies have demonstrated that dietary polyphenol intake including red wine consumption might improve endothelial function and reduce the risk of CVD(1–3). Resveratrol (trans-3,5,4’-trihydroxystilbene) is a polyphenol abundantly found in grapes, and is a key compound implicated in cardiovascular benefits associated with red wine consumption(3,4). Routine consumption of polyphenols is required for cardioprotection, suggesting that expression of specific genes in the vasculature is one of possible mechanisms of the beneficial effect of resveratrol.

NO in the vasculature is synthesised by endothelial NO synthase (eNOS) and is an important regulator of cardiovascular homoeostasis(3,5). NO prevents thrombogenic and atherogenic processes by exerting vasodilatory and anti-aggregatory effects. In animal models, the cardiovascular effects of resveratrol are mediated through NO(6–10). Therefore, whether resveratrol increases eNOS expression and NO production in endothelial cells has been investigated. Wallerath et al.(11) reported that exposure of cultured endothelial cells to resveratrol for 24–72 h up-regulates eNOS mRNA and protein expression, resulting in an increase in bioavailable NO. However, because the biological activities of resveratrol were observed between 10 and 100 μM, their experiments were not conducted under physiological conditions, in which concentrations were markedly higher than serum concentrations. Resveratrol concentrations in the blood were 20–50 nM after oral administration of 25 mg resveratrol to healthy volunteers(12,13). In addition, resveratrol at high concentrations (>10 μM) decreased cell viability and induced both apoptosis and necrosis(14,15). Nicholson et al.(16) investigated the effect of nanomolar concentrations of resveratrol on eNOS mRNA content in endothelial cells. Resveratrol treatment for 24 h increased
eNOS mRNA content. However, this increase was modest, and neither eNOS protein nor NO production was determined. Thus, it remains unclear whether resveratrol at physiological concentrations promotes NO production through eNOS induction in endothelial cells.

In the present study, we examined the effect of repeated and long-term treatment with resveratrol on functional eNOS expression and NO production in endothelial cells as a model of routine wine consumption. Here we provide evidence that resveratrol at nanomolar concentrations increases the expression of functional eNOS protein.

Materials and methods

Materials

Unless otherwise indicated, the reagents used in the present study and their sources are as follows: resveratrol, 3-(4,5-dimethyl-2-thiazoyl)-2,5-diphenyl-2H-tetrazolium bromide (MTT), N\textsuperscript{G}-nitro-L-arginine methyl ester (L-NAME; Sigma-Aldrich Company, St Louis, MO, USA); human umbilical vein endothelial cell (HUVEC) and cell culture media and supplies (Sanko Junyaku, Tokyo, Japan); porcine type I collagen (Nitta Gelatin, Osaka, Japan); MK886, Salermide and protease inhibitor mixture set III (Calbiochem, La Jolla, CA, USA); anti-eNOS antibody, anti-iNOS antibody and anti-\(\beta\)-actin antibody (BD Transduction Laboratories, Lexington, KY, USA); anti-peNOS antibody (phospho-Ser1177), secondary antibody linked to peroxidase, Phototope-HRP Western Blot Detection System and recombinant iNOS protein (Cell Signaling, Beverly, MA, USA); anti-eNOS antibody, anti-iNOS antibody and anti-\(\beta\)-actin antibody (BD Transduction Laboratories, Lexington, KY, USA); anti-peNOS antibody (phospho-Ser1177), secondary antibody linked to peroxidase, Phototope-HRP Western Blot Detection System and recombinant iNOS protein (Cell Signaling, Beverly, MA, USA); Immobilon-P membrane (Millipore, Bedford, MA, USA); ICI182780 (AstraZeneca Pharmaceuticals, London, UK); protein assay kit (Bio-Rad, Hercules, CA, USA); Sepasol-RNA (Nacalai Tesque, Kyoto, Japan). Resveratrol, Salermide, ICI182780 and MK886 were dissolved in dimethyl sulphoxide, with a final concentration of the solvent at 0.5 % in culture dishes. All other chemicals were of reagent grade.

Cell culture

HUVEC were maintained in endothelial cell basal growth-2 medium containing complete supplements (2% charcoal-treated fetal bovine serum, epidermal growth factor, basic fibroblast growth factor, vascular endothelial growth factor, insulin-like growth factor-1, heparin, hydrocortisone, ascorbic acid and GA-1000) at 37°C in 5% CO\textsubscript{2} and 95% air. Cells between passages five and ten were used in the experiments. All plates and dishes were coated with porcine type I collagen. HUVEC were seeded in 35-mm dishes for immunoblotting and NO production and in 60-mm dishes for quantitative RT-PCR. These cells were allowed to grow to approximately 50% confluence (day 0). Thereafter, the cells were incubated with resveratrol or vehicle for 1–5 d in endothelial cell basal growth-2 medium containing 10% supplements and 1 mg/ml bovine serum albumin. The medium was replaced with new medium every 24 h according to the schedule shown in Fig. 1.

Determination of NO synthase protein expression

NO synthase protein was detected by immunoblotting and quantified as previously described\textsuperscript{17}. HUVEC were lysed in 20 mM-Tris-HCl (pH 7.5), 1 mM-EDTA, 100 mM-NaCl, 10 mM-sodium orthovanadate, 10 mM-sodium fluoride, 10 mM-\(\beta\)-glycerophosphate, 1% Triton X-100, 0.5% sodium deoxycholate and protease inhibitor mixture. The lysates were placed on ice for 10 min and were then centrifuged at 10,000 rpm for 10 min at 4°C. The supernatants were recovered, and protein concentration was determined using bovine serum albumin as a standard. The supernatants (25 mg protein) were subjected to SDS-PAGE and were then blotted onto Immobilon-P membranes. The blots were incubated overnight with the primary antibody at 4°C and then probed with the peroxidase-conjugated secondary antibody. Immunoreactive proteins were visualised using an enhanced chemiluminescence kit. The intensity of each band was measured by densitometry.

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<th>Treatment</th>
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Fig. 1. Schematic diagram of repeated treatment with resveratrol. Culture medium containing vehicle or resveratrol and/or inhibitors was replaced with new medium every 24 h.
and then normalised by calculating the ratio of NO synthase to β-actin. Thereafter, relative expression was determined in comparison with the control cells.

**Determination of endothelial NO synthase mRNA expression**

Total RNA were extracted from HUVEC using Sepasol-RNA followed by DNase I digestion. Complementary DNA were synthesised using SuperScript III SuperMix (Invitrogen, Carlsbad, CA, USA). eNOS mRNA content was assessed by real-time PCR on a Roche LightCycler (Roche Diagnostics KK, Tokyo, Japan), as described previously (18). PCR primers used for eNOS and glyceraldehyde 3-phosphate dehydrogenase were 5'-CACATGTTTGCTCGGG-3' and 5'-GAGGGGCGTCTCCAGATTAAG-3', and 5'-ATGCCAGCCTCAAGGGCTAGAAAAAGA-3' and 5'-GTTGCCTGTACCAAATTCGTTGTC-3', respectively. eNOS mRNA expression was normalised to glyceraldehyde 3-phosphate dehydrogenase mRNA expression and then relative eNOS mRNA expression was determined in comparison with the control cells.

**Determination of NO production**

NO production was determined as generation of nitrite (14). HUVEC were incubated in PBS containing 10 mM-HEPES-NaOH (pH 7.4), 0.1 mM-ATP and 1 mg/ml bovine serum albumin for 30 min and then the medium was sampled. In experiments on eNOS inhibition 5 mM-L-NAME was added. NO was determined as nitrite by colorimetry following the Griess reaction. NO released into the medium was converted to nitrate and nitrite when left overnight at 4°C, and then nitrate was completely oxidised by nitrate reductase to nitrite. The amount of nitrite was measured using a NOx analyser (ENO-200, EICOM, Kyoto, Japan).

**Determination of cell viability**

Cell viability was determined by mitochondrial function assay, as described previously (14). Mitochondrial function was assessed by the colorimetric method using MTT. After treating with resveratrol for 5 d the cells were incubated in the culture medium containing 1 mg/ml MTT for 2 h at 37°C, the formed formazan was extracted with isopropanol containing 40 mM-HCl. The absorbance at 595 nm in the extract was measured.

**Statistical analysis**

Data are presented as means with their standard errors. Statistical differences in the concentration–response study were evaluated by Dunnett’s multiple comparison test. Student’s t test was used to compare the two groups. A P value of <0.05 was regarded as significant.

**Results**

In the present study, we examined whether resveratrol at physiological concentrations enhances eNOS protein expression in endothelial cells. Repeated treatment with resveratrol for 5 d resulted in an increase in eNOS protein content in HUVEC in a concentration-dependent manner (Fig. 2). Resveratrol-induced eNOS protein expression started increasing from 10 nm and significantly increased at 50 and 200 nm (P<0.01). This expression was 2.5-fold at 50 nm and 3.1-fold at 200 nm compared with the control cells. In contrast,
iNOS protein was not detected in the control or resveratrol-treated cells. Since Ca$^{2+}$-independent eNOS activation occurs by eNOS phosphorylation at Ser1177 (19, 20), the phosphorylation state of the increased eNOS protein was determined. Resveratrol increased phospho-eNOS (peNOS) content in a concentration-dependent manner.

To confirm that the increased eNOS protein was functionally active, we determined NO production after HUVEC were exposed to nanomolar concentrations of resveratrol for 5d (Fig. 3). NO released into the medium was converted to nitrite, which was then quantified. Resveratrol stimulated NO production in a concentration-dependent manner. In comparison with the control cells, a significant increase in NO production was observed at 50 nM (2.2-fold) and 200 nM (2.4-fold; P<0.05). L-NAME potently inhibited NO production in the control and resveratrol-treated cells, indicating that the measured NO was synthesised by NOS (P<0.01 in control and 200 nm-resveratrol; P<0.001 in 10 and 50 nm-resveratrol between vehicle and L-NAME).

In the present study, we further examined the time-dependent change in resveratrol-induced eNOS protein expression at 50 nM (Fig. 4). HUVEC gradually grew and reached confluence on day 5 under our culture condition. Neither cell number nor cell density differed between resveratrol- and vehicle-treated groups during the 5-d period. In fact, cell viability on day 5 was not reduced by resveratrol; 105 % at 50 nM and 94 % at 200 nM compared with the control cells. eNOS protein expression was not altered by resveratrol on day 1, whereas the expression was significantly enhanced after exposure to resveratrol for more than 3d. The increase in eNOS protein content was 2.4-fold on day 3 and 3.1-fold on day 5, compared with that in the cells on day 0 (P<0.01 in resveratrol-treated cells between day 0 and day 3, and between day 0 and day 5). eNOS protein was significantly induced even when the cells were treated with resveratrol from day 2 to day 5 (group 2–5; P<0.05), indicating that the effect of resveratrol on eNOS protein expression did not depend on cell density or number. In contrast, when the cells were treated with resveratrol for 3d followed by depletion for 2d, eNOS protein content almost returned to the control level (group 0–3–5). eNOS protein expression was not affected by vehicle treatment during the 5-d period.

Resveratrol-induced eNOS mRNA expression at 50 nM also increased in a time-dependent manner (Fig. 5). Resveratrol increased eNOS mRNA content by approximately 1.6-fold even on day 1 when eNOS protein expression was not altered. Thereafter, resveratrol significantly increased eNOS mRNA content by 2.6-fold on day 3 (P<0.05) and 2.9-fold on day 5 (P<0.01), compared with that in the cells on day 0. However, resveratrol depletion from day 3 abolished the stimulatory effect on day 5.
mRNA expression in the resveratrol (50 nM)-treated cells was only 1.6-fold of that in the control cells, consistent with the results of the present study. However, the significance of the slight increase in eNOS mRNA expression remained elusive because Nicholson et al. did not measure eNOS protein content or NO production. The present study showed that eNOS protein content was elevated from day 3, but not day 1, by resveratrol treatment at 50 nM. In addition, the increased eNOS mRNA and protein contents returned to the control levels within 2 d of resveratrol depletion from the culture medium. These results indicate that long-term and repeated treatment is required for the physiological effect of resveratrol on eNOS protein expression in endothelial cells. The accumulative effect of nanomolar concentrations of resveratrol might lead to an increase in eNOS protein level.

The amount of peNOS protein was also increased by resveratrol. However, the phosphorylation itself was not affected by resveratrol, as the ratio of peNOS content to total eNOS content was not different between the control and resveratrol-treated cells. Akt- and AMP-activated kinases are responsible for eNOS phosphorylation at Ser1177(19,20,21). Resveratrol treatment for 5 d had no effect on the activated state of these protein kinases in HUVEC (data not shown). The increase in peNOS protein might be associated with the increased eNOS protein. These results demonstrate that resveratrol-stimulated NO production is primarily due to quantitative alteration of total eNOS protein.

Resveratrol increased eNOS protein expression following an increase in eNOS mRNA content in endothelial cells,
indicating that resveratrol-stimulated induction of eNOS protein implicates eNOS gene expression. Resveratrol has multiple target molecules (25, 26), and transcriptional regulators such as ER, sirtuin and PPARα are among these targets. Combination of IC182780 and MK886 significantly reduced resveratrol-induced eNOS mRNA expression, although the inhibitors individually had no effect. In contrast, Salermide failed to affect the increased eNOS mRNA expression even in the presence of IC182780 or MK886. These results suggest that both ER and PPARα, but not sirtuin, might be involved in resveratrol-induced eNOS protein induction. Oestradiol up-regulates eNOS gene expression through ER activation in endothelial cells (27), and resveratrol as a phyto-oestrogen can activate ER in endothelial cells (20). Goya et al. (29) reported that PPARα agonist fenofibrate elevates eNOS mRNA and protein contents, resulting in an increase in NO production in endothelial cells. PPARα in endothelial cells is also activated by resveratrol (23). These previous results are consistent with the present study findings, but resveratrol-stimulated ER or PPARα alone is unable to up-regulate eNOS mRNA expression. One explanation is that the affinities of resveratrol for ER and PPARα are lower than those of the potent agonists oestradiol and fibrates. Synergy may exist between the ER and PPARα pathways for resveratrol-induced eNOS up-regulation. In addition, other mechanisms might lead to resveratrol-induced eNOS expression, because an increase in eNOS mRNA content was observed even when both ER and PPARα were inhibited.

In addition to eNOS induction, it is also possible that the antioxidant property of resveratrol might be involved in prevention of a decrease in NO bioavailability. Oxidants inhibit eNOS activity by reducing eNOS dimer stability (10,30). Resveratrol is able to stabilise eNOS dimer by scavenging oxidants. Furthermore, resveratrol up-regulates superoxide dismutase and catalase, which inhibit inactivation of NO by the oxygen radicals (10). Overall, resveratrol might exert cardiovascular protection by up-regulating eNOS protein and maintaining NO bioavailability.

We conclude that resveratrol within the physiological range increases eNOS mRNA and protein expression through ER and PPARα activation, thereby promoting NO production in endothelial cells. eNOS induction might result from the accumulative effect of nanomolar concentrations of resveratrol. The present study results could account in part for the observation that the cardiovascular benefits of red wine are experienced with routine consumption, but not with acute consumption. These findings are helpful in understanding the molecular basis of the beneficial effect of resveratrol.

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