

Differential regulation of proliferation, cell cycle control and gene expression in cultured human aortic and pulmonary artery endothelial cells by resveratrol

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Received July 13, 2010; Accepted August 9, 2010

DOI: 10.3892/ijmm_00000521

Abstract. Resveratrol is a grape polyphenol with cardioprotective attributes, supported in part by its demonstrated anti-mitogenic, apoptosis-inducing and gene modulatory activities in various cell types known to play an integral role in atherogenesis. To test whether resveratrol exerts similar effects on systemic and pulmonary vasculature, cells derived from different anatomical sites, cultured human aortic and pulmonary artery endothelial cells, respectively denoted HAECs and HPAECs, were exposed to resveratrol for assessment of effects on proliferation, cell cycle distribution, induction of apoptosis, and specific gene expression. Resveratrol inhibited cell proliferation in a time- and dosedependent manner in HAECs and HPAECs, accompanied by disruption of cell cycle control and progression as assayed by flow cytometry. Analysis of gene changes in resveratroltreated endothelial cells by RT-PCR showed suppression of nitric oxide synthase (eNOS) and preproendothelin-1 (ppET-1) expression in both cell types. To discover group gene alterations resulting from exposure to resveratrol, changes in mRNA levels were determined by human signal transduction pathway finder cDNA array analysis. The results showed that resveratrol up-regulated levels of cyclin-dependent kinase inhibitor p57, egr-1, forkhead box A2 and c-jun in HAECs, and elevated expression of cathepsin D, ICAM-1, c-jun and patched 1 in HPAECs. In addition, treatment by resveratrol also resulted in attenuated expression of bcl-xl, fibronectin-1, HIP, mdm2, PIG3 and WSB1/SWIP-1 in HAECs, and CDX1, engrailed homolog 1, FASN, fibronectin-1, forkhead box A2, Hoxa-1, hsp27, PIG3, ELAM-1/E-selectin and WSB1/SWIP-1 in HPAECs. These results suggest that resveratrol acts by distinct and overlapping signaling pathways and mechanisms in HAECs and HPAECs, further supporting the notion that the cardioactive activities and effects of this

Key words: resveratrol, human aortic endothelial cells, human pulmonary aortic endothelial cells, cell proliferation, cell cycle control

grape polyphenol are contingent upon or influenced by the vascular bed of origin.

Introduction

Atherosclerosis (AS) and pulmonary arterial hypertension (PAH) have a pronounced and significant association with cardiovascular diseases (CVD), and both are considered major causes of morbidity and mortality in developed countries (1-4). CVD currently claims the lives of 500,000 men and women annually in the USA. However, in spite of concerted efforts in education, research and prevention programs actively introduced over the past few decades to curtail CVD-related death rates, evidence of improved statistics is simply lacking. The previously identified major risk factors for CVD are cigarette smoking, hypertension, and hypercholesterolemia. Interestingly, these may not be sufficiently robust or reliable as predictors of new cases of CVD (5,6) suggesting that protective factors exist that act to counteract and mitigate the deleterious consequences of the major CVD risk factors. In support of this possibility, recent data have shown that certain diet and dietderived chemicals may play a role in reducing and ameliorating the damage induced by exposure to risk factors of CVD (7-10). Therefore, the discovery and identification of protective dietary factors and elucidation of the mechanisms by which they confer cardioprotection may be envisaged to shed light on new CVD management strategies.

Resveratrol is a polyphenol present in abundance in several foods and red wine, which has been intensively studied in recent years stemming from the 'French paradox', the population-based epidemiological phenomenon showing an inverse correlation between the risk for CVD and low-tomoderate consumption of red wine with meals, possibly attributing to phytochemicals contained in red wine, e.g., resveratrol (11-14). The cardioprotective role of resveratrol has been supported by studies from this and other laboratories showing that this grape polyphenol (i) suppresses LDL oxidation (15,16), (ii) reduces SMC proliferation (17-19), (iii) inhibits platelet aggregation (20-22), (iv) attenuates intimal thickening in rabbits fed a hypercholesterolemic diet and subjected to endothelium denudation (23,24), and (v) induces nitric oxide synthase (NOS) in cultured bovine pulmonary artery endothelial cells (BPAEC) (25-27). To date, only scant data and mechanisms currently exist regarding the effects of this grape polyphenol in human endothelial cells,

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notwithstanding that these cells are integrally involved in controlling the homeostasis of the vasculature and the initiation of atherosclerosis (12). Because vascular diseases show an anatomically defined focal distribution, it is of interest to determine whether the spatiotemporal response, regulation, and function of endothelial cells to CVD, risk and protective factors, are contingent upon the source of the vasculature from which endothelial cells originate (28-31). Similarly, it has not been established whether the cardioactive effects of resveratrol might have subtle or marked dependence on the anatomical origin of endothelial cells. In the present study, we studied the responses of HAECs and HPAECs to resveratrol, in the context of suppression of cell growth, cell cycle phase distribution, induction of apoptosis, and changes in gene expression.

Materials and methods

Materials. HAECs and HPAECs were purchased from Lonza Walkersville Inc. (Walkersville, MD, USA), and were cultured and propagated using Clonetics[®]EGM[®]-2 BulletKit[®] containing endothelial cell basal medium-2 and growth supplements, as recommended by the manufacturer (Lonza Walkersville Inc.). Cells were maintained in a CO₂ incubator and media were changed every 3-4 days. Resveratrol was dissolved in dimethyl sulfoxide (DMSO) as a 12.5 mM stock and kept in aliquots at -20°C. All other chemicals and solvents used were of analytical grade.

Cell culture and proliferation assay. HAECs and HPAECs maintained in culture media as described were grown to >70% confluence, and then split 1:4 into T-25 flasks containing fresh media. Following an overnight incubation, cells were exposed to increasing concentrations of resveratrol as specified in the text. Cell numbers were determined at the indicated times by trypan blue exclusion, as previously described (32-34).

Cell cycle analysis. HAECs and HPAECs were treated with resveratrol (0, 1, 10, and 100 μ M) for 24 or 48 h, and cell cycle phase distribution was assayed by flow cytometry as described (33,35,36). The cellular DNA content was obtained and the percentage of cells in the respective phases (G₁, S and G₂/M) of the cell cycle was quantified.

RNA extraction and reverse transcription-polymerase chain reaction (RT-PCR). Total cellular RNA was isolated from control and resveratrol treated cells, using Trizol reagent (Invitrogen, Carlsbad, CA) according to protocols provided by the manufacturer. The first strand cDNA was synthesized from 2 µg of total RNA, using Superscript[™] RNase H-reverse transcriptase (Invitrogen, Carlsbad, CA). The PCR reaction was performed using 1 μ l cDNA synthesis reaction mixture according to the following conditions, eNOS, 94°C 1 min, 60°C 1 min, 72°C 1 min, 32 cycles; ppET-1, 94°C 1 min, 60°C 1 min, 72°C 1 min, 34 cycles; GAPDH, 94°C 1 min, 60°C 1 min, 72°C 1 min, 26 cycles, and using the following primer sets: eNOS, forward and backward primer sets, 5'-CCC TTC AGT GGC TGG TAC AT-3'/5'-GAA AAC AGG AGT GAG GCT GC-3', expected size 327 bp; ppET-1, forward and backward primer sets, 5'-TCC AAG AGA GCC TTG GAG AA-3'/

5'-ATG GAA GCC AGT GAA GAT GG-3', expected size 565 bp; GAPDH, forward and backward primer sets, 5'-CCA CCC ATG GCA AAT TCC ATG GCA-3'/5'-TCT AGA CGG CAG GTC AGG TCC ACC-3', expected size 598 bp. The PCR products were resolved on 1.2% agarose gels. The relative intensity of fragments with the expected sizes was determined using the Innotech imaging system (37,38).

cDNA array analysis. Total RNA was isolated from day 2 control and 100 μ M resveratrol-treated cells. RNA (4 μ g) was used as the template for biotinylated cDNA synthesis. Hybridization of biotinylated cDNA to immobilized genespecific cDNAs and detection of hybridization signals by chemiluminescence were according to the manufacturer's protocol (Superarray, Bethesda, MD), as described (34,38,39). Briefly, membrane was prehybridized in solution containing 100 mg/ml heat-denatured, sheared salmon sperm DNA for 2 h at 60°C followed by overnight incubation with the denatured cDNA probe, with shaking at 10 rpm/min, 60°C. Membrane was washed, blocked for 40 min with solution Q provided by the manufacturer, and incubated with AP-streptavidin for 10 min at room temperature. Next, the membrane was washed and rinsed with CDP-Star chemiluminescent substrate followed by exposure to X-ray film. The data were analyzed using GEArray Analyzed software.

Results

Effects of resveratrol on growth and cell cycle distribution in cultured HAECs. To obtain information on overall cell response to resveratrol by HAECs, the growth modulatory effects of resveratrol were studied. Cells were incubated with 0 or $100 \,\mu M$ resveratrol for 24 or 48 h, or with increasing concentrations of resveratrol $(0, 1, 10 \text{ or } 100 \,\mu\text{M})$ for 48 h. Cell number was determined by trypan blue exclusion assay. Exposure to resveratrol resulted in time- and dose-dependent inhibition of growth, compared with controls. Growth suppression was ~48% at 24 h exposure to 100 μ M resveratrol (Fig. 1A). By 48 h an ~58 and ~73% diminution of cell growth was observed in cells treated with 10 and 100 μ M resveratrol, respectively (Fig. 1B). To test whether resveratrol-induced growth suppression in HAECs might involve alterations in cell cycle control, flow cytometry studies were performed. Cells treated with 100 μ M resveratrol for 24 or 48 h both showed an increase in G₁ phase cell population (~75% in control vs. ~89% in treated cells) accompanied by a concomitant reduction in the S phase cell population (~12% in control vs. ~2.5% in treated cells), and a slight increase in induction of apoptosis (from ~2.3 in control to 3.6% in treated cells) (Fig. 1C). In cells treated with 0, 1, 10 or 100 μ M resveratrol for 48 h there was a similar increase in G₁ phase cell population accompanied by a concomitant reduction in the S phase cell population by 1 and 100 μ M resveratrol treatment. However, 10 μ M resveratrol treatment resulted in a reduction in G₁ phase accompanied by an increase in S phase (Fig. 1D). These results suggest that resveratrol controls mitogenesis in HAECs.

Effects of resveratrol on growth and cell cycle distribution in cultured HPAECs. The response of HPAECs to resveratrol was also investigated with respect to growth suppression and





Figure 1. Effects of resveratrol on growth and cell cycle distribution in cultured HAECs. (A) Growth of HAECs exposed to 0 or 100 μ M of resveratrol for 24 or 48 h, respectively. Results represent the mean of triplicate experiments ± SD. (B) Growth of HAECs treated for 48 h with increasing resveratrol (1, 10, 100 μ M). Results show the mean of triplicate experiments ± SD. (C) Effects of resveratrol on the changes of cell cycle phase distribution in HAECs. Cells were treated with 0 or 100 μ M of resveratrol for 24 or 48 h, and analyzed by flow cytometry. The percentage of cells in G₁, S, and G₂ phases were determined and are presented as bar graphs. (D) Changes of cell cycle phase distribution in HAECs treated with 0, 1, 10 or 100 μ M of resveratrol for 48 h. The effects of resveratrol on cell cycle distribution, respectively, as the percentage of cells in G₁, S, and G₂ phases are presented as bar graphs.



Figure 2. Effects of resveratrol on growth and cell cycle distribution in cultured HPAECs. (A) HPAECs were exposed to 0 or 100 μ M of resveratrol for 24 or 48 h, and growth in control and treated cells are shown as the mean of triplicate experiments ± SD. (B) Exposure of HPAECs for 48 h with increasing dose (0, 1, 10, 100 μ M) of resveratrol, and results on growth are presented as the mean of triplicate experiments ± SD. (C) Effects of resveratrol on the changes of cell cycle phase distribution in HPAECs. Cells treated with 0 or 100 μ M of resveratrol for 24 or 48 h were analyzed by flow cytometry, and the percentage of cells in G₁, S, and G₂ phases are shown as bar graphs.

Genebank	Gene name	Description	Pathway	Regulation
U22398	p57Kip2	Cyclin-dependent kinase inhibitor 1C (p57, Kip2)	TGFb	Ť
X52541	egr-1	Early growth response 1	Mitogenic, creb, phospholipase C	ſ
AB028021	Forkhead box A2	Hepatocyte nuclear factor 3, ß	Hedgehog	Ť
J04111	c-jun	v-jun avian sarcoma virus 17 oncogene homolog	Mitogenic, wnt, survival, calcium and protein kinase C, phospholipase C	ſ
Z23115	bcl-xl	Homo sapiens BCL2-like 1 (BCL2L1)	Survival	Ŷ
X02761	Fibronectin-1	Fibronectin 1	Survival	Ļ
AY009317	HIP	Hedgehog-interacting protein	Hedgehog	Ļ
Z12020	mdm2	Mouse double minute 2, human homolog of p53-binding protein	p53	Ŷ
NM_004881	PIG3	Quinone oxidoreductase homolog	p53	\downarrow
NM_015626	WSB1/SWIP-1	SOCS box-containing WD protein SWiP-1 (SWIP1)	Hedgehog	Ŷ

Table I. Identity of specific genes regulated by resveratrol in HAEC.

cell cycle phase distribution. Cells were treated with different concentrations of resveratrol for variable time periods and growth was determined. Exposure to resveratrol resulted in time- and dose-dependent inhibition of proliferation, relative to controls, yielding ~42% growth suppression in 24 h, 100 μ M resveratrol-treated cells (Fig. 2A). In 48-h exposed cell cultures, an ~15, ~47 and ~62% diminution of cell growth was observed in 1, 10 and 100 μ M resveratrol treated samples (Fig. 2B). To further validate the anti-proliferative activity of resveratrol on HPAECs, the changes in cell cycle distribution were assayed by flow cytometry. Although there was no observed time-dependent difference on cell cycle distribution following treatment with 100 μ M resveratrol, exposure to the grape polyphenol nevertheless increased G₁ phase cell population (~74% in control vs. ~83.5% in treated cells), in concomitance with a comparable reduction in the S phase cell population (~12% in control vs. ~3.5% in treated cells) (Fig. 2C). Moreover, whereas cells treated with 10 μ M resveratrol for 48 h showed restriction in the S phase, treatment with 100 μ M resveratrol induced a G₁ phase arrest (Fig. 2D), suggesting that resveratrol also plays a regulatory role in the control of mitogenesis in HPAECs.

Effects of resveratrol on eNOS and ppET-1 gene expression in cultured HAECs and HPAECs. The expression of endothelial cell specific genes, respectively, eNOS and ppET-1 (40) was assayed to determine whether treatment by resveratrol elicited a differential gene change in HAECs and HPAECs. To this end, levels of eNOS and ppET-1 mRNA in control and resveratrol-treated HAECs and HPAECs were analyzed by RT-PCR. Time-dependent decreases in eNOS and ppET-1 expression were observed in 100 μ M resveratrol-treated HAECs. By contrast, in HPAECs, resveratrol only induced time-dependent inhibition in eNOS expression while the suppression of ppET-1 by resveratrol was only seen in 24-h



Figure 3. Control of eNOS and ppET-1 gene expression by resveratrol in HAECs and HPAECs. (A) Cells were treated with 0 or 100 μ M of resveratrol for 24 or 48 h, harvested and analyzed for changes in eNOS and ppET-1 expression, by RT-PCR as detailed in Materials and methods. (B) Effects of increasing concentrations of resveratrol (0, 1, 10 and 100 μ M) on eNOS and ppET-1 expression. Control and 48 h resveratrol-exposed HAECs and HPAECs were harvested and used for determination of eNOS and ppET-1 mRNA level changes by RT-PCR. In these experiments, the expression of glyceraldehyde-3-phosphate dehydrogenase (GAPDH) was used as an internal control to verify equivalent RNA loading.

treated cells (Fig. 3A). A dose-dependent study revealed that the reduction of eNOS was only seen in both HAECs and HPAECs treated with 100 μ M resveratrol for 48 h while resveratrol reduced ppET-1 expression only in HAECs but not in HPAECs (Fig. 3B). These results suggest that resveratrol differentially controls eNOS and ppET-1 expression, in a cell type-dependent manner.

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Genebank	Gene name	Description	Pathway	Regulation
M11233	Cathepsin D	Cathepsin D (lysosomal aspartyl prostease)	Estrogen, retinoic acid	î
NM_000201	ICAM-1	Intercellular adhesion molecule 1 (CD54), human rhinovirus receptor	NF- κ B, phospholipase c	↑
J04111	c-jun	v-jun avian sarcoma virus 17 oncogene homolog	Mitogenic, wnt, survival, calcium and protein kinase C, phospholipase C	ſ
U43148	Patched 1	Patched (Drosophila) homolog 1	Hedgehog	Ť
NM_001804	CDX1	<i>Homo sapiens</i> caudal type homeo box transcription factor 1 (CDX1)	Retinoic acid	Ŷ
NM_001426	Engrailed homolog 1	Engrailed homolog 1	Hedgehog; Retinoic acid	Ŷ
U26644	FASN	Fatty acid synthase	Insulin	Ŷ
X02761	Fibronectin-1	Fibronectin 1	Survival	Ŷ
AB028021	Forkhead box A2	Hepatocyte nuclear factor 3, ß	Hedgehog	↓
NM_005522	Hoxa-1	Homo sapiens homeo box A1 (HOXA1)	Retinoic acid	Ŷ
Z23090	Hsp27	Heat shock 27 kD protein	Stress	Ŷ
NM_004881	PIG3	Quinone oxidoreductase homolog	p53	↓
M30640	ELAM-1/E-selectin	Human endothelial leukocyte adhesion molecule 1 (ELAM1) mRNA	LDL	Ŷ
NM_015626	WSB1/SWIP-1	SOCS box-containing WD protein SWiP-1 (SWIP1)	Hedgehog	Ŷ

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Effects of resveratrol on expression of cell signal transduction genes in HAECs and HPAECs. To define group gene changes in HAECs and HPAECs that respond to exposure to resveratrol treatment, human Signal Transduction PathwayFinder cDNA GEArray was used since it was designed to explore aggregate gene expression changes associated with 18 signal transduction pathways. To test the feasibility and validity of this approach, cells were treated with 100 μ M resveratrol for 2 days and used in the analysis. In HAECs, p57Kip2, egr-1, forkhead box A2 and c-jun showed up-regulation with treatment by 100 μ M of resveratrol. In contrast, 6 genes (bcl-xl, fibronectin-1, HIP, mdm2, PIG3 and WSB1/SWIP-1) having an integral role on different facets of control of cell signaling were down-regulated by resveratrol (Table I). In HPAECs, resveratrol resulted in increased expression of cathepsin D, ICAM-1, c-jun and patched 1 gene expression, while concurrently suppressing the levels of 10 genes, respectively, CDX1, engrailed homolog 1, FASN, fibronectin-1, forkhead box A2, Hoxa-1, hsp27, PIG3, ELAM-1/E-selectin and WSB1/SWIP-1 (Table II). These results provide further evidence that there is a differential control of signal transduction by resveratrol and this control is cell type-dependent.

Discussion

Resveratrol shows cardioprotective attributes based on *in vitro* and *in vivo* data obtained using tissue culture cells and experiments involving animals (12,22,41,42). However, relatively fewer studies have addressed the activities of resveratrol on vascular components, including endothelial cells

derived from humans, particularly those derived from different anatomical sites. In this study, we investigated whether resveratrol exerts similar effects in two normal human endothelial cells, HAECs and HPAECs, in terms of their growth and gene regulatory activities. Results of this study provide several hitherto unreported observations lending further support for the cardioprotective potential of this grape polyphenol: (i) resveratrol significantly and differentially inhibits the proliferation of both HAECs and HPAECs (Figs. 1 and 2), (ii) resveratrol alters cell cycle distribution in HAECs and HPAECs (Figs. 1 and 2), (iii) resveratrol inhibits the expression of eNOS and ppET-1 in HAECs but only exerted transient inhibitory effect on ppET-1 expression in HPAECs (24 but not 48 h of exposure to resveratrol) (Fig. 3A), (iv) resveratrol induces distinct and overlapping changes in signal transduction pathways in HAECs and HPAECs. Taken together, these results are consistent and support the interpretation that resveratrol exerts cardioactive affects on normal endotheial cells, possibly targeting differential mechanistic control on cell signaling pathways that impinge on the control of cell growth, thus befitting the rubric of efficacy as a cardioprotective agent.

The vascular smooth muscle and endothelial cells both play a significant role in the maintenance of vascular tone in the adult blood vessel (12,22,40). Under certain pathological conditions aberrant cell proliferation in smooth muscle or artery endothelial cells may lead to disorder of vascular remodeling as an antecedent to the pathological sequelae of CVD. Previously, we have shown that resveratrol inhibited human aortic smooth muscle cell proliferation (40). The results of this study show that resveratrol also suppressed proliferation of HAECs and HPAECs in concert with disruption of cell cycle progression. Accordingly, restriction of HAECs growth by resveratrol could attenuate the development of AS plaques. Similarly, control of proliferation of structurally and functionally altered, subclincal abnormal pulmonary endothelial cells could prevent formation of plexiform lesions associated with remodeled pulmonary vasculature found in pulmonary hypertensive specimens (40,43-45) both of which may contribute towards a potentially efficacious preventive strategy for CVD.

Impaired nitric oxide (NO) synthesis has been reported in AS and the primary source of vascular NO is likely to come from endothelial cells (12,22). Therefore, in this study we investigated the effects of resveratrol on eNOS expression. Treatment of HAECs and HPAECs by resveratrol resulted in down-regulation of eNOS gene expression, in contrast to our previous observation showing that resveratrol induced eNOS protein expression in BPAECs (26). While the significance and basis for these discordant observations remain to be investigated in detail, they clearly point to complex, cell type and species-dependent control of eNOS, by exposure to cardioprotective cues, such as resveratrol. In addition to eNOS, endothelin (ET) also plays an important role in pathogenesis of CVD particularly in congestive heart failure. Accordingly, in this study we also investigated the effects of resveratrol on ppET-1 mRNA expression. Our results showed that resveratrol inhibited ppET-1 expression. Moreover, the inhibition of ppET-1 was more pronounced in HAECs than HPAECs, further supporting that the cardioprotective effects of resveratrol may be cell type-dependent.

Since the differentially effective growth response in HAECs and HPAECs elicited by resveratrol may be attributed to functionally related gene changes, studies employing the signal transduction pathway finder cDNA array were performed to test this possibility, as results derived from array analysis are expected to provide a panoramic mechanistic framework relevant to the resveratrol induced signaling changes. Using this approach, distinct and/or overlapping signal transduction pathways involved in control HAECs and HPAECs cell growth and gene expression by resveratrol were identified, validating that resveratrol indeed triggered cell type-dependent gene regulatory effects (Tables I and II). Notably, in HAECs, this grape polyphenol elicited growth response and gene control changes, in part mediated through Hedgehog, p53 and survival pathway, whereas by contrast, resveratrol induced changes in HPAECs possibly through the Hedgehog and retinoic acid pathways.

In conclusion, the results of this communication suggest that resveratrol controls proliferation of human endotheial cells irrespective of their anatomical origin, albeit might exert endothelial cell type-specific affects utilizing distinct and preferential signaling pathways. Overall, though, the data lend credence to the categorization of resveratrol as an example of diet-derived phytochemicals capable of restricting endothelial cell growth and hence exhibits the potential of being developed as an efficacious preventive strategy for CVD.

Acknowledgements

Studies reported herein were supported in part by the Intramural Sponsored Research Program of New York Medical College and by Phillip Morris USA Inc. and Phillip Morris International (to J.M.W.).

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