

# **Allicin Reverses Diabetes-Induced Dysfunction of Human Coronary Artery Endothelial Cells Daniel Horuzsko<sup>1</sup>, Margeaux LaCavera<sup>1</sup>, Handong Ma<sup>1</sup>, Yan Wu<sup>1</sup>, Shu Zhu<sup>1</sup>, Richard White<sup>1</sup>**

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### ABSTRACT

Cardiovascular disease (CVD) is the leading cause of death in the United States, and is the major source of morbidity and mortality associated with diabetes mellitus. Because the incidence of diabetes continues to increase, reducing the risk of CVD in diabetes will continue to be a major focus of cardiovascular research. An early manifestation of diabetesinduced CVD is dysfunction of the vascular endothelium, as indicated by depressed production of NO. Our findings now demonstrate depressed activity of endothelial nitric oxide synthase (eNOS) in diabetes, and suggest that treating human coronary artery endothelial cells with allicin, the major bioactive organosulfur component in garlic extract, can restore NO production in these cells. Coronary artery endothelial cells (Lonza) were obtained from control (HCAEC) or diabetic (DHCAEC) donors, and NO production was measured by fluorescence microscopy via 4,5-diaminofluorescein diacetate. On average, NO production was depressed by 12.9% in DHCAEC compared to controls. Treating these cells for 20 minutes with 4µM allicin restored NO production by 32.9%. Further, immunoblot studies revealed that diabetes decreased expression of eNOS protein by 20.3%; however, allicin was able to reverse this effect of diabetes. On average, eNOS expression was increased by 26% by overnight exposure to 5µM allicin. Taken together, these data indicate that allicin improves endothelium-dependent NO production in diabetes by enhancing the expression and/or activity of eNOS in human coronary artery endothelial cells. These studies further suggest that this improved endothelial function likely contributes to the established health benefits of garlic consumption (e.g., lowering blood pressure), and also suggests a natural means of reducing the devastating consequences of diabetes on CVD. Future experiments are needed to identify the mechanism of allicin action on eNOS and in vascular endothelial cells.

### INTRODUCTION

Diabetes mellitus is a consistent rising public health burden in the 21<sup>st</sup> century, and remains a major risk factor for the development of CVD. Currently, diabetes affects over 125 million people in the western world, and estimates of its prevalence are expected to surpass 300 million by 2025 due to a combination of poor diet and lifestyle. Natural garlic (Allium) sativum) supplementation has been utilized as a remedy for thousands of years in treating various chronic diseases. Allicin (2-propene-1-sulfonothioic acid S-2 propenyl ester, diallyl thiosulfinate) is the primary active component in garlic, and current literature supports its therapeutic effects on lowering blood pressure, decreasing blood glucose levels, reducing atherosclerotic plaque formation, and modulating inflammation. This organosulfur compound rapidly crosses the cellular membrane and reacts with sulfhydryl groups on cellular proteins or with glutathione. It has been demonstrated that allicin increases intracellular levels of glutathione and phase II detoxifying enzymes that include glutamatecysteine ligase and heme oxygenase-I in vascular endothelial cells. We tested the hypothesis that allicin could reverse diabetes-induced inhibition of NO production in human coronary artery endothelial cells, possibly by decreasing oxidative stress to lower CVD risk.

### METHODS

<u>Cell Culture</u>: Human coronary artery endothelial cells (HCAEC) from healthy, Type-I, and Type II diabetic cell lines(DHCAEC-I; DHCAEC-II) were grown in EBM-2 basal media supplemented with EGM-2 MV (Lonza). Passage 4 cells were thawed and used for treatment. Cells were changed to phenol-free media 24 hours prior to imaging. Allicin Treatment: Cells were incubated in 10µM 4,5-Diaminofluorescein diacetate (DAF-2DA) for 15 minutes at 37°C. Cells were then washed with KREBS solution and incubated for 20 minutes at 37° C. Afterwards, cells were incubated with L-NAME and images were obtained with EVOS microscope at 10X, 20X, 40X and analyzed using ImageJ software. Immunoblotting: Cell lysates (20µg of protein) were analyzed by 8% Tris Glycine SDS polyacrylamide gel electrophoresis and transferred to PVDF membrane. The membranes were blocked with 5% milk + TBST 0.1% for 1 hour and then probed overnight with a antieNOS antibody (1:10000). Membranes were then incubated with species-specific HRPconjugated secondary antibody (1:5000) followed by detection with ECL substrate. Quantitation of proteins was determined by densitometry analysis of scanned films using ImageJ software.

Mass Spectrometry: Proteins were extracted from cell lines and equal amounts were immunoprecipitated with anti-eNOS antibody or control goat IgG. The immune complexes were immobilized on the protein A/G ultralink resin. Trypsin digestion and MS analysis were performed at the proteomics and metabolomics core facility at the Georgia Cancer Center. Scaffold (version Scaffold\_4.8.9, Proteome Software, Inc.) was used to validate LC-MS and LC-MS/MS-based peptides and protein identification. Peptide identifications were accepted if they could be established at >95.0% probability as specified by the Peptide Prophet algorithm.



RESULTS

analysis, respectively.

