Research Paper

Hypoxia mediated release of endothelial microparticles and increased association of S100A12 with circulating neutrophils

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Microparticles are released from the endothelium under normal homeostatic conditions and have been shown elevated in disease states, most notably those characterised by endothelial dysfunction. The endothelium is sensitive to oxidative stress/status and vascular cell adhesion molecule-1 (VCAM-1) expression is upregulated upon activated endothelium, furthermore the presence of VCAM-1 on microparticles is known. S100A12, a calcium binding protein part of the S100 family, is shown to be present on circulating leukocytes and is thought a sensitive marker to local inflammatory process, which may be driven by oxidative stress. Eight healthy males were subjected to breathing hypoxic air (15% O2, approximately equivalent to 3000 metres altitude) for 80 minutes in a temperature controlled laboratory and venous blood samples were processed immediately for VCAM-1 microparticles (VCAM-1 MP) and S100A12 association with leukocytes by flow cytometry. A pre-hypoxic blood sample was used for comparison. Both VCAM-1 MP and S100A12 association with neutrophils were significantly elevated post hypoxic breathing later declining to levels observed in the pre-test samples. A similar trend was observed in both cases and a correlation may exist between these two markers in response to hypoxia. These data offer evidence using novel markers of endothelial and circulating blood responses to hypoxia.

Introduction

During periods of hypoxia, cells are metabolically compromised resulting in cellular dysfunction that can eventually lead to cell death through metabolic failure. Neutrophils bound to the endothelial surface accelerate endothelial cell dysfunction and tissue destruction through a variety of cytotoxic mechanisms. In addition to physically plugging capillaries,¹ activated neutrophils mediate a variety of harmful responses, such as production of reactive oxygen species (ROS), cytotoxic enzymes and cytokines.² Many of these also recruit additional neutrophils and enhance their adhesiveness to the vascular endothelium. The attachment of neutrophils

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Previously published online as an Oxidative Medicine and Cellular Longevity E-publication: http://www.landesbioscience.com/journals/oximed/article/7611 to the vascular endothelium is mediated by adhesion glycoprotein complexes expressed both on the surface of the neutrophil and on the surface of the endothelial cell. ROS, in large quantities, disrupt the redox balance of cells resulting in oxidative stress and damage to membrane structures (possibly leading to membrane microparticle release), protein damage and cause the activation of signal transduction pathways that lead to apoptosis.³

Microparticles (MP) are cell membrane fragments released into the circulation to a certain extent in healthy individuals under normal homeostatic conditions⁴ and release is increased under stress conditions and disease. MP are released through either cellular activation or apoptosis.⁵ We recently observed a correlation between vascular cell adhesion molecule-1 (VCAM-1) positive MP (VCAM⁺ MP) released from the endothelium and oxidant status,⁶ as measured by thiobarbituric acid reactive substances (TBARS), which is a measure of lipid peroxidation. Previous to this observation, a circadian rhythm in VCAM⁺ MP was observed in healthy males, that along with additional data acquired within this group allowed a normal homoeostatic range to be established.⁷

VCAM-1 is an adhesion molecule, expressed on activated endothelium, being involved in leukocyte attachment and transmigration and is a therapeutic target in inflammatory disease.⁸ Endothelial MP offer an insight into the state of the endothelium as they carry surface antigens particular to the cell of origin and furthermore these MP can be routinely analysed with specific antibodies by flow cytometry.^{5,9} The endothelium is sensitive to oxidative stress¹⁰ and produces nitric oxide (NO) via induction of inducible nitric oxide synthase (iNOS) and constitutively expressed endothelial nitric oxide synthase (eNOS). The endothelium may become activated in times of oxidative stress and this can result in cellular remodelling and the release of MP.¹¹ Diseases, particularly those characterised by endothelial dysfunction, have been shown to result in an elevated concentration of endothelial derived MP.⁵

S100A12, a calcium binding protein belonging to the S100 family, acts as a ligand for the multiligand cell surface receptor for advanced glycation end products $(RAGE)^{12}$ and soluble RAGE (sRAGE). S100 proteins have been implicated in intracellular and extracellular regulatory processes and been shown to mediate inflammatory responses.¹³ S100A12 is secreted by activated neutrophils and inflamed tissue.¹⁴ S100A12 has been shown to interact with RAGE present on the endothelium to trigger an activated state (through NF κ B pathway) resulting in induction of VCAM-1 and

enhanced expression of intercellular adhesion molecule-1 (ICAM-1) in HUVECS that can aid in further recruitment of leukocytes to the injured endothelial cells.¹²

The aim of this study was to quantify the effects of oxidative stress, induced by breathing air equivalent to approximately 3,000 metres (m) altitude (15% oxygen)¹⁵ on circulating VCAM⁺ MP concentration, thus gaining an insight into the endothelial response to hypoxia and furthermore to correlate this response with a novel stress marker, S100A12, associated with the circulating blood cell population of neutrophils and monocytes.

Results

Venous blood samples were taken on numerous days, prior to the hypoxic breathing, to establish a normal range of values in healthy male subjects over a 24 hour (h) period. This data was added to our previously published data that demonstrated a circadian rhythm and resulted in an overall mean of 1350 VCAM⁺ MP per microlitre (μ L) of platelet

poor plasma (PPP), with a mean coefficient of variation of 44% (n = 190). VCAM⁺ MP levels are higher in the morning hours, reaching a minima late afternoon and again in the night. For this reason, the hypoxic (15% Oxygen) treatment was scheduled for 09:00, after which a decrease in VCAM⁺ MP is normally observed. There was a significant effect for time (F = 11.9; p < 0.001) for VCAM⁺ MP (Fig. 1), where post-hypoxic breathing (11:00), there was a large increase from 1020 \pm 263 to 3600 \pm 1040 per µL PPP (p = 0.001), a value well outside the established normal range. Thereafter, VCAM⁺ MP decreased towards the normal range, but at 14:00 was still significantly higher than at 08:00 (p = 0.010). After this time no significant changes were observed in relation to the pre-test value.

There was a significant effect for time (F = 7.3; p < 0.001) for TBARS (Fig. 1), where TBARS demonstrated a small non-significant increase from 08:00 to 14:00, and thereafter, a marked downward trend. TBARS at 20:00 was significantly lower than at 11:00 (p = 0.013) and 14:00 (p = 0.029) and significantly lower at 17:00 than 14:00 (p = 0.047).

S100A12 association with monocytes (S100A12_{mono}) and neutrophils (S100A12_{neutro}) was assessed by flow cytometry. Typical forward scatter/side scatter and fluorescent antibody labelling of neutrophils and monocytes is shown in Figure 2. There was a significant effect for time for S100A12_{neutro} (F = 9.1; p < 0.001) (Fig. 3), where a marked increase was observed immediately after hypoxic breathing (p < 0.001). The S100A12_{neutro} remained elevated at 14:00 (p = 0.044). The effect of time on S100A12_{mono} (Fig. 3) was not statistically significant (F = 1.3; p = 0.29).

Figure 2. Typical flow cytometry profiles showing (top) forward scatter/ side scatter with Netrophils and monocytes gated accordingly and (bottom) fluorescence profiles showing gated (black) neutrophils and (grey) monocytes incubated with either (filled) goat serum or (line) anti-human EN-RAGE.



Figure 1. Time dependent profile for VCAM⁺ MP and TBARS before and after hypoxic breathing. The data are represented as the mean and standard error of the mean (n = 8). HB = time during which subjects breathed hypoxic air.





Figure 3. Time dependent profile for S100A12 expression on monocytes and neutrophils before and after hypoxic breathing. The data are represented as the mean and standard error of the mean (n = 8). HB = time during which subjects breathed hypoxic air.

Significant relationships were observed between VCAM⁺ MP and S100A12_{neutro} ($r_s = 0.39$; p = 0.044) and between S100A12_{mono} and S100A12_{neutro} ($r_s = 0.74$; p < 0.001).

Discussion

Endothelial MP have been proposed as a sensitive blood borne marker of endothelial state.⁴ The endothelium is known to be affected by oxidative stress and the subjects in this study were exposed to a significant oxidative stress by hypoxic breathing and subsequent re-oxygenation. This resulted in an increase in plasma TBARS, a measure of lipid peroxidation, and a marker of oxidative stress. Hypoxia leads to numerous cellular responses including generation of ROS and production of hypoxia-inducible factor-1.16 Hypoxic breathing led to a significant increase in circulating VCAM+ MP (Fig. 1), demonstrating a novel measure of endothelial response to hypoxia. Whether or not these MP came from localised activated endothelium or hypoxia mediated endothelial activation is debatable. Previously we have reported a circadian variation in VCAM+ MP,⁷ with an early morning peak decreasing into the afternoon. Conversely, the data presented here shows a large increase from early morning, after hypoxic breathing to levels well outside the normal range.

The synthesis of S100A12 by neutrophil granulocytes has been shown.¹⁷ The role of S100A12 has been implicated in the innate immune system in response to infection. The exact nature of S100A12 ligand interaction is unclear, it has been shown able to bind both itself and a range of other molecules and was suggested to act as a molecular chaperone in the presence of calcium.¹⁸ RAGE been identified as a receptor for S100A12.¹² This RAGE interaction leads to initiation of pro-inflammatory signalling cascades via MAP-Kinase and NF κ B, resulting in increased adhesion molecule

expression.¹⁹ Increased serum concentrations of S100A12 have been shown in patients with severe bacterial infections.²⁰ Furthermore, serum S100A12 maybe of use as a diagnostic marker in inflammatory bowel disease,²¹ arthritis²² and asthma.²³ In inflammatory disease high concentrations of S100A12 have also been shown at local sites of inflammation such as synovial fluid,²⁴ inflamed bowel mucosa²⁵ and atherosclerotic lesions.²⁶

S100A12 interaction with RAGE is capable of activation of endothelial, lymphocyte and macrophage cells directly.13 S100A12 has been postulated to be a novel marker of localised inflammatory processes in part due to its overexpression at local inflammatory sites.¹⁹ As S100A12 is a ligand for RAGE it may be postulated that there is either an increase in RAGE expression upon the neutrophil surface following hypoxia or a higher degree of circulating S100A12 is bound to the receptor. RAGE expression has previously been shown to be affected by hypoxia via hypoxiainducible factor-1.27 Interaction between RAGE and its ligands has also been demonstrated to upregulate VCAM-1 expression on endothelium and promote other proinflammatory pathways.²⁸ Furthermore endothelial cells release RAGE specific ligands in response to hypoxia,29 which could account for the observed increase

of neutrophil associated S100A12.

The amount of \$100A12 associated with both neutrophils and monocytes appear correlated in this study, suggesting a similar response to hypoxia, although the increase in the S100A12_{mono} subject mean was not found significant. The increase in S100A12_{neutro} following hypoxic breathing suggests S100A12 may also be a marker of oxidative stress in the circulating blood cell population. A significant correlation was observed between both endothelium and neutrophil reaction to hypoxia via release of VCAM⁺ MP and $S100A12_{neutro}$, (p = 0.044). This may be expected as both cell types have intrinsic roles in hypoxia (especially I/R injury) and localised inflammation, where VCAM-1 mediates firm attachment of neutrophils to the endothelium. A recent study found significant correlations between serum concentrations of advanced glycation products (AGE) and VCAM-1 and the soluble form of S100A12 in diabetic patients.³⁰ It has already been suggested that serum levels of cell surface markers may in fact be MP associated and not entirely freely soluble,⁵ furthermore elevated endothelial MP concentration is established in diabetes.³¹ Therefore these data³⁰ may well include VCAM+ MP released from dysfunctional endothelium in diabetic patients and their correlation with S100A12, as observed in the present study. In addition advanced oxidation protein products have recently been shown to induce overexpression of VCAM-1 on endothelial cells via S100A12 mediated signalling.^{32,33} Induction of ROS generation and VCAM-1 expression was effectively prevented by blocking S100A12 binding to the endothelial surface.³²

The effects of hypoxia on the vasculature are well characterised and here we highlight two potentially novel markers which link the blood response and endothelial response within healthy human subjects. Further research is necessary to elucidate the specific cellular mechanism and consequences behind these observations.

Table 1	Sequence of experimental events
Time (h)	Experimental event
07:15	Subjects report to laboratory
08:00	Pre-hypoxia blood sample
09:00	Hypoxic breathing initiated
10:20	Hypoxic breathing ended
11:00	1 st post-exposure blood sample
14:00	2 nd post-exposure blood sample
17:00	3 rd post-exposure blood sample
20:00	4 th post-exposure blood sample

Materials and Methods

Eight healthy male volunteers (18–24 years old) participated in this study. All subjects gave written informed consent and ethical approval was granted by the Departmental ethics committee at the University of Hull. Subjects reported to a temperature-controlled laboratory at 07:15 on the day of the study and a pre-test blood sample was drawn at 08:00. Venous blood was obtained by venipuncture into potassium EDTA tubes (Vacuette[®], Greiner, UK) for analysis of both S100A12 and TBARS or sodium citrate tubes (Vacuette[®], Greiner, UK) for MP analysis pre- and post-exposure according to Table 1. Exposure consisted of breathing hypoxic air (15% Oxygen, air mix) produced by a hypoxicator for 80 minutes (min), through masks. The hypoxic exposure started at 09:00 and lasted 80 min; 15% Oxygen is equivalent to approximately 3000 m altitude.¹⁵

MP quantification. Citrated blood was analysed according the method of Hugel et al.³⁴ Platelet rich plasma was obtained by centrifugation (180 xg, 10 min) and this was further centrifuged (13,000 xg, 2 min) to remove platelets. The now platelet poor plasma 25 μ L was then incubated with either Isotype matched IgG negative control FITC conjugate (4 μ L, AbD Serotec, UK) or IgG VCAM-1 specific FITC conjugate (4 μ L, AbD Serotec, UK) for 30 min. Quantification was achieved by adding counting beads (25 μ L, Caltag medsystems, UK) and 0.2 μ m filtered PBS (200 μ L) prior to flow cytometry.

S100A12 analysis. EDTA blood (100 μ L) was mixed with Erythrolyse (2 mL, AbD Serotec, UK) to selectively lyse red blood cells. The white cells were then pelleted by centrifugation (400 xg, 5 min), washed and resuspened in PBS. Cells (2 x 10⁵) were incubated with either goat serum, to control for non-specific binding, or goat anti-human EN-RAGE antibody (RnD Systems, UK). Cells were incubated with antibody for one hour before washing with PBS, centrifugation (400 xg, 5 min) and incubating (30 min) with a secondary, FITC conjugated anti-goat IgG antibody (AbD Serotec, UK). Cells were washed as above and resuspended in a final volume of 300 μ L PBS prior to flow cytometry analysis.

Flow cytometry. For data acquisition a BD FACSCalibur (BD Biosciences, UK) was used and analysed using CellQuest software. For S100A12, 10,000 cells were counted and 50,000 events were counted for MP analysis. For S100A12 analysis leukocytes were gated according to forward scatter/side scatter characteristics and data is presented as mean fluorescence intensity (MFI) on the FL-1 scale of the cells incubated with anti-human EN-RAGE minus the MFI obtained with goat serum.

TBARS analysis. EDTA plasma was obtained via centrifugation of the EDTA Vacuette[®] (10 min, 1500 xg) and stored at -80°Centigrade until analysed. Plasma TBARS was then analysed using a commercially available kit (Zeptometrix) following manufacturers' instructions and are expressed as malondialdeyde equivalents.

Statistical analysis. All statistical analyses were completed using SPSS for Windows version 16.0 (SPSS Inc., Chicago, IL). Normality for each dependent variable for each time point was checked using Q-Q plots and deemed plausible in each instance. Sample descriptive statistics are reported as means and standard deviations. The effect of time on TBARS expression and S100A12 expression on both monocytes (S100A12_{mono}) and neutrophils (S100A12_{neutro}) was analyzed using one-way analysis of variance for repeated measures. The Huynh-Feldt corrected p value was reported where the sphericity assumption was violated (as identified by a significant Mauchly's test statistic). The VCAM⁺ MP data showed considerably less variance at 08:00 in comparison with later time points. The VCAM⁺ MP data was therefore analyzed using a linear mixed model incorporating an autoregressive repeated covariance structure. Sidak post hoc pairwise comparisons were used in the event of significant main effects. The relationship between VCAM+ MP and $S100A12_{neutro}$ and between $S100A12_{mono}$ and $S100A12_{neutro}$ were analyzed using the Spearman correlation. Two-tailed statistical significance was as accepted as p < 0.05.

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6



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