Differential Effect of LPS on Glucose, Lactate and Inflammatory Markers in the Lungs of Normal and Diabetic Mice

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Abstract

Elevation of blood glucose results in increased glucose in the fluid that lines the surface of the airways and this is associated with an increased susceptibility to infection with respiratory pathogens. Infection induces an inflammatory response in the lung, but how this is altered by hyperglycemia and how this affects glucose, lactate and cytokine concentrations in the airway surface liquid is not understood. We used Wild Type (WT) and glucokinase heterozygote (GK+/-) mice to investigate the effect of hyperglycemia, with and without LPS-induced inflammatory responses, on airway glucose, lactate, inflammatory cells and cytokines measured in Bronchoalveolar Lavage Fluid (BALF).

We found that glucose and lactate concentrations in BALF were elevated in GK+/- compared to WT mice and that there was a direct correlation between blood glucose and BALF glucose concentrations. LPS challenge increased BALF inflammatory cell numbers and this correlated with decreased glucose and increased lactate concentrations although the effect was less in GK+/- compared to WT mice. All cytokines measured (except IL-2) increased in BALF with LPS challenge. However, concentrations of TNFα, INFγ, IL-1β and IL-2 were less in GK+/- compared to WT mice. This study shows that the normal glucose/lactate environment of the airway surface liquid is altered by hyperglycemia and the inflammatory response. These data indicate that inflammatory cells utilize BALF glucose and that production of lactate and cytokines is compromised in hyperglycemic GK+/- mice.

Keywords: Airway; Glucose; Hyperglycemia; Inflammation; Metabolism

Abbreviations

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<tr>
<td>WT</td>
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<td>GK</td>
<td>Glucokinase</td>
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Introduction

There is mounting evidence that raised blood glucose concentrations (hyperglycemia) as a result of diabetes mellitus or severe illness (stress hyperglycemia) are associated with increased susceptibility to respiratory infections, particularly when associated with underlying respiratory disease[1-4]. Glucose concentration in the fluid that lines the luminal surface of the airways (airway surface liquid, ASL) is in normal situations low at ~0.4mM. However,
we have shown that elevation of blood glucose leads to increased glucose concentration in ASL (up to ~4.0mM) in nasal and bronchial secretions in patients[13,5,6]. Glucose was also elevated in Bronchoalveolar Lavage Fluid (BALF) from the lungs of a mouse model of diabetes (the leptin receptor deficient db/db mouse) compared to wild type[7]. Elevation of glucose was associated with increased Staphylococcus aureus and Pseudomonas aeruginosa in the airway and we have shown that sugars in the ASL can be utilized by these bacteria as a nutrient source for growth[8-10].

Inflammatory stimuli increase the permeability of the epithelium to glucose, elevating glucose diffusion from blood into the ASL[9]. Inflammatory stimuli also increase activation and genesis of inflammatory cells and their recruitment to the airway lumen[7,11]. Thus, glucose in the ASL potentially supplies a local nutrient source for inflammatory cells of the respiratory tract. Hyperglycemia has also been shown to modify inflammatory responses in the lung[12,13]. We hypothesized that the availability of glucose in the ASL and its metabolism by inflammatory cells could contribute to this effect. In addition, if inflammatory cells utilize glucose this would reduce glucose concentration in ASL and affect the concentration of other metabolites such as lactate.

Here, we investigated the relationship between blood glucose and glucose concentration in the BALFs of normal (WT) and hyperglycemic (glucokinase deficient, GK−/−) mice and studied how this relationship, the concentration of lactate, production of cytokines and the inflammatory responses changed when the lungs were challenged with bacterial lipopolysaccharide.

Materials and Methods

Male and female (17 weeks old, average weight 24±3g) heterozygote GK−/− mice on C57/B6 background and wild type C57/B6 mice were used in the study[14]. Animals were kept in a facility with 12h light/dark cycle at 21±2°C and with 55±15% relative humidity. Water and normal rodent chow were available ad libitum.

WT and GK−/− mice were randomized into treatment groups of 4, 24 and 48 hours’ post LPS challenge versus no challenge (indicated as time 0) using 4-5 animals per group. Experiments were repeated for 0 and 48 hours to obtain additional samples for cytokine analysis and results from these studies were pooled. LPS from P.aeruginosa (Sigma-Aldrich, USA) was previously frozen in 1mg/ml aliquots and diluted to the concentration needed to give 0.0875μg/g mouse in 50μl (based on the average weight of the group) and given by intranasal dosing at time 0. In brief, animals were anaesthetized with isoflurane 4-5% (O2 1.2l/min) prior to administration of the LPS solution to one nostril, which was subsequently inhaled naturally. Mice were then returned to their cages when retaining consciousness.

Body weight was recorded once daily from 0-48 hours after administration of LPS to follow the wellbeing of the animals. Blood was collected from the vena saphena in conscious mice for glucose evaluation after 4 hours of fasting. At 4, 24 or 48 hours after LPS administration, the animals were terminated with an overdose of pentobarbital (0.2ml of 100mg/ml i.p.). The lungs of each animal were subjected to Bronchoalveolar Lavage (BAL). In brief, the trachea was exposed and a catheter inserted into the trachea through a cut between two upper cartilage rings below the larynx. The catheter was secured with a silk suture and three volumes of 0.3ml saline were instilled, gently aspirated, pooled and weighed.

The BAL fluid (BALF) was centrifuged (SorvallRotanta 46R, HettichZentrifugen, Tullingen, Germany, 1200rpm (314g), 10min, 4°C). The pellet was re-suspended in 0.25ml of PBS and the total and differential cell count was performed using SYSMEX XT-1800i Vet. (SYSMEX, Kobe Japan). Cell count is expressed as the amount of cells per ml of recovered BALF. The supernatant was used to measure glucose and lactate on the ABX Pentra 400 (Horiba ABX Medical, Kyoto, Japan) according to the manufacturer’s protocol. Inflammatory cytokines in the supernatant were analyzed by means of an MSD 10-plex (K15048D-1 MSD mouse Pro inflammatory panel (V-plex), Mesoscale Discovery, Maryland, US). At termination, blood from behind the eye was collected in EDTA tubes and blood glucose was measured directly using Accu-check (Roche, Bromma, Sweden). Plasma lactate was assayed using theABX Pentra 400. All experiments were approved by the local Ethical committee in Gothenburg (184-2012).

Statistical Analysis

Values are reported as the mean±SEM. Statistical analysis was performed using analysis of variance (ANOVA) tests followed by Bonferroni multiple comparison post hoc tests (Graph-Pad Prism) or Student’s t test. P values of <0.05 were considered statistically significant.

Results

Characterization of Airway Glucose Metabolism

Fasting blood glucose was higher in GK−/− than Wild Type (WT) mice (15.2±4.9 compared to 9.4±0.19mM, p<0.0001, n=25) (Figure 1a). Blood glucose remained higher in GK−/− mice compared to WT 48 hours after treatment with LPS although the concentration of glucose was reduced to 11.7±0.6 and 7.7±0.2mM, respectively (p<0.0001, n=17). There was no significant difference in plasma insulin levels between WT and GK−/− mice (0.59±0.12 and 0.47±0.1ng/ml, respectively, n=9). There were also no significant changes in blood lactate concentration in WT or GK−/− at any of the time points tested (Figure 1b).

Glucose concentration in the BALF of GK−/− mice was significantly higher than that of WT mice (0.15±0.2 compared to 0.03±0.004mM, p<0.0001, n=19). It remained higher in the GK−/− mice following LPS challenge, but at significantly lower levels for both mouse types (p<0.001 at 24 and 48 hours, n=4 and 14, respec-
We could not detect any measurable glucose in BALF from WT mice at 24 hours (Figure 1c). There was a significant relationship between blood glucose/BALF glucose concentration prior to treatment with LPS and independent of mouse type (p<0.0001, n=44). However, this relationship was lost 48 hours after treatment with LPS (p=0.05, F=6, Df=73) (Figure 2a).

Lactate concentration in the BALF of GK+/− mice was significantly higher than WT in the absence of an inflammatory stimulus (0.1±0.008 compared to 0.07±0.007mM, p<0.05, n=19) (Figure 1d). As expected, the concentration of lactate in the BALFs of GK−/− and WT mice increased after treatment with LPS. Less so in the GK−/− mice, only reaching significance after 48 hours (p<0.01, n=16), whereas lactate concentration was significantly increased at 24 and 48 hours in WT mice (p=0.0001, n=3 and n=16, respectively). There were also higher concentrations of lactate in WT compared to GK+/- mice at 48 hours (p<0.01, n=15) whereas the difference at 24 hours was borderline significant (p=0.059, n=4). Thus, less lactate was produced in the lungs of GK−/− mice after an inflammatory stimulus than in WT mice (Figure 1d).

Impact of Hyperglycemia on Airway Response to Lipopolysaccharide

Prior to LPS treatment, GK+/- mice had similar numbers of inflammatory cells (total White Blood Cells (WBC) including lymphocytes, macrophages, neutrophils and eosinophils) in the BALF as compared to WT. Treatment of GK+/- with LPS elicited an increase in inflammatory cells recovered from the BAL which was maximal at 24-48 hours. The absolute numbers of lymphocytes and neutrophils were similarly increased in WT and GK+/- at 4, 24 and 48 hours versus untreated controls. Macrophages in the BALF were not elevated until 24 hours’ post LPS treatment and to a similar extent in WT and GK−/− mice (Figure 3a-c). Interestingly, 24 hours’ post LPS treatment, eosinophils were significantly reduced in GK−/− compared to WT mice (p<0.01, n=3) indicating that the elevation of blood glucose in these mice could influence eosinophil recruitment and/or survival (Figure 3d).

There was no relationship between blood lactate and BALF lactate but there was a small positive relationship between blood glucose and BALF lactate (p<0.01, n=41) at baseline (Figure 2b). This relationship was completely reversed 48 hours after treatment with LPS where lactate in the BALF decreased with increasing blood glucose (p<0.01, n=29) (Figure 2b), similarly for both mouse types.
most numerous cells in BALF, indicating that these were the main source of lactate. There was an indication that the slope of the line was reduced and that less lactate was produced by the inflammatory cells in GK+/- compared to WT mice, however, we were unable to attribute significance to this observation (p=0.08). Interestingly, eosinophils in WT mice showed a positive association with lactate (p<0.0001, n=39) but this was not observed in GK+/- mice (Figure 1d and Figure 3d).

Figure 3: Numbers of lymphocytes (a), neutrophils (b), macrophages (c) and eosinophils (d) in BALF from wild type (WT) (white symbols) and hyperglycaemic GK+/- mice (black symbols) before (circles) and after treatment with LPS for 4 (triangles), 24 (inverted triangles) and 48 hours (squares). Data are shown as individual measurements. Horizontal bars show mean±SEM. Significantly different from untreated control, *p<0.05. Significantly different from WT at similar time point, †p<0.05.

Hyperglycemic Airways and Secretion of Cytokines in Response to Respiratory Challenge

As the greatest differences in the parameters tested were identified at 48 hours’ post LPS challenge, we then investigated cytokine concentration in the BALFs of challenged and unchallenged animals. There was almost no production of cytokines into the BALF of untreated WT and GK+/- mice, consistent with the lack of inflammatory cells. LPS challenge for 48 hours significantly increased the concentration of TNFα, IL-1β, IL-6, IL-8 and INFγ(Figure 4a-f). IL-2 concentrations were lower than other cytokines and showed no elevation with LPS treatment (Figure 4d). There was significantly less IL-2, IL-1β and TNFα in the BALFs of GK+/- compared to WT mice 48 hours after LPS treatment (p<0.01, n=8-10 respectively).

Figure 4: Concentration of cytokines, TNFα(a), INFγ(b), IL-1β(c), IL-2 (d), IL-6 (e) and IL-8 (f) present in BALF from wild type (WT) (white symbols) and hyperglycaemic GK+/- mice (black symbols) before (circles) and after treatment with LPS for 48 hours (squares). Data are shown as individual measurements. Horizontal bars show mean±SEM. Significantly different from untreated control, *p<0.05. Significantly different from WT at similar time point, †p<0.05.

Discussion

The present study demonstrates increased respiratory glucose concentration in a further hyperglycemic mouse model of human diabetes, in line with previous reports including human disease[1,3,7,10,15]. Also, here, we show a positive and significant correlation of airway glucose levels with increasing blood glucose concentrations, but no impact on inflammatory cells or cytokine release in the unchallenged situation. Exposure to Lipopolysaccharide (LPS) completely changes the blood/BALF glucose relationship. Glucose concentration falls and lactate rises as inflammatory cell numbers increase. We propose that inflammatory cells utilize the available glucose and increase the production of lactate. This effect is altered with hyperglycemia. Lactate concentration is lower and goes in hand with less cytokine production and recruitment of eosinophils. These data indicate a defective/suppressed response in hyperglycemic lungs to respiratory infections.

We have used a mouse model of hyperglycemia (the glucokinase heterozygous knockout model, GK+/-) which exhibit impaired
glucose tolerance and diminished insulin secretion in response to glucose resulting in raised blood glucose concentration[16,17]. These mice do not exhibit obesity or hyperinsulinemia that is associated with the leptin and leptin receptor deficient models of type II diabetes such as the ob/ob and db/db mouse[18,19]. In this mouse, and similar to our findings in vivo and in vitro, raised blood glucose is associated with increased glucose in Bronchoalveolar Lavage Fluid / Airway Surface Liquid (BALF / ASL) and there is a positive relationship between blood glucose concentration and glucose in BALF. It is difficult to calculate the volume of mouse lung ASL and thus the absolute concentration of glucose without using dilution markers. However, a mouse of the weight used in these experiments would likely have a surface area of 70cm²[20,21]. Airway surface liquid has been proposed to be a depth of 8-35µm depending on the radius of the airway [21,22] and alveolar fluid depths are much smaller at 0.1µm [23]. A contribution of 50:50 alveolar: airway surface liquid to BALF glucose would give an estimate of 0.48mM glucose in ASL from WT mice and 1.5mM in GK⁻⁺ mice. Even if it is assumed that most of the sampling is from the distal lung i.e. 90:10 alveolar: airway, the resultant glucose concentrations are still calculated to be significantly less than corresponding blood glucose in accordance with our previous in vivo and in vitro studies [3,9].

Treatment of mice with LPS induced a robust inflammatory response as indicated by the elevation in inflammatory cells and pro-inflammatory cytokines in the BALF. Inflammatory cell numbers after LPS challenge were similar in GK⁻⁺ compared to WT mice except for eosinophils which were reduced. Similarly, eosinophils but not neutrophils, were reduced in ovalbumin challenged lungs of diabetic ob/ob mice[24]. No difference in lung neutrophils was shown in diabetic Zucker rat lungs after tracheal instillation of LPS or in diabetic Wistar rats compared to wildtype [12,13]. Thus, hyperglycemia appears to effect eosinophil recruitment to the lung.

LPS challenge decreased BALF glucose over 48 hours. Although, blood glucose declined with the inflammatory response [25] this was unlikely to be the cause of the change in BALF glucose as the relationship between blood and BALF glucose was lost and BALF glucose concentrations were much lower than expected for the corresponding blood glucose after LPS treatment. There was however, a clear inverse relationship between glucose concentration and number of inflammatory cells/lactate in BALF. A previous study in rats showed that tracheal instillation of LPS increased glucose utilization by White Blood Cells (WBC) in the airways [26]. This was mostly explained by increased numbers of neutrophils which had high glucose utilization and a doubling of glucose utilization by activated macrophages [26]. In our study, neutrophils and macrophages were the predominant cells post LPS treatment. In rats, glucose utilization by activated macrophages and PMNs increased to ~7.5 and 1.75nmols/min/10⁶ cells, respectively.

We had approximately 2 x 10⁶ macrophages plus neutrophils in the mouse BALF 48 hours after LPS treatment. If rat and mouse models are similar, then glucose utilization would be ~18nmole/min. WT BALF contained 35nmoles of glucose which would be depleted rapidly by LPS activated inflammatory cells. Thus, utilization of glucose by inflammatory cells would account for the reduced glucose concentration in the BALF after LPS challenge. This hypothesis now needs to be tested to establish if this is indeed the case.

Glucose remained detectable in the BALF (except in WT 24 hours post LPS treatment) and glucose concentrations were persistently higher in GK⁻⁺ mice. We previously showed that LPS increased glucose permeability across ex vivo mouse trachea and in vitro which would increase glucose diffusion into ASL [9]. We therefore suggest that glucose in ASL is metabolized by inflammatory cells and that increased glucose permeability during the inflammatory response increases glucose diffusion into ASL and this process may help sustain inflammatory cell function. During hyperglycemia, the increased gradient for glucose diffusion would drive even more glucose into ASL resulting in the higher residual BALF glucose we observed after LPS treatment in GK⁻⁺ mice.

The increase in inflammatory cells correlated with elevation of pro-inflammatory cytokines and lactate. Lactate is involved in inflammatory cell signaling and is released at sites of inflammation by activated inflammatory cells. Its production is dependent on cellular glucose metabolism by inflammatory cells such as neutrophils[27-29]. However, we also observed a small increase in BALF lactate (but not in cytokines or inflammatory cells) in untreated GK⁻⁺ compared to WT mice. We have evidence that hyperglycemia elevates epithelial lactate secretion. This may therefore reflect a contribution from airway epithelial cells to lactate in BALF[8,30-32].

Here we provide evidence that LPS elevated lactate concentration in the BALF of GK⁻⁺ mice was reduced compared to WT. Consistent with our results, peritoneal neutrophils from streptozotocin-treated diabetic rats exhibited reduced lactate production[33]. In these neutrophils, the effect was reversed by insulin treatment. Insulin has also been shown to alter other LPS-induced inflammatory responses in diabetic rats[34], yet the GK⁻⁺ mice used in this, and two other studies, had normal circulating plasma insulin levels[14,16]. Thus, the role of luminal glucose concentration and insulin on inflammatory cell responses in these mice and in the lung, needs further investigation.

LPS-treated GK⁻⁺ mice also exhibited decreased TNFα, IL-1β and IL-2 in the BALF compared to WT. Lower levels of TNFα and IL-1β were reported 4-6 hours after LPS challenge also in diabetic Wistar rats compared to WT[13,35]. Similar to our findings, in one study this was independent of changes in neutrophils although in the other, this was not the case[13,35]. TNFα and IL-1β are se-
creted by activated macrophages. Whilst, we did not see a significant reduction of macrophages in BALF, there is evidence that macrophage secretory function was impaired in LPS-challenged streptozotocin-induced diabetic mice[36].

IL-2 is secreted by αβT-cells which are involved in endotoxin-induced airway inflammation[37]. We cannot explain why IL-2 was reduced in the GK<sup>−/−</sup> mice after LPS treatment. Genes involved in IL-2 signaling are implicated in the development of type I diabetes and circulating levels are reduced in this disease[38]. We did not measure plasma IL-2 and so cannot confirm whether this was the case in GK<sup>−/−</sup> mice. IL-2 is further involved in modifying cytokine secretion from other inflammatory cells. For example, human airway macrophages can be induced to produce TNFα in response to IL-2[39]. In addition, eosinophil degranulation is thought to be triggered by IL-2[38]. Thus, it is possible that the reduction of IL-2 in LPS-treated GK<sup>−/−</sup> mice leads to the reduction in the secretion of other cytokines and reduced eosinophil numbers at 24 hours.

**Conclusion**

Taken together, we show that acute LPS challenge in the mouse lung results in a reduction of BALF glucose concentration and elevation of inflammatory cells but that there is altered inflammatory cell responses in the lungs of GK<sup>−/−</sup> mice. We propose that glucose in the lung lumen is utilized by inflammatory cells during the inflammatory response. We showed previously that hyperglycemia results in elevated ASL glucose and that this promotes the growth of respiratory pathogens in the lung. We also proposed a model for glucose homeostasis across the airway [7,9,30,40]. We now propose a more complex model whereby, during infection, the inflammatory response competes with the airway epithelium and bacteria for glucose in the in the lung. Hyperglycemia elevates glucose concentration in the lung lumen resulting in a bigger nutrient pool and an altered immune response. Recent findings in the upper airway also showed that glucose in the airway surface liquid may suppress anti-microbial peptide secretion[41,42]. Thus, the balance that favors neutralization of pathogens in normal lungs may shift to favor pathogen evasion/survival in lungs exposed to hyperglycemia. Such changes may underlie the increased susceptibility to infection in patients with respiratory disease and diabetes (COPD/CF).

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**Authorship**

Deborah L Baines, Emma H Baker, John D Taylor had the original concept for this work. Annika Åstrand, Cecilia Nagorny Holmberg, James P Garnett, Cecilia Wingren and Gaëll Mayer designed the study, acquired and analyzed the data. Deborah L Baines, Annika Åstrand, Cecilia Nagorny Holmberg, Gaëll Mayer and John D Taylor interpreted the data and compiled the work for publication.

**References**