• CLINICAL RESEARCH •

Influence of acute hyperglycemia in human sepsis on inflammatory cytokine and counterregulatory hormone concentrations

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Abstract

AIM: In human sepsis, a prominent component of the hypermetabolite is impaired glucose tolerance (IGT) and hyperglycemia. Elevations in plasma glucose concentration impair immune function by altering cytokine production from macrophages. We assessed the role of glucose in the regulation of circulating levels of insulin, glucagon, cortisol, IL-6 and TNF- α in human sepsis with normal or impaired glucose tolerance.

METHODS: According to the results of intravenous glucose tolerance test, forty patients were classified into two groups: control group (n=20) and IGT group (n=20). Plasma glucose levels were acutely raised in two groups and maintained at 15 mmol/L for 3 hours. Plasma insulin, glucagon and cortisol levels were measured by radioimmunoassay, the levels of TNF- α and IL-6 were detected by ELISA.

RESULTS: In IGT group, the fasting concentrations of plasma glucose, insulin, glucagon, cortisol, IL-6 and TNF- α levels were significantly higher than those in control group (P < 0.05). During clamp, the control group had a higher average amount of dextrose infusion than the IGT group (P < 0.01). In control group, plasma insulin levels rose from a basal value to a peak at an hour (P < 0.05) and maintained at high levels. Plasma glucagon levels descended from a basal value to the lowest level within an hour (P < 0.01) and low levels were maintained throughout the clamp. In IGT group, plasma insulin was more significantly elevated (P < 0.01), and plasma glucagon levels were not significantly declined. Plasma cortisol levels were not significantly changed in two groups. In control group, plasma IL-6 and TNF- α levels rose (P<0.01) within 2 hours of the clamp and returned to basal values at 3 hours. In IGT group, increased levels of plasma cytokine lasted longer than in control group (3 hours vs. 2 hours, P<0.05), and the cytokine peaks of IGT group were higher (P < 0.05) than those of control group.

CONCLUSION: Acute hyperglycemia pricks up hyperinsulinemia and increases circulating cytokine concentrations and these effects are more pronounced in sepsis with IGT. This suggests a potential modulation of immunoinflammatory responses in human sepsis by hyperglycemia. Yu WK, Li WQ, Li N, Li JS. Influence of acute hyperglycemia in human sepsis on inflammatory cytokine and counterregulatory hormone concentrations. *World J Gastroenterol* 2003; 9(8): 1824-1827

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INTRODUCTION

Severe human sepsis is associated with hypermetabolic stress response, and affects protein, carbohydrate and lipid metabolism throughout the body^[1-7]. A prominent component of hypermetabolic stress response is hyperglycemia and impaired glucose tolerance (IGT). The mechanisms of stress hyperglycemia are well known^[8]. Counterreglulatory hormone and excess cytokine result in insulin resistance, and many hospitalized patients are insulin deficient for a variety of reasons (e.g. old age, pancreatitis, hypothermia, hypoxemia). Excess dextrose infusion is an often-overlooked contributor to hyperglycemia^[9].

The harm of acute hyperglycemia in stress response patients has been demonstrated in several clinical and experimental conditions^[10-12]. An increased susceptibility to infections in the presence of hyperglycemia has long been known in patients with diabetes^[10,11,13]. And recent investigations have demonstrated that elevations in plasma glucose concentration impair immune function by altering cytokine production from macrophages, diminishing lymphocyte proliferation, and depressing intracellular bactericidal activity of leukocytes^[14-17]. Furthermore, in the issue of circulation, Esposito K. argued hyperglycemia acutely increased circulating cytokine concentrations, and this effect was more pronounced in group with IGT^[18]. But former researches focused on normal human being, diabetes or critically ill patients, etc. and rarely concentrated on severe human sepsis.

The present study was to test whether circulating levels of hormones and cytokines are regulated by glucose levels in sepsis, and to measure serum insulin, glucagon, cortisol, TNF- α , and IL-6 concentrations during acute hyperglycemia in septic patients with normal or impaired glucose tolerance (IGT).

MATERIALS AND METHODS

Subjects

The subjects of this study were the patients admitted from January 5 to November 1, 2002 to the medical SICU of Jinling Hospital in Nanjing. Forty patients (23 men, 17 women) with a mean (s.d.) age of 52.2 (15.6) years with abdominal or pelvic sepsis were studied. Patients with diabetes mellitus, trauma, human immunodeficiency virus disease, end-stage renal disease, end-stage hepatic disease, the patients receiving immunosuppressive agents were excluded. Forty patients were divided into two groups according to normal (control group, n=20) or impaired (IGT group, n=20) glucose tolerance (Table 1). The patients' glucose tolerance was assessed by intravenous glucose tolerance test (IVGTT). After an overnight fast each patient underwent IVGTT: a 50 % glucose solution (glucose 0.5 gm/kg of body weight) was injected into the femoral vein

for 2 minutes. Patients in IGT group had a 2-hour plasma glucose value between 7.7 mmol/L and 11 mmol/L, and the control group had normal glucose tolerance (2-hour plasma glucose value below 7.7 mmol/L). Informed consent was obtained from all participating patients or their surrogates.

Severity of sepsis and underlying diagnosis

Sepsis was defined by the American College of Chest Physicians-Society of Critical Care Medicine consensus statement by an identifiable site of infection and evidence of a systemic inflammatory response manifested by at least three of the following criteria: (1) temperature, >38 $^{\circ}$ C or <36 $^{\circ}$ C; (2) heart rate, >90 beats per minute; (3) respiratory rate, >20 breaths per minute; (4) white blood cell count, >12 000/mm³ or <4 000/mm^{3[19]}. Patients meeting enrollment criteria were entered within 24 h. The cause of sepsis was severe acute pancreatitis (n=11), colorectal anastomotic dehiscence (n=9), perforated diverticular disease (n=9), gastroduodenal perforation (n=7), gallbladder perforation (n=3) and spontaneous splenic abscess (n=1). Sepsis severity was scored using the method of Elebute and Stoner^[20]. This scoring procedure takes into account the site of infection, bacteriology, body temperature, secondary effects (e.g. jaundice) and various haematological and biochemical variables, such as white cell count and plasma albumin concentration.

Study protocol

After 12-hour fast overnight, the patients were placed in a supine comfortable position with the sickroom temperature between 20 °C and 24 °C. Intravenous lines were inserted into a large antecubital vein of one arm for infusions and into a dorsal vein of the contralateral arm for blood sampling. Patency was preserved in a slow saline infusion (0.9 % NaCl). After withdrawal of baseline blood samples, plasma glucose concentrations were acutely raised with a bolus injection of 0.25 g/kg glucose followed by a varying 30 % glucose infusion to achieve steady-state plasma glucose concentrations of about 15 mmol/L for 180 minutes.

Analysis

Samples for analysis of plasma glucose were collected in tubes containing a trace of sodium fluoride. Plasma glucose was determined according to the glucose oxidase method with an autoanalyzer (Beckman Instruments). Serum samples for measuring hormone and cytokine level were stored at -80 °C until assay. Commercially available kits were used for radioimmunoassay of plasma insulin, glucagon and cortisol concentrations. Serum concentrations of TNF- α , IL-6 were determined in duplicate with commercially available kits (R&D Systems). Dilution curves of serum samples were parallel to those of standard. Intra-assay and interassay coefficients of variation were 3.8 % and 5.8 % for TNF- α , 2.8 % and 3.3 % for IL-6.

Statistical analysis

Results were given as mean \pm SD. One-way ANOVA was used to compare baseline data, followed by Scheffé's test for pairwise comparisons. Multiple comparison tests were made with ANOVA, followed by post hoc analysis (Student-Newman-Keuls test) to locate the significant difference indicated by ANOVA. A value of *P*<0.05 was considered statistically significant. Data were analysed using Statistical Package for the Social Sciences computer software (SPSS 11.0).

RESULTS

During the 7-month study period 46 patients were collected,

however analysis was limited to 40 patients with complete and available data. IGT group had higher admission severity scores than control group (APACHE \parallel , 10 (7-15) *vs.* 7 (4-10), sepsis score, 13 (8-19) *vs.* 8 (6-13); *P*<0.05). In IGT group, the levels of fasting plasma glucose, insulin, glucagon, cortisol, IL-6 and TNF- α levels were significantly higher than those in control group (Table 1). During the clamp, plasma glucose became stabilized at 15 mmol/L with oscillations not exceeding 5 % of the prefixed value. The control group had a higher average amount of dextrose infusion than IGT group (0.68±0.31 g/kg *vs.* 0.42±0.16 g/kg; *P*<0.01).

Table 1 Details of control and IGT groups

Variable	Control group (<i>n</i> =20)	IGT subject (n=20)
Age, y	41±4	42±6
Sex, M/F, n	13/7	12/8
Body mass index, kg/m2	21.11±1.22	20.12±1.43
Plasma glucose, mmol/L	5.22±0.89	7.23±0.73ª
Plasma insulin, pmol/L	72.00±14.83	82.95±10.23ª
Plasma glucagon, pmol/L	80.75±12.98	$90.90{\pm}15.54^{a}$
Plasma cortisol, mmol/L	0.68±0.11	$0.79{\pm}0.12^{\mathrm{b}}$
IL-6, pg/ml	3.18 ± 0.64	3.63 ± 0.43^{a}
TNF-α, pg∕ml	4.66±0.70	$5.99{\pm}0.76^{\rm b}$
APACHE score	7(4-10)	10(7-15) ^a
Sepsis score	10(8-16)	15(10-25) ^a

All data were the mean (s.d.), except APACHE \parallel score and sepsis score which were the median (range). APACHE, acute physiology and chronic health evaluation. ^a*P*<0.05 *vs.* control group. ^b*P*<0.01 *vs* control group.

Counterregulatory hormone

During the clamp, plasma insulin levels increased from a basal level of 72.00 ± 14.83 pmol/L to a peak of 83.40 ± 14.29 pmol/L within one hour (*P*<0.05) and maintained at high levels in control group. Whereas, plasma insulin levels increased more significantly in IGT group (*P*<0.01) (Figure 1). In control group, plasma glucagon levels decreased from a basal value of 80.75 ± 12.98 pmol/L to the lowest level of 74.70 ± 11.40 pmol/L within an hour (*P*<0.01) and low levels maintained, and was not significantly declined in IGT group during the entire observation period (Figure 2). Plasma cortisol was not significantly changed in two groups (Figure 3).

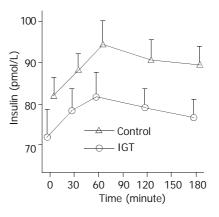


Figure 1 Circulation insulin levels during hyperglycemia clamps in 20 patients of control group (\bigcirc - \bigcirc) and in 20 patients of IGT group (\triangle - \triangle). Mean ± S.E.M. in human sepsis. Plasma insulin levels rose from a basal value to a peak within an hour (*P*<0.01) and high levels maintained in two groups.

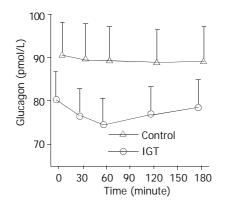


Figure 2 Circulation glucagon levels during hyperglycemia clamps in 20 patients of control group (\bigcirc - \bigcirc) and in 20 patients of IGT group (\triangle - \triangle). Mean ± S.E.M. in human sepsis. In control group, plasma glucagon levels decreased from a basal value to the lowest level within half an hour (*P*<0.01) and a low level maintained, and was not significantly declined in IGT group in the entire observation period.

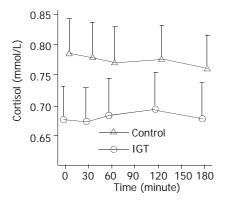


Figure 3 Circulation cortisol levels during hyperglycemia clamps in 20 patients of control group (\bigcirc - \bigcirc) and in 20 patients of IGT group (\triangle - \triangle). Mean ± S.E.M. in human sepsis. There were no significant changes in two groups.

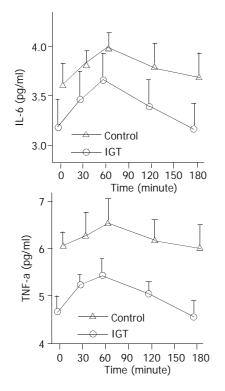


Figure 4 Circulation IL-6, TNF- α levels during hyperglycemia clamps in 20 patients of control group (\bigcirc - \bigcirc) and in 20

patients of IGT group $(\triangle - \triangle)$. Mean ± S.E.M. in human sepsis. In two groups, plasma IL-6, TNF- α levels rose from a basal value to a peak within 1 hour (*P*<0.01). In IGT group, increased level of plasma TNF- α , IL-6 during the clamping lasted longer than in control group (3 hours vs. 2 hours; *P*<0.05).

Inflammatory cytokine

In control group, plasma IL-6 levels increased from a basal value of 3.18 ± 0.64 pg/mL to a peak of 3.67 ± 0.57 pg/mL within 1 hour (P<0.01) and returned to basal level at 3 hours. Fasting plasma TNF- α levels were 4.66 ± 0.70 pg/mL, they peaked at 1 hour (5.4 ± 0.64 pg/mL, P<0.01), and returned to baseline at 3 hours. In IGT group, increased levels of plasma cytokine during the clamp lasted longer than in control group (3 hours *vs.* 2 hours; P<0.05) (Figure 4, 5).

DISCUSSION

In this study, we found that IGT group had higher admission severity scores than control group, and had high plasma concentrations of counterregulatory hormones and inflammatory cytokines. During the clamp, IGT group had a less average amount of dextrose infusion than control group. In IGT patients, a short term of hyperglycemia after glucose infusion failed to adjust the plasma concentration of counterregulatory hormones to maintain glucose homeostasis. Acute hyperglycemia in control and in IGT patients induced an increase in plasma IL-6, TNF- α concentrations and insulin levels, and the effect was amplified by IGT group. These results indicate that hyperglycemia in human sepsis with IGT is more easy to be revoked, and plays a potential modulation role in immunoinflammatory responses.

The mechanisms for "stress IGT" are well known. Human sepsis is accompanied by a marked increase in plasma concentration of counterregulatory hormones, i.e. glucagons, epinephrine, cortisol and growth hormone that affect glucose homeostasis. These hormones can lead to significant reductions in insulin sensitivity through poorly understood mechanisms likely related to alterations in insulin signal pathway^[21]. Counterregulatory hormones also enhance lipolysis and level of free fatty acids (FFA) which may contribute additional defects to the defective insulin action. Cytokine, TNF- α and IL-6 may exert their influence indirectly by stimulating counterregulatory hormone secretion and by direct action themselves^[22-24]. TNF- α and IL-6 individually and synergistically increase net glucose flux through resistance to insulin actions in muscle and liver via poorly understood postreceptor mechanisms^[25]. Hepatic insulin resistance leads to ongoing glucose production even in hyperglycemia^[26]. Peripheral insulin resistance decreases sketetal muscle glucose uptake and reduces glucose clearance, which leads to the development of IGT and even hyperglycemia.

After glucose infusion, in normal group, glucagon and glucose concentrations reduced by host insulin were sufficient and inhibited hepatic gluconeogenesis and glycogenolysis and prevented glucose production^[27]. In contrast, in IGT group, glucose infusion failed to suppress endogenous glucose production despite accompanying hyperinsulinemia. Using stable isotopes, it was demonstrated that hepatic glucose production was -150 % of the normal resting post-absorptive values of healthy subjects in spite of provision of total parenteral nutrition with dextrose at rates exceeding the basal energy expenditure^[28].

Glucose-based nutrition in septic patients may cause marked hyperinsulinemia due to peripheral insulin resistance^[29]. Hyperinsulinemia associated with glucose-based nutrition in sepsis might augment proinflammatory cytokine production and stress response in septic patients^[30]. The present findings also demonstrated that acute hyperglycemia affected concentration of plasma cytokines in human sepsis, and this effect was more pronounced in IGT patients. *In vitro* studies using supraphysiological glucose concentration (>22 mmol/l) showed an increase in TNF- α and IL-6 secretion from healthy human monouclear cells^[31]. Furthermore, increased synthesis of TNF- α has been reported both in rat uterine cells cultured *in vitro* with increasing concentrations of glucose^[32] and in placental tissue explants from women with gestational diabetes incubated with high glucose (25 mmol/l)^[33]. Human monocytes produced by IL-6 in healthy volunteers increased during 24hour incubation in high-glucose medium^[34]. These findings are in accordance with our observations *in vivo*, suggesting a potential modulation of immunoinflammatory response by carbohydrates.

In summary, IGT in sepsis is associated with marked changes in plasma concentrations of counterregulatory hormones and inflammatory cytokines, and these changes partially account for the fact that IGT easily develops acute hyperglycemia during glucose infusion. Acute hyperglycemia pricks up hyperinsulinemia and increases circulating cytokine concentrations. This suggests a potential role of hyperglycemia in inflammatory responses in human sepsis.

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