

Blood amino acids profile responding to heat stress in dairy cows

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Objective: The objective of this experiment was to investigate the effects of heat stress on milk protein and blood amino acid profile in dairy cows.

Methods: Twelve dairy cows with the similar parity, days in milk and milk yield were randomly divided into two groups with six cows raised in summer and others in autumn, respectively. Constant managerial conditions and diets were maintained during the experiment. Measurements and samples for heat stress and no heat stress were obtained according to the physical alterations of the temperature-humidity index.

Results: Results showed that heat stress significantly reduced the milk protein content ($p < 0.05$). Heat stress tended to decrease milk yield ($p = 0.09$). Furthermore, heat stress decreased dry matter intake, the concentration of blood glucose and insulin, and glutathione peroxidase activity, while increased levels of non-esterified fatty acid and malondialdehyde ($p < 0.05$). Additionally, the concentrations of blood Thr involved in immune response were increased under heat stress ($p < 0.05$). The concentration of blood Ala, Glu, Asp, and Gly, associated with gluconeogenesis, were also increased under heat stress ($p < 0.05$). However, the concentration of blood Lys that promotes milk protein synthesis was decreased under heat stress ($p < 0.05$).

Conclusion: In conclusion, this study revealed that more amino acids were required for maintenance but not for milk protein synthesis under heat stress, and the decreased availability of amino acids for milk protein synthesis may be attributed to competition of immune response and gluconeogenesis.

Keywords: Amino Acids; Heat Stress; Insulin; Milk Protein

INTRODUCTION

Global warming is one of the biggest challenges of current century [1]. Hyperthermal environment induces heat stress to dairy cows in summer and causes huge losses to the dairy economy. The decreased milk production and conception rate of dairy cows during summer depicts the heat stress on dairy cows [2]. In addition, the components of milk are strongly affected by heat stress. The greater number of somatic cells counted in milk during summer also shows that hyperthermic environment severely affects the quality of milk [3]. Heat stress also reduces the milk fat content of Holstein and Jersey cows [2]. Furthermore, it is found that a hyperthermal environment could also reduce the milk protein content via reduction of casein concentration [1].

Nutritional conditions and metabolism of dairy cows are affected by heat stress. For instance, alteration in post-absorptive metabolism of heat-stressed cows fails to enlist glucose “sparing” mechanisms [4]. Furthermore, because of reduction of feed intake and rise of maintenance requirement, heat-stressed cows may decrease the nutrient availability for milk production [5]. Heat stress has been reported to interfere with nitrogen metabolism and cause nitrogenous repartition in dairy cow, and decreases milk protein content while increasing milk urea concentration [6]. Amino acids of blood are known to synthesize the major components of milk protein in bovine mammary glands. Many studies indicate marked changes of circulating amino acids under catabolic conditions [7]. However, limited information is available concerning changes of circul-

ating amino acids under heat stress. Therefore, the objective of this study was to investigate the effect of heat stress on milk protein content and blood amino acid profile in dairy cows.

MATERIALS AND METHODS

Animals and management conditions

All procedures were carried out in strict accordance with the recommendations in the Guide for Guidelines for Experimental Animals of the Ministry of Science and Technology (Beijing, China). The protocol was approved by the Animal Care and Use Committee of Institute of Animal Science, Chinese Academy of Agricultural Sciences.

The experiment was conducted at the Sino-farm (Beijing, China). Twelve Holstein cows were assigned to two groups in a completely random design. Heat stress group (HS and $n = 6$) were performed in summer, and non-heat stress group (NS and $n = 6$) were performed in autumn. Parity, day in milk (DIM) and milk yield were similar between treatments, the DIM of these cows was 164 ± 19 d in HS and 175 ± 31 d in NS, the average values of milk yield were 29.7 ± 5.2 kg/d for HS cows and 28.5 ± 1.8 kg/d for NS cows, and the average values of parity were 1.7 ± 1.0 in HS and 1.7 ± 1.0 in NS. In addition, constant managerial conditions (milking frequency, feeding frequency, and time) and diets were maintained during the experiment. Basal ration was formulated to meet or exceed nutrient requirement recommendations [8]. Ingredients and chemical composition of the basal diet are presented in Table 1. All cows were housed in a free stall barn and were assigned to 12 electronic gate feeders (RIC system, Insentec, Marknesse, Netherlands) with individual identification transponders fitted to each cow to record the intake of each individual meal. Cows were fed the total mixed ration (TMR) daily at 0700 and 1600 h and the gate feeders were supplied with TMR to allow for 10% refusals. Cows suffered HS and NS more than three weeks before sampling, respectively. Measurements and samples were collected from July 21 to August 11, 2014 and from October 20 to November 10, 2014.

Measurements and analyses

To measure the environmental conditions inside the barn, temperatures and relative humidity were recorded daily (0600, 1400, and 2200 h) using thermometer and hygrometer in one instrument panel (Beijing Yaguang Equipment Co. Ltd., Beijing, China). The temperature-humidity index (THI) was calculated using the following formula: $THI = 1.8 \times T + 32 - 0.55 \times (1 - RH) \times (T \times 1.8 + 32 - 58)$, where RH was relative humidity expressed as fraction of the unit and T was ambient temperature expressed as degrees Celsius [1]. Rectal temperature and respiration rate of dairy cows were measured at 0700 and 1400 h twice a week. Rectal temperatures were measured using a glass mercury thermometer (Nasco, Ft. Atkinson, WI, USA), and respiration rate was determined by counting numbers of flank movements/min for 120 s.

Table 1. Ingredient and chemical composition of the basal diet

Items	Content
Ingredient (% of DM)	
Alfalfa hay	10.0
Corn silage	25.0
Oat hay	12.9
Ground corn	3.9
Soybean meal	8.6
Rapeseed meal	1.2
Extruded soybean	1.1
Dry distillers grain	7.5
Steam-flaked corn	13.1
Beet pulp	4.8
Whole cottonseed	6.6
Condensed molasses fermentation solubles	2.1
EB100 ¹⁾	0.7
XP ²⁾	0.3
Salt	0.2
Limestone	0.6
Sodium bicarbonate	0.7
Potassium bicarbonate	0.3
Premix ³⁾	0.3
Total	100
Chemical composition (%)	
NE _L ⁴⁾ (Mcal/kg)	1.67
CP	16.04
EE	5.61
NDF	36.00
ADF	23.78
Ca ⁴⁾	0.57
P ⁴⁾	0.41
Ash	7.30

DM, dry matter; NE_L, net energy for lactation; CP, crude protein; EE, ether extract; NDF, neutral detergent fiber; ADF, acid detergent fiber.

¹⁾ EB100 is a mainly saturated free fatty acid fat supplement.

²⁾ XP is a mainly saturate free amino acids supplement.

³⁾ Contained (per kilogram of DM) a minimum 250,000 IU of vitamin A; 65,000 IU of vitamin D; 2,100 IU of vitamin E; Fe 400 mg; Cu 540 mg; Zn 2,100 mg; Mn 560 mg; Se 15 mg; I 35 mg; and Co 68 mg.

⁴⁾ Determined by prediction of CPM-DairyBetav3.

Samples of TMR were collected weekly and dried for 48 h at 60°C. Samples were ground through a 1-mm sieve in a Cyclotec 1093 Mill (Tecator 1093, Hoganäs, Sweden), and residual dry matter was determined by drying at 105°C for 3 h. Chemical composition was determined according to AOAC for ash (method 942.05), crude protein (method 990.13), neutral detergent fiber (method 2002.04), acid detergent fiber (method 973.18), and ether extract (method 920.39) [9].

Cows were milked three times daily (at 0700, 1400, and 1900 h). Milk yield was recorded at each milking, and milk samples (approximately 50 mL) were collected weekly and combined at a ratio of 4:3:3 by volume (this ratio reflecting the milk yield of three times). Each milk sample was treated with potassium dichromate as a preservative and then stored at 4°C for subsequent analysis of protein using mid-infrared spectrophotometry (BactoScan FC

CombiFoss 6000, Foss Electric, Hillerød, Denmark).

On days 21 of sample collection period, duplicate blood samples were collected from the median coccygeal vein before the morning feeding into tubes with and without anticoagulant. The samples were centrifuged (3,000 g for 15 min at 4°C) to separate the plasma or serum, which were stored at -70°C for further analysis. Serum was analyzed for non-esterified fatty acid (NEFA) [10]. The concentrations of glucose and insulin in plasma were analyzed using the assay kits (Beijing Furui BioTech Ltd., Beijing, China). The levels of malondialdehyde (MDA) and the activities of glutathione peroxidase (GSH-Px), and superoxide dismutase (SOD) in plasma were determined using the assay kits (Nanjing Jiancheng Biochemical Reagent Co., Nanjing, China). Plasma amino acid profile was determined by automatic amino-acid analyzer (Hitachi, Ltd. Tokyo, Japan).

Statistical analysis

All statistical analyses were performed using the statistical software package SAS 9.3 (SAS Institute Inc., Cary, NC, USA). Milk yield, dry matter intake (DMI), rectal temperature, respiration rate, milk protein content and yield were analyzed using PROC MIXED procedure of SAS. The model included treatment, week and treatment by week interaction as fixed effects and cow as a random effect. Blood glucose, NEFA, insulin, MDA, GSH-Px, SOD, and amino acids concentration were analyzed using the general linear model procedure of SAS. The p<0.05 was considered statistically significant, and tendencies were indicated if the p value was between 0.05 and 0.10.

RESULTS

Temperature-humidity index, rectal temperature and respiration rate

The Figure 1 shows the course of daily THI during the experimental period. The THI averages were 78.1 in summer (July 21 to August 11) and 54.1 in autumn (October 20 to November 10), respectively. Compared with group NS, rectal temperature of group HS was higher at 0600 h and 1400 h (p<0.05; Table 2). Furthermore, respiration rate of group HS was also higher at 0600 h and 1400 h (p<0.05).

Milk yield, dry matter intake, and milk protein

Compared with NS, cows in HS tended to have lower milk yield (p = 0.09; Table 3). The DMI in HS was lower than that in NS (21.0 vs 17.9 kg/d, p<0.05). Milk protein content was also lower in HS (p<0.05). However, there was no difference in milk protein yield between two groups (p>0.05).

Antioxidant biomarkers

Cows of group HS had higher plasma MDA level than those in group NS (p<0.05; Table 4). However, cows in HS had lower plasma GSH-Px activity compared with NS (p<0.05). No difference was observed in SOD activity between NS and HS (p>0.05).

Blood glucose, non-esterified fatty acid, and insulin

Compared with group NS, cows of group HS had lower glucose and insulin concentrations (p<0.05; Table 5). However, cows of group HS had higher NEFA concentrations than those in group

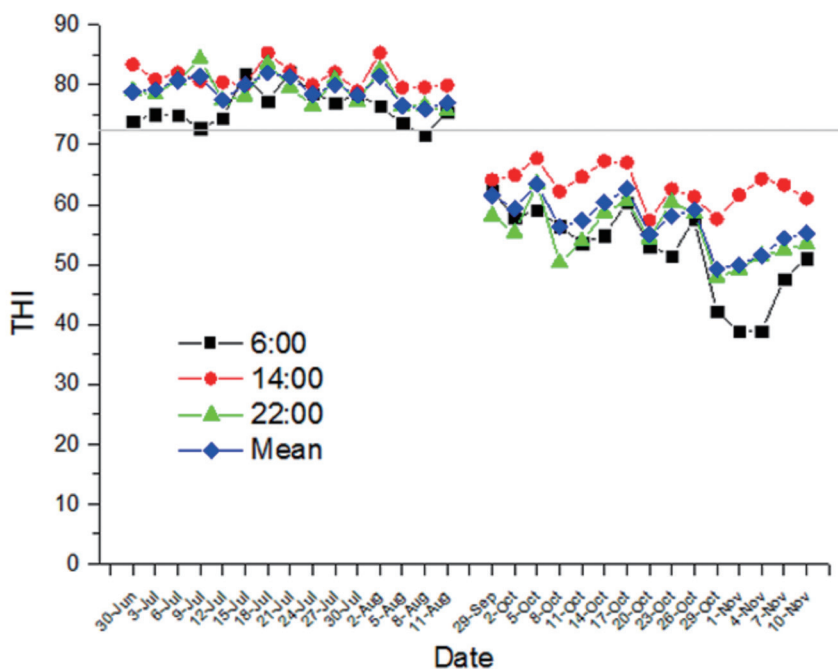


Figure 1. Temperature-humidity index (THI) at 0600, 1400, 2200 h and mean during the experiment periods.

Table 2. Rectal temperature (RT) and respiratory rate (RR) of cows observed in summer and autumn

Items	Treatment		SEM	p-value		
	NS ¹⁾	HS ²⁾		Treatment	Week	Treatment×week
RT (°C)						
0700 h	38.39	38.59	0.051	0.02	0.27	0.11
1400 h	38.41	38.91	0.099	<0.01	0.05	0.20
RR (breath/min)						
0700 h	30.0	58.6	4.52	<0.01	0.07	0.14
1400 h	36.3	73.4	4.98	<0.01	0.21	0.27

NS, non-heat stress group; HS, heat stress group; SEM, standard error of the mean.

¹⁾ Cows were raised in autumn. ²⁾ Cows were raised in summer.

Table 3. Milk yield, DMI, and milk protein content and yield of cows observed in summer and autumn season

Items	Treatment		SEM	p-value		
	NS ¹⁾	HS ²⁾		Treatment	Week	Treatment×week
Milk yield (kg/d)	29.3	25.1	1.55	0.09	0.64	0.58
DMI (kg/d)	21.0	17.9	0.71	0.01	0.92	0.19
Milk protein,						
Content (%)	3.50	3.24	0.082	<0.05	0.86	0.95
Yield (kg/d)	1.05	0.87	0.076	0.16	0.48	0.34

NS, non-heat stress group; HS, heat stress group; SEM, standard error of the mean; DMI, dry matter intake.

¹⁾ Cows were raised in autumn. ²⁾ Cows were raised in summer.

NS (p<0.05).

Plasma concentration of amino acids

It was observed that concentration of some amino acids in plasma was affected by HS (Table 6). Compared with group NS, cows of group HS had higher concentrations of total amino acids, Ala, Glu, Thr, and Gly (p<0.05). Cows tended to have higher Asp concentration in HS (p = 0.08). Only Lys concentration was lower in group HS (p<0.05). In addition, there were no significant differences in Arg, Asn, Leu, Ile, Phe, Trp, Met, Gln, Tyr, and Val concentrations between NS and HS (p>0.05).

DISCUSSION

The THI is considered as an indicator of heat stress in dairy cows. It is traditionally believed that dairy cows begin to be affected when THI exceeds 72. In the present experiment, the THI aver-

ages exceeded 72 in summer (July 21 to August 11), indicating that dairy cows were subjected to heat stress. In addition, respiration rate and rectal temperature at 1400 h exceeded 60 breath/min and 38.9°C in summer, respectively. According to Cheng et al [11], dairy cows experienced heat-stressed status during the summer in our study.

Exposure to hyperthermic environment could be responsible for oxidative stress of dairy cows. The results from our study showed that heat-stressed cows had higher level of MDA and lower activity of GSH-Px. It was consistent with previous reports, which showed that hyperthermic environment increased lipid peroxidation and inhibited the activity of antioxidant enzyme [12]. Zuo et al [13] has reported that hyperthermic environment could stimulate intracellular and extracellular superoxide formation. Hydroxyl radical generated by superoxide can attack membrane lipid and initiate lipid peroxidation [14]. Excessive accumulation of MDA may inhibit the activities of antioxidant

Table 4. The oxidative index and antioxidant enzyme activity collected in summer and autumn season

Items	Treatment		SEM	p-value
	NS ¹⁾	HS ²⁾		
MAD (nmol/mL)	2.81	3.75	0.200	0.01
GSH-Px (U/L)	121.5	88.5	6.77	0.01
SOD (U/mL)	121.9	111.2	6.55	0.28

NS, non-heat stress group; HS, heat stress group; SEM, standard error of the mean; MAD, malondialdehyde; GSH-Px, glutathione peroxidase; SOD, superoxide dismutase.

¹⁾ Cows were raised in autumn.

²⁾ Cows were raised in summer.

Table 5. Blood glucose, non-esterified fatty acid (NEFA) and insulin collected in summer and autumn

Items	Treatment		SEM	p-value
	NS ¹⁾	HS ²⁾		
Glucose (mg/dL)	34.5	29.5	1.12	<0.01
NEFA (µmol/L)	144.0	245.8	22.08	0.02
Insulin (µIU/mL)	11.27	6.14	1.273	0.02

NS, non-heat stress group; HS, heat stress group; SEM, standard error of the mean; NEFA, non-esterified fatty acid.

¹⁾ Cows were raised in autumn.

²⁾ Cows were raised in summer.

Table 6. Plasma amino acids collected in summer and autumn season

Items	Treatment		SEM	p-value
	NS ¹⁾	HS ²⁾		
Total amino acids (μmol/L)	947.8	1,198.7	38.38	<0.01
Ala	111.1	176.7	12.69	<0.01
Arg	4.70	4.05	0.541	0.42
Asn	15.7	15.2	1.30	0.82
Asp	3.90	5.30	0.483	0.08
Gln	105.3	113.3	8.11	0.24
Glu	14.6	22.0	1.82	0.02
Leu	135.2	137.1	5.86	0.83
Ile	80.3	77.0	6.31	0.72
Lys	3.23	2.16	0.228	0.01
Met	17.6	19.6	0.96	0.19
Phe	34.1	37.7	1.96	0.24
Thr	39.2	51.4	2.95	0.02
Trp	22.6	27.8	2.10	0.11
Tyr	37.0	41.6	2.60	0.25
Val	163.0	167.4	9.03	0.74
Gly	99.2	180.7	12.60	<0.01

NS, non-heat stress group; HS, heat stress group; SEM, standard error of the mean.

¹⁾ Cows were raised in autumn.

²⁾ Cows were raised in summer.

enzymes [15]. Our results suggested that heat stress might impair antioxidative ability via the depletion of enzymatic antioxidants.

It was observed that heat stress decreased DMI of dairy cow in our study. It was in accordance with a previous report, which reported that DMI was the most sensitive to air temperature [16]. However, DMI and digestive requirements increase with the increase of milk yield of dairy cow. Thus, inadequate feed intake caused by heat stress is responsible for decreased milk production. Moreover, Scott et al [17] has suggested that the DMI affects blood glucose concentration and subsequent conversion to lactose in the mammary gland. Lactose is the main osmotic regulator of lactation which can improve milk production. In the present study, heat-stressed cows had lower concentration of blood glucose, which could explain the lower milk yield in summer. In addition, NRC [8] has suggested that maintenance expenditure at 35°C increase by 20% over thermoneutral condition, thus increasing the cow's energy expenditure, often at the expense of milk yield. Per THI unit increase would decrease 0.88 kg milk yield during hot weather [16]. Our results indicated that the decline of DMI and increase of maintenance expenditure could be a major cause of reducing production of heat-stressed cows.

Our results showed that heat stress reduced insulin level of dairy cows, which might be due to lower concentration of blood glucose. De Rensis and Scaramuzzi [18] has reported that heat stress decreased DMI of dairy cows and inhibited nutrient absorption, resulting in low concentration of blood glucose and insulin. Moreover, reduction of DMI in heat-stressed dairy cows prolongs the period of negative energy balance and leads to increased concentration of NEFA [18]. Elevated NEFA causes pancreas β -cell

apoptosis and decreases insulin secretory capacity [19]. Excessive NEFA might have inhibited insulin secretion in our study. In addition, a reduction of insulin level with increasing temperature is due to the reduction in heat production [20]. However, pair-feeding experiment of heat stress has shown that heat-stressed cows have higher insulin level [21]. These inconsistent results might be due to different DMI of thermal-neutral cows.

In the present experiment, total amino acid concentrations in plasma was increased in heat-stressed dairy cows. Due to nutrient deficiency, the capacity of amino acid catabolism is likely reinforced in the liver, and amino acid concentrations are increased in plasma [22]. Additionally, higher concentrations of Glu, Gly, and Asp were observed in heat-stressed dairy cows. These results are consistent with the reports of Ai et al [23], who found that amino acids (Glu, Gly, Asp, and Val) for gluconeogenesis were higher in heat-stressed cows. When plasma glucose level is low, more limited utilization of amino acids for gluconeogenesis is observed in heat-stressed cows [6]. In addition, previous study has reported that Ala can regulate gluconeogenesis and glycolysis to ensure glucose production during periods of food deprivation [24]. We found higher Ala concentration in heat-stressed dairy cows, indicating that gluconeogenesis was likely reinforced under heat stress. It is reported that Thr is involved in immune function [25]. Higher concentrations of Thr would be associated with immune response under heat stress. However, lower Lys concentration of heat-stressed cows was observed in the present study. Lys is inadequately synthesized by the body relative to needs and needs be provided from the diet. Lower Lys concentration could be due to reduction of DMI and Lys intake in our study. The decreased Lys concentrations may limit milk protein synthesis [26]. Our results suggested that amino acids may be less used to synthesize milk protein under heat stress, but rather involved in immune response and gluconeogenesis.

Seasonal variation of milk protein level is influenced by various factors such as dietary factors, managerial factors, and climatic conditions. Constant managerial conditions and diets were maintained during the trial periods, which facilitated the observation of the effect of climatic conditions on milk protein. Heat-stressed dairy cows had lower milk protein content in our study, which is defined as "heat-stressed milk protein decrease syndrome". We found that reduction of milk protein content might be due to low level of circulating insulin. Burgos et al [27] has reported that insulin is a potent anabolic hormone and which can promote mammary protein synthesis. In addition, competition for amino acids that supply both milk protein and maintenance metabolism could be a possible reason of the decreased milk protein content under heat stress. Dairy cows could utilize more intensive amino acids as precursors for gluconeogenesis during heat stress [28]. Immune response and gluconeogenesis depleted a large number of amino acids, thus decreased the availability of amino acids for milk protein synthesis in our study. Similarly, Cowley et al [6] also has reported that the synthesis of the heat shock proteins may

decrease the availability of amino acids for milk protein synthesis. However, Naderi et al [29] has suggested that substituting beet pulp for corn silage in the diet of heat-stressed cows results in a high milk protein content, corresponding to low rumen ammonia-nitrogen concentration and milk urea concentration. Heat-stressed cows fed the protein containing high Lys could increase milk yield and milk protein yield [30]. The exact mechanisms need to be further studied.

CONCLUSION

In summary, the results from the current study indicate that heat stress reduces milk protein content, glucose and insulin concentrations in blood, and changes amino acids profile of dairy cows, suggesting that more amino acids are required for maintenance (immune response and gluconeogenesis) but not for milk protein synthesis under heat stress, which provides scientific experimental bases for how to alleviate the detrimental effect of heat stress in dairy cow production.

CONFLICT OF INTEREST

We certify that there is no conflict of interest with any financial organization regarding the material discussed in the manuscript.

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