# Effects of Cooling and Supplemental Bovine Somatotropin on Milk Production relating to Body Glucose Metabolism and Utilization of Glucose by the Mammary Gland in Crossbred Holstein Cattle

<sup>1</sup>Siravit Sitprija, <sup>2</sup>Somchai Chanpongsang and <sup>1</sup>Narongsak Chaiyabutr <sup>1</sup>Department of Veterinary Physiology, <sup>2</sup>Department of Animal Husbandry, Faculty of Veterinary Science, Chulalongkorn University, Bangkok 10330, Thailand

Abstract: Problem statement: The low milk yield and shorter persistency of lactation of dairy cattle is the major problem for the dairy practices in the tropics. High environmental temperatures and rapid decline of plasma growth hormone level can influence milk production. Regulation of the milk yield of animals is mainly based on the mechanisms governing the quantity of glucose extracted by the mammary gland for lactose biosynthetic pathways. The mechanism(s) underlying the effects of cooling and supplemental bovine somatotropin on milk production relating to body glucose metabolism and intracellular metabolism of glucose in the mammary gland of crossbred Holstein cattle in the tropics have not been investigated to date. Approach: Ten crossbred 87.5% Holstein cows were divided into two groups of five animals each. Animals were housed in Normal Shade barn (NS) as non-cooled cows and cows in the second group were housed in barn which was equipped with a two Misty-Fan cooling system (MF) as cooled cows. Supplementation of recombinant bovine Somatotropin (rbST) (POSILAC, 500 mg per cow) were performed in both groups to study body glucose metabolism and the utilization of glucose in the mammary gland using a continuous infusion of [3-<sup>3</sup>H] glucose and [U-<sup>14</sup>C] glucose as markers in early, mid and late stages of lactation. **Results:** Milk yield significantly increased in both groups during supplemental rbST with a high level of mammary blood flow. Body glucose turnover rates were not significant different between cooled and non-cooled cows whether supplemental rbST or not. The glucose taken up by the mammary gland of both non-cooled and cooled cows increased flux through the lactose synthesis and the pentose cycle pathway with significant increases in NADPH formation for fatty acid synthesis during rbST supplementation. The utilization of glucose carbon incorporation into milk appeared to increase in milk lactose and milk triacylglycerol but not for milk citrate during supplemental rbST in both non-cooled and cooled cows in early and mid lactation. Conclusion: The present study demonstrated that local changes for biosynthetic capacity within the mammary gland would be a factor in identification of the utilization of substrates in the rate of decline in milk yield. The proportion of glucose was metabolized less for lactose synthesis, but metabolized more via the Embden-Meyerhof pathway and the tricarboxylic acid cycle as lactation advanced to late lactation in both cooled and non-cooled cows whether supplemental rbST or not.

Key words: rbST, glucose metabolism, mammary gland, crossbred Holstein cow

# INTRODUCTION

The low milk yield and short lactation period of either pure exotic or crossbred dairy cattle is still the major problem for the dairy practices in the tropics. The mechanisms that limit the rate of milk yield and shorter lactation persistency as lactation advances in crossbred dairy cattle in tropics are unclear. It is not only animal genetics that are considered but other factors, for example, high environmental temperatures and hormonal factors can influence milk production (Collier *et al.*, 1982). The study in 87.5% crossbred Holstein cattle (HF) showed rapid decline in the concentration of plasma bovine Somatotropin (bST), which would accompany with a reduction in both mammary blood flow and milk yield as lactation progressed to mid and late lactation (Chaiyabutr *et al.*, 2000a). Many studies demonstrated the efficacy of bST for improvement in milk yield (Breier *et al.*, 1991; Burton *et al.*, 1994). Long term exogenous recombinant

**Corresponding Author:** Narongsak Chaiyabutr, Department of Veterinary Physiology, Faculty of Veterinary Science, Chulalongkorn University, Bangkok 10330, Thailand

bovine Somatotropin (rbST) in 87.5% HF cow increased in milk yield which accompanied with an increase in the rate of mammary blood flow, but the stimulant effect for milk yield was less in late lactation despite a high level of mammary blood flow (Chaiyabutr *et al.*, 2007). It is not known which factors are the cause and which factors are the effects for such reduction.

It is well recognized that prolonged high ambient temperatures can affect either directly or indirectly to the ability of dairy cow to synthesize milk production. Environmental modifications such as fans and sprinklers (Fike et al., 2002) or evaporative cooling system (Chan et al., 1997; Chaiyabutr et al., 2008a) can be used to alleviate of severe heat stress and increase in milk production in dairy cattle. In addition to environmental modifications, other technologies can increase milk production in dairy cattle, for example, by supplementation of dietary methionine (Yang et al., 2010), or the application of exogenous bovine somatotropin (West et al., 1991; West, 1994). Somatotropin is known to play a role in responsible for galactopoietic and contributing to homeostasis and homeorhesis in ruminants (Bauman and Currie, 1980). However, few data are available on the combined effects of high environmental temperatures and bST administration on mechanisms of milk secretion.

Glucose is known to be the principal precursor of lactose synthesis. Lactose is the major osmotic factor of milk synthesis and is required in proportion to the amount of milk produced (Linzell and Peaker, 1971). The regulation of the milk yield is mainly based on the quantity of glucose extracted by the mammary gland and converted into lactose. The rapid decline in lactose biosynthetic pathways has been shown to account for a short persistency of lactation as lactation advanced to mid and late lactation in 87.5% HF cows in either without rbST administration (Chaiyabutr et al., 2000b) or long-term administration of rbST (Chaiyabutr et al., 2008b). During long-term administration of rbST, the utilization of glucose in the mammary gland was metabolized less for lactose synthesis and the pentose phosphate pathway but metabolized more via the Embden-Meyerhof pathway as lactation advances (Chaiyabutr et al., 2008b). Cows treated with bST does not only increase efficiencies of milk yields, but also increase heat production, which was probably due to increased metabolic activity associated with higher milk yield (West, 1994). The high heat production in rbST treated-cows with high ambient temperatures would induce high heat stress. The further study would be established whether short persistency of lactation occurring in 87.5% HF cows is due to the effect of high

ambient temperatures or the less stimulant effect of bovine somatotropin or combination of both of these factors during lactation advances. Environmental modifications are needed to minimize the effects of heat stress and that will maintain potentially increased milk yields because of bST technology (West, 1994). Few data are available concerning glucose utilization by the udder, under the effect of cooling and supplementation of rbST in crossbred Holstein cattle. Therefore, the present study was designed to investigate the mechanisms of milk secretion relating to intracellular metabolism of glucose in the mammary gland body glucose metabolism during and rbST supplementation in 87.5% HF cows housing under mistyfan cooling system.

# MATERIALS AND METHODS

Animals and managements: Ten primiparous, crossbred 87.5% Holstein cattle were randomly divided into two groups of five animals each. Animals in the control group were housed in the Normal Shade (NS) in individual stall, while animals in the experimental group were housed in shade with using mister and fans cooling to reduce the environmental temperature (MF). The MF barn had two sets of misters and fans cooling system, which each system consisted of a 26 inch diameter blade fan circulating 7,200 ft<sup>3</sup> min<sup>-1</sup> of air, with oscillation coverage of 180°. The amount of water discharged from 4 spray heads was 7.5 L  $h^{-1}$  and side of mist droplet 0.01 mm. Animal were exposed to MF for 45 min at 15 min intervals from 0600-1800 h. At night, animals were exposed to MFC for 15 min at 45 min intervals from 1800-0600 h. Animals in each group were fed with the same ration of TMR (Table 1) twice daily throughout the experiment. Each day, the diet was given in equal portion at about 0600 and 1700 h when animal were milked. Water was available at all times. All animals were weighed monthly throughout the experiment.

The study was performed under a protocol approved by ethic committee of Faculty of Veterinary Science, Chulalongkorn University. The procedures used in the present study were formulated to comply with international standards and are in accordance with the principles and guidelines of the National Research Council of Thailand.

**Experimental procedures:** The diagram illustrating the time course of the experiment is shown in Fig. 1. The experiments were carried out throughout lactating periods in each group. The experiment in each group was divided into 3 stages, namely early-(Days 65-95 postpartum), mid-(Days 125-155 postpartum) and late lactating stages (Days 185-215 postpartum).



Fig. 1: Diagram illustrating the time course of studies in each cow supplemented with rbST at different stages of lactation. Pretreat = timed study for pre-treatment; Treat = timed study for treatment

Table 1: Ingredients and nutrient compositions of the TMR diet

Ingredients	Kg (as fed basis)
Pine apple waste	50.0
Soybean meal	23.0
Rice bran	3.0
Cotton seed	20.0
Lime stone	1.4
Di-calcium phosphate	1.4
Sodium bicarbonate	0.3
Potassium chloride	0.1
Mineral and vitamin premix	0.8
Total	100.0
Nutrient compositions	
Dry matter (%)	39.1
Ash (% DM)	7.3
Organic matter (% DM)	92.7
Crude protein (% DM)	18.0
Acid detergent fiber (% DM)	20.1
Neutral detergent fiber (% DM)	33.9

The pretreatment study was conducted on the starting day of each lactating stage. At the end of the pretreatment, within the same day, the animal was injected with the first dose subcutaneous injection of 500 mg of recombinant bovine Somatotropin (rbST) (POSILAC, Monsanto, USA). Subsequently, the animal was injected with two consecutive doses injections of rbST in every 2 weeks. Thereafter, within 2 days after the third injection, the treatment study was conducted. The pretreatment, 3 doses of injections and the treatment periods were performed during the first 30 days and the same procedures were followed for each lactating stage. During the last 30 days of each lactating stage, no experiments were conducted in order to allow the milk yield from the effect of rbST treatment to return to the control level (Kirchgessner et al., 1991).

Rectal temperature and respiration rate of individual animals were determined at the same time as recording ambient temperature. Ambient temperature and relative humidity were measured weekly throughout the experiment. The Temperature Humidity Index (THI) was calculated according to West (1994), where: THI = db-(0.55-0.55RH) (db-58) with db = dry bulb temperature ( $^{\circ}F$ ) and RH = relative humidity. On

each specified day, measurements of mammary blood flow, glucose metabolism and the utilization of glucose by the mammary gland were carried out at around 10.00 h. Both ear vein and milk vein were catheterized with the non-radiopaque intravenous catheters, gauge 18G (Surflo, Terumo Europe N.V., Belgium) under local anesthesia for infusion of solution. An arterial blood sample was collected from the coccygeal vessel by venipuncture with a # 21 needle into heparinized tube. In the present study, blood sample from either tail vessel can be accounted for arterial blood sample. Since, it has been demonstrated in dairy cattle by Emery et al. (1965) that concentrations metabolites such as glucose and acetate in tail arterial blood were similar to those of tail venous blood. Blood samples from arterial and mammary venous blood in heparinized tube were kept in crushed ice and then centrifuge at 3000 rpm for 30 min at 4°C. Plasma samples were collected and frozen at -40°C in aliquots until time of assays for measurements the concentration of metabolites.

Glucose turnover measurements: The study of glucose kinetic and efficiency of glucose utilization by the mammary gland during rbST supplementation in both cooled and non-cooled cows were performed at different stages of lactation. Glucose kinetic studies were carried out by using continuous infusion of both [U-<sup>14</sup>C]-glucose and [3-<sup>3</sup>H]-glucose solution as described previously by Chaiyabutr et al. (1998). Briefly, at about 10.00 h of the specified day, a priming dose of radioactive glucose in 20 mL of sterile NSS containing 30 µCi [3-<sup>3</sup>H]-glucose and 15 µCi[U-<sup>14</sup>C]glucose was administered intravenously via the ear vein catheter and followed by a continuous infusion of 1 mL min<sup>-1</sup> of normal saline solution (0.9%) containing 0.7  $\mu$ Ci mL<sup>-1</sup> of [U-<sup>14</sup>C]-glucose and 1.5  $\mu$ Ci mL<sup>-1</sup> of [3-<sup>3</sup>H]-glucose for 3 h. (Peristatic pump; EYLA Model 3). During the last 1 h (1200-1300 h) of continuous infusion, three sets of blood samples were collected at

20 min intervals. A venous blood sample was collected from the milk vein via a catheter while an arterial blood sample was collected from the coccygeal vessel by venipuncture with a # 21 needle. Blood samples in heparinized tubes were kept in crushed ice for chemical studies. Milk secretion was recorded for the last 1 h of continuous infusion. Milk samples were used for measurement of radioactive glucose incorporation into other milk components. Milk yield was recorded by weight.

**Mammary blood flow measurement:** Blood flow through half of the udder were performed in duplicated by dye dilution technique using dye T-1824 (Evans blue) by a short term continuous infusion into the milk vein as described by Chaiyabutr *et al.* (1997). The rate of blood flow through half of the udder was calculated from plasma sample and the value of packed cell volume using the equation derived by Thompson and Thomson (1977). Quarter milking showed that the yields of the two halves of the udder were similar. Mammary blood flow was therefore calculated by doubling the flow measured in one milk vein (Bickerstaffe *et al.*, 1974). Packed cell volume was measured after centrifugation of the blood in a microcapillary tube.

**Chemical methods:** Radiochemicals for [U-<sup>14</sup>C]glucose and [3-<sup>3</sup>H]-glucose were obtained from the Radiochemical Center, Amersham Bucks, UK. The specific activity of labeled plasma glucose was determined by the method described by Chaiyabutr and Buranakarl (1989). The plasma glucose concentration was measured using enzymatic oxidation in the presence of glucose oxidase (Human GmBH, Germany). Plasma free fatty acid were determined by colorimetry after plasma extraction with chloroform, heptane and methanol and TAN solution (Wang et al., 2004). Plasma triacylglycerol concentration was determined by enzymatic colorimetric test (Triglyceride liquicolor, Wiesbaden, Germany). The concentration of milk lactose was determined by spectrophotometry (Teles et al., 1978). Lactose radioactivity was determined after isolation by the hydrolysis method (Wood et al., 1965). Milk fatty acids was extracted from 1 ml of an aliquot thawed milk in 2 mL of Dole's solution (Dole, 1956), (iso-propanol 40: n-heptane 10: 1N H<sub>2</sub>SO<sub>4</sub> 1, v/v) shaking in water bath for 30 min. After 1 ml hexane and 1ml H<sub>2</sub>O was added to the vial and shaking, the upper layer containing fatty acids was transferred into two vials for radioactivity assay and for determination of milk fatty acids concentration. Milk extraction solution in counting vial with a scintillation cocktail was measured radioactivity of <sup>14</sup>C and <sup>3</sup>H-fat by liquid scintillation counter. Other portion of milk extraction was used to determine milk fatty acids concentration by colorimetry according to Wang et al. (2004) using chloroform, heptane and methanol and TAN solution containing 1-(2-Thiazolylazo)-2-naphthol (Sigma-Aldrich). Milk fatty acid profiles were determined by gas chromatography (GC-2010 Gas Chromatograph, Shimazu) after extraction by chloroform and methanol (Christopherson and Glass, 1969) in comparison with the appropriate internal standard of pentadecanoic acid (C:15:0 fatty acid). The concentration of milk citrate was determined by spectrophotometry from tricarboxylic acid filtrate (White and Davies, 1963). Citrate radioactivity was determined after isolation by anion exchange chromatography (Hardwick et al., 1963).

**Calculations:** According to Chaiyabutr *et al.* (1980; 2008b), body glucose metabolism and intramammary glucose metabolism can be calculated as follow.

The glucose turnover rate in the whole animal (T), expressed as  $\mu$ mol min<sup>-1</sup>, was calculated from the equation:

$$T = I/G_A$$

Where:

- I = Rate of infusion of  $[U^{-14}C]$  glucose or  $[3^{-3}H]$ glucose ( $\mu$ Ci min<sup>-1</sup>)
- $G_A$  = Specific activity of <sup>14</sup>C- or <sup>3</sup>H-glucose in arterial plasma at equilibrium ( $\mu$ Ci  $\mu$ mol<sup>-1</sup>)

Recycling of glucose carbon in the whole animal, expressed as % glucose turnover, was calculated from the equation:

$$Recycling = (T_3 - T_{14}) \times 100/T_3$$

Where:

- $T_3$  = Reversible turnover of glucose calculated from [3-<sup>3</sup>H] glucose
- $T_{14}$  = Irreversible turnover of glucose calculated from  $[U^{-14}C]$  glucose

The metabolic glucose clearance rate in the whole animal ( $C_G$ ), expressed as mL min<sup>-1</sup>, was calculated from the equation:

$$C_G = T_3 / P_{AG}$$

Where:

 $T_3$  = Reversible turnover of glucose calculated from  $3^{-3}H$  glucose (µmol min<sup>-1</sup>)

 $P_{AG}$  = Arterial plasma glucose concentration (µmol mL<sup>-1</sup>)

Uptake of glucose by the udder (U<sub>G</sub>), expressed as  $\mu$ mol min<sup>-1</sup>, was calculated from the equation:

 $U_G = MPFx (P_A P_V)$ 

Where:

MPF = Mammary plasma flow (mL min<sup>-1</sup>)

- $P_A$  = Concentration of glucose in coccygeal arterial plasma (µmol mL<sup>-1</sup>)
- $P_V$  = Concentration of glucose of plasma from milk vein (µmol mL<sup>-1</sup>)

The milk component output (MO), expressed as  $\mu$ mol min<sup>-1</sup>, was calculated from the equation:

 $MO = Ms \ x \ Cc/1000$ 

Where:

Ms = Milk secretion rate (Ml min<sup>-1</sup>)

Cc = Concentration of components in milk ( $\mu$ mol L<sup>-1</sup>)

Incorporation (A) of radioactivity from glucose into milk components was calculated from the equation:

 $A = M_A/G_A \ge t$ 

Where:

- A = Incorporation of radioactivity from glucose into milk components ( $\mu$ mol min<sup>-1</sup>)
- $M_A$  = Total activity of <sup>3</sup>H or <sup>14</sup>C in the milk components ( $\mu$ Ci)

 $G_A$  = Specific activity of <sup>14</sup>C-or <sup>3</sup>H-glucose in arterial plasma at equilibrium ( $\mu$ Ci  $\mu$ mol<sup>-1</sup>)

t = Time of infusion (min)

Requirement of NADPH for fatty acid synthesis (P) in the mammary gland, expressed as  $\mu$ mol min<sup>-1</sup>, was calculated from the equation:

$$P_{\text{NADPH}} = \sum [FFA_n \times (n-2)]$$

Where:

n = Chain length of the fatty acid (6-16)

 $FFA_n$  = Output in milk of fatty acid chain length n ( $\mu$ mol min<sup>-1</sup>)

Values for  $FFA_n$  were calculated from all medium chain length fatty acids and 30% of  $C_{16}$ -fatty acids (Annison and Linzell, 1964).

Net metabolism of Glucose Phosphorylation (G<sub>6</sub>P), expressed as  $\mu$ mol min<sup>-1</sup>, was calculated from the equation:

 $G_{6}{}^{P} = U_{G} \text{-} L$ 

Where:

 $U_G = Mammary glucose uptake (\mu mol min<sup>-1</sup>)$ 

L = Output of lactose in milk ( $\mu$ mol min<sup>-1</sup>)

Net metabolism of glucose (B) to the galactose or glucose moiety of lactose, expressed as  $\mu$ mol min<sup>-1</sup>, was calculated from the equation:

B = L

where, L output of lactose in milk ( $\mu$ mol min<sup>-1</sup>).

Metabolism of glucose via the pentose phosphate pathway (PC) was calculated from the equation:

Y = 3 PC/(1+2PC)

where, Y specific yield of  ${}^{14}CO_2$  from (1- ${}^{14}C$ ) glucose via the pentose phosphate pathway (Katz and Wood, 1963).

If the NADPH formed via PC were used exclusively for reductive biosynthesis of fatty acids, the <sup>3</sup>H-incorporation from [3-<sup>3</sup>H] glucose into fatty acids would equal the <sup>14</sup>CO<sub>2</sub> released from [1-<sup>14</sup>C] glucose via the pentose phosphate pathway (Katz *et al.*, 1974). Metabolism of glucose via PC was therefore calculated from the equation:

Z = 3 PC/(1+2PC)

where, Z (Total <sup>3</sup>H in milk fatty acid)/t×G<sub>A</sub>×(U<sub>G</sub>-L)

Net metabolism of glucose 6-phosphate via PC ( $G_{PC}$ ), expressed as  $\mu$ mol min<sup>-1</sup>, was calculated from the equation:

 $G_{PC} = G_6 P \times PC$ 

Net metabolism of glucose 6-phosphate via the Embden-Meyerhof pathway ( $G_E$ ), expressed as  $\mu$ mol min<sup>-1</sup>, was calculated from the equation:

 $G_{\rm E} = G_{\rm 6p} - (B + G_{\rm PC})$ 

The  ${}^{3}\text{H}/{}^{14}\text{C}$  ratio in the plasma and related product was calculated from the equation:

 ${}^{3}\text{H}/{}^{14}\text{C}$  glucose =  ${}^{3}\text{H}/{}^{14}\text{C}$  in plasma glucose relative to  ${}^{3}\text{H}/{}^{14}\text{C}$  ratio of 1 in the infusion

<sup>3</sup> H/ <sup>14</sup> C lactose	=	${}^{3}\text{H}/{}^{14}\text{C}$ in milk lactose relative to ${}^{3}\text{H}/{}^{14}\text{C}$ ratio of 1 in the
<sup>3</sup> H/ <sup>14</sup> C galactose	=	infusion $2(^{3}\text{H}/^{14}\text{C} \text{ lactose})-(^{3}\text{H}/^{14}\text{C})$
<sup>3</sup> H/ <sup>14</sup> C citrate	=	glucose) ${}^{3}\text{H}{}^{14}\text{C}$ in milk citrate relative to ${}^{3}\text{H}{}^{14}\text{C}$ ratio of 1 in the
<sup>3</sup> H/ <sup>14</sup> C triacyglycerol	=	infusion ${}^{3}\text{H}/{}^{14}\text{C}$ in milk triacyglycerol relative to ${}^{3}\text{H}/{}^{14}\text{C}$ ratio of 1 in
		the infusion

**Statistical analysis:** Individual cow data in each stage of lactation were adjusted for covariate effects with the data from pretreatment period before start of treatment period. The statistic analyses were performed using General Linear Model procedure of statistical software package SPSS (SPSS for windows, V14.0; SPSS Inc., Chicago, IL, USA) to study either effects or interaction effects of treatment and housing. The model used for each parameter analysis was:

$$\begin{split} Y_{ijk} &= \mu + A_l + H_i + A(H)_{il} + B_j + (HB)_{ij} + A(HB)_{ijl} + \\ Cov_k + e_{iikl} \end{split}$$

Where:

- B<sub>j</sub> = Treatment effect (rbST) as a split plot (j = with and without rbST supplementation)
- $(HB)_{ij}$  = Interaction effect between treatment and house

 $Cov_k$  = Covariate effect  $e_{ijk}$  = Residual error

The statistical significant differences of environmental parameters between NS and MF barn were determined by unpaired t-test. Statistical significance was declared at p<0.05 and trends were declared at 0.05 .

#### RESULTS

Ambient temperature, relative humidity, Temperature Humidity Index (THI) respiratory rate and rectal temperature: Mean values of measurements at experimental site during periods of studies for daily temperatures, humidities and THI are shown in Table 2. The rectal temperature and respiratory rate are shown in Table 3. Average values of ambient temperature in the barn during the daytime in the morning (0900 h) between NS barn and MF barn were not significantly different, while ambient temperatures during 1400 h at NS barn were significantly higher than that of MF barn. The high relative humidity was apparent at morning and it decreased onwards from morning to evening in both NS and MF barns, whereas relative humidity in MF barn was significantly higher than that of NS barn.

Table 2: Ambient temperature, Relative humidity and temperature humidity index in Normal Shade (NS) barn and shade plus Misty-Fan cooling (MF) barn

			Treatments		
Parameter	Stage of lacta	ition	NS (n = 5)	MF (n = 5)	p-value
Ambient	Early	0900 h	27.8±0.67	27.3±0.68	0.275
Temperature (°C)		1400 h	34.9 ±1.29	30.7±1.38	0.001
• · ·	Mid	0900 h	28.3±0.67	27.3±0.70	0.052
		1400 h	33.3±1.13	30.4±0.89	0.002
	Late	0900 h	$28.4 \pm 0.90$	27.4±0.80	1.000
		1400 h	$32.4 \pm 1.29$	$29.0 \pm 1.09$	0.002
Relative humidity (%)	Early	0900 h	79.3±3.67	86.4±3.08	0.011
		1400 h	53.5±5.62	67.9±7.91	0.011
	Mid	0900 h	78.7±2.83	84.4±2.86	0.013
		1400 h	58.8±3.04	73.5±5.88	0.001
	Late	0900 h	75.4±3.94	82.4±3.52	0.018
		1400 h	60.9±6.71	$71.8\pm8.81$	0.059
Temperature	Early	0900 h	78.6±0.73	78.6±0.88	1.000
Humidity Index (THI)		1400 h	85.4±1.10	81.9±1.67	0.001
	Mid	0900 h	79.3±0.88	78.4±0.99	0.176
		1400 h	84.0±0.97	82.4±1.17	0.046
	Late	0900 h	78.9±1.10	78.3±1.01	0.395
		1400 h	83.2±0.97	79.9±1.01	0.001

Mean  $\pm$  SD = Standard Deviation of the mean

			NS		MFC			<sup>1</sup> Effect		
	Stages of									
Parameter	lactation		Pre	rbST	Pre	rbST	SEM	rbST	MFC	rbSTxMFC
Rectal temperature (°C)										
-	Early	0900 h	38.5	38.9	38.1	38.5	0.08	0.003	0.032	1.000
		1400 h	39.4	40.0	39.0	39.4	0.21	0.061	0.037	0.817
	Mid	0900 h	38.5	39.0	38.0	38.3	0.08	0.001	0.024	0.258
		1400 h	39.7	40.1	38.6	39.5	0.13	0.002	0.002	0.090
	Late	0900 h	38.5	38.8	38.0	38.3	0.10	0.015	0.007	0.723
		1400 h	39.2	39.9	38.4	38.8	0.16	0.016	0.015	0.309
Respiratory rate (breaths/min	)									
	Early	0900 h	40.0	42.5	35.0	38.0	0.56	0.003	0.003	0.670
		1400 h	73.0	82.3	55.5	68.0	4.13	0.039	0.023	0.708
	Mid	0900 h	41.2	45.8	36.4	40.4	0.66	0.001	0.022	0.663
		1400 h	73.6	77.2	49.0	57.6	1.96	0.018	0.001	0.294
	Late	0900 h	40.5	44.5	37.0	41.5	1.44	0.025	0.101	0.868
		1400 h	71.5	80.0	54.3	59.3	1.09	0.001	0.019	0.159

Table 3: Ambient temperature, Relative humidity, temperature humidity index, mean values of rectal temperature and respiratory rate of crossbred Holstein cows housing in Normal Shade (NS) and shade plus Misty-Fan Cooling (MFC)

SEM = Standard error of the mean.<sup>1</sup> P-values for the effects; MFC = Misty-Fan Cooling effect, rbST = rbST effect, MFC x rbST = interaction effect of MFC and rbST

THI values at the MF barn in afternoon were significantly lower than that of NS barn. Cows in both groups exposed to high THI values (78.3-85.4) in both barns. Rectal temperature recording in the morning and afternoon (0900-1400 h) of cooled and non-cooled cows were significant different whether rbST injection or not. The rectal temperatures of cooled cows were lower than those of non-cooled cows during afternoon (1400 h). There were significant increases in rectal temperature and respiration rate by the effect of supplemental rbST in different parts of the day. The respiratory rates of cooled cows were significantly lower than those of non-cooled cows throughout experimental periods.

Milk yield, milk compositions and its secretion: Milk yield, milk compositions and its secretion of cooled and non-cooled cows are shown in Table 4. It is obvious that milk yield was significantly increased by rbST of both cooled and non-cooled cows, but it decreased as lactation advances. Milk lactose concentrations were not affected by rbST supplementation as compared with pretreatment in both groups or among stages of lactation in the same group. The ratio of lactose output/glucose uptake were not different in comparison between cooled and non-cooled cows whether supplemental rbST or not in each stage of lactation, but it showed tendency to decrease as lactation advances., The milk lactose secretion was significantly increased by rbST in both cooled and non-cooled cows in eac stage of lactation. The milk citrate concentration was significantly increased during supplemental rbST in early lactation, while its significantly decreased in late lactation in both cooled and non-cooled cows. However, during early and mid lactation, the secretions of milk citrate were significantly increased by rbST in

both cooled and non-cooled cows. The concentration triacylglycerol in milk had tendency to increase during supplemental rbST, but a significant increases were apparent in early lactation in both groups. The secretions of milk triacylglycerol were significantly increased in both cooled and non-cooled cows during rbST supplementation in each stage of lactation.

Mammary plasma flow, plasma glucose concentration, mammary glucose uptake and percentage of glucose extraction: The utilization of glucose across the mammary gland during rbST supplementation in both cooled and non-cooled cows are shown in Table 5. Mammary plasma flow of both cooled and non-cooled cows were significantly increased by rbST administration in each stage of lactation. During rbST supplementation mammary glucose uptake increased in each stage of lactation in both cooled and non-cooled cows. The mammary glucose uptake of both non-cooled and cooled cows were significantly increased during supplemental rbST in mid and late lactations by average 37% and 34%, respectively. Plasma glucose concentrations were not affected by rbST throughout lactation in both cooled and non-cooled cows. There were no significant changes in A-V concentration differences for glucose across the mammary gland during rbST supplementation in each stage of lactation. The percentage of glucose extraction was not influenced by the supplementation of rbST in both groups.

**Glucose turnover and related variables:** Simultaneous estimation of the total glucose entry rate using  $3-[^{3}H]$  glucose infusion and the utilization rate of glucose using  $[U^{-14}C]$  glucose infusion of both cooled and non-cooled cows supplemental rbST are shown in Table 6.

		NS		MFC			<sup>1</sup> Effect		
Parameter	Stages of lactation	Pre	rbST	Pre	rbST	SEM	rbST	MFC	rbSTxMFC
Milk vield (kg dav <sup>-1</sup> )									
	Early	13.39	15.43	14.82	15.84	0.31	0.001	0.684	0.140
	Mid	11.13	13.10	13.79	15.73	0.54	0.003	0.269	0.549
	Late	10.31	11.77	11.29	15.00	0.61	0.003	0.372	0.101
Lactose concentration (mmol L <sup>-1</sup> )									
	Early	132.5	135.60	133.20	134.00	1.51	0.231	0.833	0.467
	Mid	129.3	130.60	130.50	131.20	1.29	0.658	0.536	0.195
	Late	130.5	129.70	131.70	133.60	1.68	0.769	0.338	0.448
Milk lactose secretion (µmol min <sup>-1</sup> )									
	Early	1230.3	1458.40	1367.70	1471.30	36.12	0.002	0.716	0.123
	Mid	999.2	1188.00	1249.70	1497.50	48.38	0.003	0.225	0.100
	Late	936.6	1066.30	1028.60	1392.50	57.35	0.003	0.347	0.075
Lactose output/Glucose uptake (%)									
	Early	65.1	70.40	60.10	57.70	5.60	0.799	0.537	0.510
	Mid	62.8	48.80	66.40	54.60	5.90	0.678	0.994	0.702
	Late	40.8	34.40	36.70	43.90	3.00	0.903	0.668	0.055
Milk citrate concentration (mmol L <sup>-1</sup> )									
	Early	4.24	4.54	4.22	4.85	0.15	0.014	0.305	0.303
	Mid	4.7	4.71	5.67	5.78	0.11	0.575	0.016	0.645
	Late	4.74	4.14	5.24	4.38	0.15	0.001	0.042	0.404
Milk citrate secretion (µmol min <sup>-1</sup> )									
	Early	39.51	48.81	43.81	52.95	1.74	0.001	0.578	0.965
	Mid	36.21	42.16	54.48	67.00	2.82	0.011	0.078	0.277
	Late	33.72	33.90	41.39	45.56	2.26	0.364	0.228	0.402
Milk triacylglycerol concentration (mm	ol L <sup>-1</sup> )								
	Early	42.36	48.50	45.95	58.66	3.87	0.041	0.361	0.420
	Mid	58.77	56.92	57.53	64.99	3.12	0.395	0.699	0.174
	Late	61.13	69.36	54.67	66.41	5.61	0.113	0.503	0.762
Milk triacylglycerol secretion (µmol mi	n <sup>-1</sup> )								
	Early	374.05	491.3	483.14	632.77	40.09	0.010	0.105	0.697
	Mid	446.56	510.5	519.84	710.27	48.33	0.030	0.1	0.227
	Late	433.02	569 36	415 35	688 55	81 45	0.036	0.59	0.425

Table 4: Milk yield, milk compositions and secretion for milk lactose, milk citrate and milk triacylglycerol during rbST administration at different stages of lactation of cows housing in Normal Shade (NS) and shade plus Misty-Fan Cooling (MFC)

SEM = Standard Error of the Mean; <sup>1</sup>P-values for the effects; MFC = Misty-Fan Cooling effect; rbST = rbST effect; MFC x rbST = Interaction effect of MFC and rbST

Table 5: Mammary plasma flow, arterial plasma glucose concentration, mammary glucose uptake and percentage of glucose extraction during rbST administration at different stages of lactation of cows housing in Normal Shade (NS) and shade plus Misty-Fan Cooling (MFC)

	U		U		,	/	1	2	<u> </u>
		NS		MFC			<sup>1</sup> Effect		
	Stages of								
Parameter	lactation	Pre	rbST	Pre	rbST	SEM	rbST	MFC	rbSTxMFC
Mammary plasma flow (mL min <sup>-1</sup> )									
	Early	3748.00	4030.00	3923.00	5024.00	186.00	0.006	0.561	0.060
	Mid	3139.00	3871.00	3164.00	4141.00	303.00	0.023	0.822	0.696
	Late	2817.00	3843.00	3389.00	3792.00	185.00	0.005	0.676	0.131
Plasma glucose (µmol mL <sup>-1</sup> )									
	Early	3.73	3.51	3.64	3.48	0.10	0.098	0.883	0.763
	Mid	3.55	3.40	3.52	3.67	0.10	0.992	0.719	0.159
	Late	3.49	3.52	3.82	3.77	0.09	0.918	0.286	0.646
A-V (µmol mL <sup>-1</sup> )									
4	Early	0.66	0.67	0.76	0.61	0.08	0.485	0.858	0.261
	Mid	0.62	0.58	0.74	0.72	0.07	0.810	0.480	0.605
	Late	0.78	0.86	0.81	0.80	0.08	0.552	0.650	0.352
Mammary glucose uptake (µmol min <sup>-1</sup> )									
	Early	2299.00	2651.00	2438.00	2653.00	212.00	0.168	0.766	0.632
	Mid	1879.00	2437.00	1881.00	2745.00	355.00	0.042	0.624	0.982
	Late	2183.00	3235.00	2475.00	2936.00	253.00	0.051	0.530	0.203
Percentage of mammary glucose extraction (	%)								
	Early	16.70	18.60	19.30	16.90	1.58	0.984	0.816	0.164
	Mid	17.10	16.70	19.60	18.80	1.57	0.696	0.461	0.398
	Late	22.20	24.40	21.50	21.50	1.62	0.530	0.901	0.373
					22				

SEM = Standard Error of the Mean;<sup>1</sup> p-values for the effects; MFC = Misty-Fan Cooling effect; rbST = rbST effect; MFC x rbST = Interaction effect of MFC and rbST

		NS		MFC			<sup>1</sup> Effect		
Parameter lact	iges of tation	Pre	rbST	Pre	rbST	SEM	rbST	MFC	rbSTxMFC
Glucose turnover									
[U- <sup>14</sup> C] glucose (µmol min <sup>-1</sup> )									
Ear	rly	3377.4	3974.2	4547.8	4029.8	432.70	0.930	0.236	0.234
Mi	d	4380.8	4388.2	5851.6	5144.4	510.40	0.512	0.135	0.504
Lat	te	4000.2	4302.8	5426.6	5428.8	414.90	0.723	0.110	0.727
[3- <sup>3</sup> H] glucose (µmol min <sup>-1</sup> )									
Ear	rly	4631.0	5064.6	5252.2	5032.4	539.80	0.848	0.703	0.562
Mie	d	5493.4	5488.8	7926.6	6026.4	598.10	0.150	0.134	0.152
Lat	te	5309.2	5824.2	6707.0	8188.2	973.20	0.335	0.199	0.633
Glucose-C-recycling (%)									
Ear	rly	24.9	22.1	21.0	19.3	5.71	0.698	0.347	0.927
Mie	d	19.5	20.4	26.2	15.9	2.83	0.140	0.696	0.082
Lat	te	28.5	24.2	18.6	30.1	4.58	0.453	0.776	0.126
Plasma glucose clearance (mL min <sup>-1</sup> )									
Ear	rly	1403.9	1614.3	1391.9	1434.2	163.20	0.461	0.681	0.620
Mie	d	1603.0	1588.3	2357.3	1643.3	182.50	0.081	0.192	0.092
Lat	te	1437.4	1737.8	1845.2	1881.2	283.40	0.252	0.264	0.866
Non mammary glucose utilization (umol	min <sup>-1</sup> )								
Ear	rlv	2331.9	2413.6	2814.6	2379.1	659.40	0.754	0.898	0.746
Mi	ď	3614.1	3052.4	4965.4	3281.3	406.70	0.014	0.207	0.336
Lat	te	3126.2	2589.7	4231.6	3929.6	636.40	0.713	0.228	0.793
Non mammary glucose utilization (%)									
Ear	rlv	49.6	47.8	52.5	44.2	7.00	0.395	0.754	0.787
Mi	d	65.4	59.1	74.3	53.8	3.32	0.001	0.903	0.207
Lat	te	58.6	44.6	58.9	55.9	4.49	0.223	0.538	0.301
Body weight (kg)									
Ear	rlv	358.8	380.8	363.8	380.2	6.54	0.019	0.908	0.680
Mi	d	378.8	386.8	381.8	411.4	3.67	0.001	0.586	0.019
Lat	te	391.0	400.2	418.6	427.4	6.16	0.182	0.297	0.975

Table 6: Glucose turnover rate, glucose-C-recycling, plasma glucose clearance, non-mammary glucose utilization and body weight during rbST administration at different stages of lactation of Holstein cows housing in Normal Shade (NS) and shade plus Misty-Fan Cooling (MFC)

SEM = Standard Error of the Mean;  $^{1}$  p-values for the effects; MFC = Misty-Fan Cooling effect, rbST = rbST effect; MFC x rbST = Interaction effect of MFC and rbST

The glucose entry and utilization rates were not affected by rbST in both cooled and non-cooled cows. The recycling of glucose-C showed no differences between cooled and non-cooled cows whether supplemental rbST or not. Plasma glucose clearance remained unchanged during rbST administration in both cooled and non-cooled cows. Both absolute values and percentage of utilization of glucose of non-mammary tissues of both cooled and non-cooled cows increased as lactation advanced, but it was significantly decreased by rbST during mid lactation. The body weights of both cooled and non-cooled cows whether supplemental rbST or not increased stepwise as lactation advances.

Utilization of glucose carbon in the mammary gland: Glucose uptake and incorporation into related products of lactose, citrate and triacylglycerol are shown in Table 7. Absolute values of the utilization of glucose carbon to milk lactose were increased by rbST in early and mid lactation of both cooled and noncooled cows, while it decreased in late lactation. However, the percentage of utilization of glucose carbon for synthesis of milk lactose was not affected by rbST in early and mid lactation, but the significant decrease was apparent in late lactation of both cooled and non-cooled cows. The absolute values and percentage of utilization of glucose carbon for synthesis of milk citrate were significantly lower in rbST-treated cows during mid and late lactation in both cooled and non-cooled cows. During supplementation of rbST, the utilizations of glucose carbon for synthesis of milk triacylglycerol were higher in both cooled and noncooled cows in all stages of lactation.

Glucose metabolisms in different metabolic pathways in the udder: The effects of supplemental rbST and cooling on intracellular glucose metabolisms in the mammary gland are shown in Table 8. The incorporation of <sup>3</sup>H from [3-<sup>3</sup>H] glucose into fatty acids and the flux through the pentose phosphate pathway were increased by supplemental rbST in both cooled and non-cooled cows. These flux were also increased because of lactation advances neither cooling system nor rbST. Correction for the lower <sup>3</sup>H/<sup>14</sup>C ratio in presenting intracellular glucose 6-phosphate still gave high flux values as lactation advances and during supplemental rbST in both cooled and non-cooled cows. The present results for the net metabolism of glucose 6-phosphate via the pentose phosphate pathway has been defined as glucose 6-phosphate metabolized according to the equation (Katz and Wood, 1963):

glucose 6-phosphate g lyceraldehyde 3-phosphate + 3CO<sub>2</sub>

Table 7: Utilization of glucose carbon in the udder during rbST administration at different stages of lactation of Holstein cows housing in Normal Shade (NS) and shade plus Misty-Fan Cooling (MFC)

		NS		MFC			<sup>1</sup> Effect		
	Stages of								
Parameter	lactation	Pre	rbST	Pre	rbST	SEM	rbST	MFC	rbSTxMFC
[ <sup>14</sup> C] Glucose incorporati	ion (µmol min <sup>-1</sup> ) iı	nto:							
Milk lactose									
	Early	1102.90	1633.50	1809.70	2372.50	254.96	0.009	0.323	0.601
	Mid	1280.20	1405.70	1738.70	2113.40	237.83	0.856	0.602	0.452
	Late	1369.70	1034.00	1661.60	874.10	223.11	0.059	0.652	0.675
Milk triacylglycerol									
	Early	78.18	135.81	165.25	236.94	38.69	0.163	0.010	0.760
	Mid	126.85	217.62	197.00	197.66	29.82	0.244	0.660	0.251
	Late	154.49	205.94	118.87	231.53	51.25	0.226	0.930	0.638
Milk citrate									
	Early	25.45	21.19	23.78	16.41	4.58	0.793	0.822	0.597
	Mid	25.06	17.20	16.50	8.84	3.67	0.013	0.136	0.922
	Late	25.43	16.67	20.81	18.13	2.59	0.052	0.704	0.024
Percentage of glucose can	rbon appearing as:	:							
Milk lactose									
	Early	52.50	64.60	73.60	88.60	34.93	0.261	0.080	0.718
	Mid	81.30	58.50	91.90	77.90	19.45	0.628	0.377	0.831
	Late	58.60	32.40	57.80	28.30	11.34	0.012	0.199	0.429
Milk triacylglycerol									
	Early	3.70	7.60	8.20	11.40	1.98	0.227	0.011	0.577
	Mid	6.90	10.30	11.60	7.60	1.74	0.424	0.354	0.649
	Late	6.70	6.40	4.90	8.40	2.28	0.632	0.655	0.590
Milk citrate									
	Early	1.46	1.09	0.98	0.750	0.38	0.556	0.714	0.853
	Mid	1.56	0.91	1.07	0.380	0.22	0.005	0.135	0.468
	Late	1.19	0.58	0.70	0.620	0.12	0.013	0.385	0.017

 $\overline{SEM} = Standard Error of the Mean; ^1p-values for the effects; MFC = Misty-Fan Cooling effect; rbST = rbST effect; MFC x rbST = Interaction effect of MFC and rbST$ 

 Table 8: Glucose metabolism in different metabolic pathway in the udder during rbST administration at different stages of lactation of Holstein cows housing in Normal Shade (NS) and shade plus Misty-Fan Cooling (MFC)

Stages of parameterStages of lactationPre rbSTPre PrerbSTSEM rbSTMFCMFCMFKFlux through the pentose phosphate pathway calculated as "H incorporation into milk fatty acid			NS		MFC			<sup>1</sup> Effect		
Parameter         Inclution         Pre         rbS1         Pre         rbS1         SEM         rbS1         MFC         rbS1xMPC           Flux through the pentose phabep attributed as "H incorporation into milk fatty acid (equivalent µmol of glucose min")	<b>D</b>	Stages of		1.077			(F) (			
Flux through the pentose phosphate pathway calculated as "4"           ***********************************	Parameter	lactation	Pre	rbST	Pre	rbST	SEM	rbST	MFC	rbSTxMFC
"H incorporation into milk fatty acid (equivalent µmol of glucose min <sup>-1</sup> ) Early 156.50 234.70 236.70 326.90 34.08 0.039 0.269 0.865 Mid 212.70 294.20 300.00 344.10 64.36 0.358 0.250 0.779 Late 421.30 412.90 376.00 282.90 68.33 0.479 0.415 0.552 Corrected <sup>3</sup> H incorporation into milk fatty acid (equivalent µmol of glucose min <sup>-1</sup> ) Early 237.27 280.81 273.54 420.42 65.64 0.185 0.406 0.454 Mid 280.22 352.92 406.02 551.58 77.09 0.195 0.096 0.649 Late 537.57 572.53 472.91 420.03 103.24 0.933 0.501 0.682 Net metabolism of glucose 6-phosphate via the pentose phosphate pathway (µmol min <sup>-1</sup> ) Early 70.10 102.68 97.13 143.49 18.04 0.060 0.336 0.713 Mid 94.30 131.29 126.93 134.54 34.27 0.534 0.586 0.680 Mid 94.30 131.29 126.53 134.54 34.27 0.534 0.385 0.622 Net metabolism of glucose 6-phosphate via the pentose phosphate pathway (µmol min <sup>-1</sup> ) Early 11.00 13.20 11.40 16.80 4.04 0.374 0.778 0.699 Mid 13.10 14.20 14.50 14.90 4.24 0.544 0.374 0.778 0.699 Mid 13.10 14.20 16.50 4.709 6.20 2.00 0.195 0.409 0.952 Metabolism of glucose 6-phosphate via the pentose fore (%) Early 81.60 90.10 6.77.0 74.90 10.10 0.459 0.317 0.951 Mid 87.10 87.40 97.20 78.30 16.30 0.583 0.972 0.571 Mid 87.10 87.40 97.20 78.30 16.30 0.583 0.972 0.571 Early -115.80 -103.40 -91.50 -216.60 111.40 0.977 0.748 0.892 Mid -216.60 124.40 43.00 -25.60 114.0 0.974 0.748 0.892 Mid -216.60 124.0 43.00 -25.60 114.0	Flux through the pentose ph	10sphate pathway ca	alculated as							
	'H incorporation into milk	fatty acid								
Early         156.50         234.70         236.70         326.90         34.08         0.039         0.269         0.865           Mid         212.70         294.20         300.00         344.10         64.36         0.358         0.250         0.779           Late         421.30         412.90         376.00         282.90         68.33         0.479         0.415         0.552           Corrected <sup>3</sup> H incorporation into milk fatty acid           572.53         472.91         420.42         65.64         0.185         0.406         0.454           Mid         280.22         352.92         406.02         551.58         77.09         0.195         0.096         0.649           Late         537.57         572.53         472.91         420.03         0.324         0.933         0.501         0.682           Net metabolism of glucose 6-phosphate via the             0.561         0.680         0.536         0.713           Mid         94.30         131.29         126.93         134.54         34.27         0.534         0.586         0.680           Late         18.80         170.62         155.0         14.90         4.24<	(equivalent µmol of glucose	min <sup>-1</sup> )								
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $		Early	156.50	234.70	236.70	326.90	34.08	0.039	0.269	0.865
Late         421.30         412.90         376.00         282.90         68.33         0.479         0.415         0.552           Corrected <sup>3</sup> H incorporation into milk fatty acid         Early         237.27         280.81         273.54         420.42         65.64         0.185         0.406         0.454           Mid         280.22         352.92         406.02         551.58         77.09         0.195         0.096         0.669           Late         537.57         572.53         472.91         420.03         103.24         0.933         0.501         0.682           Net metabolism of glucose 6-phosphate via the         Early         70.10         102.68         97.13         143.49         18.04         0.060         0.336         0.713           Mid         94.30         131.29         126.93         134.54         34.27         0.534         0.355         0.622           Net metabolism of glucose 6-phosphate via the pentose phosphate pathway (%)         Early         11.00         13.20         11.40         16.80         4.04         0.374         0.778         0.699           Mid         13.10         14.20         16.50         14.90         4.24         0.544         0.347         0.406		Mid	212.70	294.20	300.00	344.10	64.36	0.358	0.250	0.779
Corrected <sup>4</sup> H incorporation into milk fatty acid (equivalent µmol of glucose min <sup>-1</sup> ) Early 237.27 280.81 273.54 420.42 65.64 0.185 0.406 0.454 Mid 280.22 352.92 406.02 551.58 77.09 0.195 0.096 0.649 Late 537.57 572.53 472.91 420.03 103.24 0.933 0.501 0.682 Net metabolism of glucose 6-phosphate via tbe pentose phosphate pathway (µmol min <sup>-1</sup> ) Early 70.10 10.26.8 97.13 143.49 18.04 0.060 0.336 0.713 Mid 94.30 131.29 126.93 134.54 34.27 0.534 0.586 0.680 Late 181.80 170.62 155.28 106.89 36.28 0.435 0.355 0.622 Net metabolism of glucose 6-phosphate via tbe pentose phosphate pathway (%) Early 11.00 13.20 11.40 16.80 4.04 0.374 0.778 0.699 Mid 13.10 14.20 16.50 14.90 4.24 0.544 0.347 0.406 Late 12.50 8.70 9.70 6.20 2.60 0.195 0.409 0.952 Metabolism of glucose 6-phosphate via tbe gentose two perturbed with the glucose (%) Early 11.00 87.40 97.20 78.30 16.30 0.583 0.972 0.571 Late 12.50 8.70 9.70 6.20 2.60 0.195 0.409 0.952 Metabolism of glucose 6-phosphate via the glucose (%) Mid 87.10 87.40 97.20 78.30 16.30 0.583 0.972 0.571 Late 56.00 42.70 57.0 74.90 10.10 0.459 0.317 0.951 Mid 87.10 87.40 97.20 78.30 16.30 0.583 0.972 0.571 Late 12.40 91.00 91.10 -345.90 -226.60 168.20 0.243 0.351 0.597 Metabolism of glucose 6-phosphate via Embertway (µmol/min) Early -115.80 -103.40 -197.50 -216.40 111.40 0.977 0.748 0.892 Mid -213.60 91.10 -345.90 -226.60 168.20 0.243 0.351 0.597 Metabolism of glucose 6-phosphate via Embden-Weverfor pathway (%) Early -115.80 -103.40 -197.50 -216.40 111.40 0.971 0.748 0.892 Mid -213.60 91.10 -345.90 -226.60 168.20 0.243 0.351 0.597 Metabolism of glucose 6-phosphate via Embden-Weverfor pathway (%) Early -22.50 -42.10 -43.00 -25.90 11.40 0.914 0.949 0.146 Mid -23.60 -12.40 -61.60 2.500 13.00 4.10 0.555 0.582 0.050		Late	421.30	412.90	376.00	282.90	68.33	0.479	0.415	0.552
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Corrected <sup>3</sup> H incorporation	into milk fatty acid								
Early       237.27       280.81       273.54       420.42       65.64       0.185       0.406       0.454         Mid       280.22       352.92       406.02       551.58       77.09       0.195       0.096       0.649         Late       537.57       572.53       472.91       420.03       103.24       0.933       0.501       0.682         Net metabolism of glucose 6-phosphate via the	(equivalent µmol of glucose	min <sup>-1</sup> )								
Mid       280.22       352.92       406.02       551.58       77.09       0.195       0.096       0.649         Late       537.57       572.53       472.91       420.03       103.24       0.933       0.501       0.682         Net metabolism of glucose 6-phosphate via the pentose phosphate pathway (µmol min <sup>-1</sup> )              Early       70.10       102.68       97.13       143.49       18.04       0.060       0.336       0.713         Mid       94.30       131.29       126.93       134.54       34.27       0.534       0.355       0.622         Net metabolism of glucose 6-phosphate via the pentose phosphate pathway (%)               Early       11.00       13.20       11.40       16.80       4.04       0.374       0.778       0.699         Mid       13.10       14.20       16.50       14.90       2.260       8.70       9.70       2.60       0.419       0.374       0.778       0.699         Mid       13.10       14.20       16.50       14.90       2.260       2.60       0.195       0.409       0.952         Metabolism of glucose 6-phosphate via the gala		Early	237.27	280.81	273.54	420.42	65.64	0.185	0.406	0.454
Late         537.57         572.53         472.91         420.03         103.24         0.933         0.501         0.682           Net metabolism of glucose 6-phosphate via the pentose phosphate pathway (µmol min <sup>-1</sup> )         572.53         472.91         420.03         103.24         0.933         0.501         0.682           Mid         94.30         131.29         126.93         134.54         34.27         0.534         0.586         0.680           Late         181.80         170.62         155.28         106.89         36.28         0.335         0.622           Net metabolism of glucose 6-phosphate via the pentose phosphate pathway (%)         U         U         U         U         U           Early         11.00         13.20         11.40         16.80         4.04         0.374         0.778         0.699           Mid         13.10         14.20         16.50         14.90         4.24         0.544         0.347         0.406           Late         12.50         8.70         9.70         6.20         2.60         0.195         0.91           Mid         87.10         87.40         97.20         78.30         16.30         0.583         0.972         0.571		Mid	280.22	352.92	406.02	551.58	77.09	0.195	0.096	0.649
Net metabolism of glucose 6-phosphate via the pentose phosphate pathway (µmol mi <sup>-1</sup> )       97.13       143.49       18.04       0.060       0.336       0.713         Mid       94.30       131.29       126.93       134.54       34.27       0.534       0.586       0.680         Late       181.80       170.62       155.28       106.89       36.28       0.435       0.355       0.622         Net metabolism of glucose 6-phosphate via the pentose phosphate pathway (%)       11.40       16.80       4.04       0.374       0.406         Late       12.50       8.70       9.70       6.20       2.60       0.195       0.409       0.952         Mid       13.10       14.20       16.50       14.90       4.24       0.544       0.347       0.406         Late       12.50       8.70       9.70       6.20       2.60       0.195       0.409       0.952         Mid       81.60       90.10       67.70       74.90       10.10       0.459       0.317       0.951         Late       12.50       87.40       97.20       78.30       16.30       0.583       0.972       0.571         Late       56.00       42.70       58.90       58.30       11.20<		Late	537.57	572.53	472.91	420.03	103.24	0.933	0.501	0.682
pentose phosphate pathway (µmol min <sup>-1</sup> )       Early       70.10       102.68       97.13       143.49       18.04       0.060       0.336       0.713         Mid       94.30       131.29       126.93       134.54       34.27       0.534       0.586       0.680         Late       18.180       170.62       155.28       106.89       36.28       0.435       0.586       0.620         Net metabolism of glucose 6-phosphate via the pentose phosphate pathway (%)         Early       11.00       13.20       11.40       16.80       4.04       0.374       0.778       0.699         Mid       13.10       14.20       16.50       14.90       4.24       0.544       0.347       0.406         Late       12.50       8.70       9.70       6.20       2.60       0.195       0.409       0.921         Midi       13.10       14.20       16.50       14.90       4.24       0.544       0.347       0.406         Late       12.50       8.70       9.70       6.20       2.60       0.317       0.951         Mid       13.10       143.49       143.49       16.30       0.583       0.972       0.571	Net metabolism of glucose 6	-phosphate via the								
Early         70.10         102.68         97.13         143.49         18.04         0.060         0.336         0.713           Mid         94.30         131.29         126.93         134.54         34.27         0.534         0.586         0.680           Late         181.80         170.62         155.28         106.89         36.28         0.435         0.355         0.629           Net metabolism of glucose 6-phosphate via the pentose phosphate pathway (%)         Early         11.00         13.20         11.40         16.80         4.04         0.374         0.778         0.699           Mid         13.10         14.20         16.50         14.90         4.24         0.544         0.347         0.406           Late         12.50         8.70         9.70         6.20         2.60         0.195         0.409         0.952           Metabolism of glucose 6-phosphate via the glactose moiety of lactose (%)         Early         81.60         90.10         67.70         74.90         10.10         0.459         0.317         0.951           Mid         87.10         87.40         97.20         78.30         16.30         0.583         0.972         0.571           Late         56.00	pentose phosphate pathway	(µmol min <sup>-1</sup> )								
Mid         94.30         131.29         126.93         134.54         34.27         0.534         0.586         0.680           Late         181.80         170.62         155.28         106.89         36.28         0.435         0.355         0.622           Net metabolism of glucose 6-phosphate via the pentose t		Early	70.10	102.68	97.13	143.49	18.04	0.060	0.336	0.713
Late         181.80         170.62         155.28         106.89         36.28         0.435         0.355         0.622           Net metabolism of glucose 6-phosphate via the pentose phosphate pathway (%)                Early         11.00         13.20         11.40         16.80         4.04         0.374         0.778         0.699           Mid         13.10         14.20         16.50         14.90         4.24         0.544         0.347         0.406           Late         12.50         8.70         9.70         6.20         2.60         0.195         0.409         0.952           Metabolism of glucose 6-phosphate via the galactose moiety of lactose (%) <td></td> <td>Mid</td> <td>94.30</td> <td>131.29</td> <td>126.93</td> <td>134.54</td> <td>34.27</td> <td>0.534</td> <td>0.586</td> <td>0.680</td>		Mid	94.30	131.29	126.93	134.54	34.27	0.534	0.586	0.680
Net metabolism of glucose 6-phosphate via the pentose phosphate pathway (%)           Early         11.00         13.20         11.40         16.80         4.04         0.374         0.778         0.699           Mid         13.10         14.20         16.50         14.90         4.24         0.544         0.347         0.406           Late         12.50         8.70         9.70         6.20         2.60         0.195         0.409         0.952           Metabolism of glucose 6-phosphate via the galactose moiety of lactose (%)           Early         81.60         90.10         67.70         74.90         10.10         0.459         0.317         0.951           Mid         87.10         87.40         97.20         78.30         16.30         0.583         0.972         0.571           Late         56.00         42.70         58.90         58.30         11.20         0.553         0.515         0.581           Metabolism of glucose 6-phosphate via Embden-Meyrehof pathway (µmol/min)         Early         -115.80         -103.40         -197.50         -216.40         111.40         0.977         0.748         0.892           Mid         -213.60         91.10         -345.90         -226.60		Late	181.80	170.62	155.28	106.89	36.28	0.435	0.355	0.622
Early         11.00         13.20         11.40         16.80         4.04         0.374         0.778         0.699           Mid         13.10         14.20         16.50         14.90         4.24         0.544         0.347         0.406           Late         12.50         8.70         9.70         6.20         2.60         0.195         0.409         0.952           Metabolism of glucose 6-phosphate via the galactose moiety of lactose (%)         Early         81.60         90.10         67.70         74.90         10.10         0.459         0.317         0.951           Mid         87.10         87.40         97.20         78.30         16.30         0.553         0.515         0.588           Metabolism of glucose 6-phosphate via Embden-Meyenhop pathway (µmol/min)         Early         -115.80         -103.40         -197.50         -216.40         111.40         0.977         0.748         0.892           Mid         -213.60         91.10         -345.90         -226.60         168.20         0.243         0.351         0.597           Late         124.30         465.70         321.3         111.60         139.30         0.649         0.713         0.083           Metabolism of glucose 6-phos	Net metabolism of glucose 6	-phosphate via the	pentose phosp	ohate pathway (	%)					
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$		Early	11.00	13.20	11.40	16.80	4.04	0.374	0.778	0.699
Late         12.50         8.70         9.70         6.20         2.60         0.195         0.409         0.952           Metabolism of glucose 6-phosphate via the galactose moiety of lactose (%)                Early         81.60         90.10         67.70         74.90         10.10         0.459         0.317         0.951           Mid         87.10         87.40         97.20         78.30         16.30         0.583         0.972         0.571           Late         56.00         42.70         58.90         58.30         11.20         0.553         0.515         0.588           Metabolism of glucose 6-phosphate via Embden-Meyerhof pathway (µmol/min)                  0.977         0.748         0.892           Mid         -213.60         91.10         -345.90         -226.60         168.20         0.243         0.351         0.597           Late         124.30         465.70         321.3         111.60         139.30         0.649         0.713         0.083           Metabolism of glucose 6-phosphate via Embden-Meyerhof pathway (%)           22.50 <td></td> <td>Mid</td> <td>13.10</td> <td>14.20</td> <td>16.50</td> <td>14.90</td> <td>4.24</td> <td>0.544</td> <td>0.347</td> <td>0.406</td>		Mid	13.10	14.20	16.50	14.90	4.24	0.544	0.347	0.406
Metabolism of glucose 6-phosphate via the galactose moiety of lactose (%)           Early         81.60         90.10         67.70         74.90         10.10         0.459         0.317         0.951           Mid         87.10         87.40         97.20         78.30         16.30         0.583         0.972         0.571           Late         56.00         42.70         58.90         58.30         11.20         0.553         0.515         0.588           Metabolism of glucose 6-phosphate via Embder-Meyerhof pathway (µmol/min)         Early         -115.80         -103.40         -197.50         -216.40         111.40         0.977         0.748         0.892           Mid         -213.60         91.10         -345.90         -226.60         168.20         0.243         0.351         0.597           Late         124.30         465.70         321.3         111.60         139.30         0.649         0.713         0.083           Metabolism of glucose 6-phosphate via Embder-Meyerhor pathway (%)         Early         -22.50         -42.10         -43.00         -25.90         11.40         0.914         0.949         0.146           Mid         -25.60         -12.40         -61.10         -61.40         21.60		Late	12.50	8.70	9.70	6.20	2.60	0.195	0.409	0.952
Early         81.60         90.10         67.70         74.90         10.10         0.459         0.317         0.951           Mid         87.10         87.40         97.20         78.30         16.30         0.583         0.972         0.571           Late         56.00         42.70         58.90         58.30         11.20         0.553         0.515         0.588           Metabolism of glucose 6-phosphate via Embden-Meyerhof pathway (µmol/min)         Early         -115.80         -103.40         -197.50         -216.40         111.40         0.977         0.748         0.892           Mid         -213.60         91.10         -345.90         -226.60         168.20         0.243         0.351         0.597           Late         124.30         465.70         321.3         111.60         139.30         0.649         0.713         0.083           Metabolism of glucose 6-phosphate via Embden-Meyerhof pathway (%)         Early         -22.50         -42.10         -43.00         -25.90         11.40         0.914         0.949         0.146           Mid         -25.60         -12.40         -61.10         -61.40         21.60         0.774         0.370         0.762           Late	Metabolism of glucose 6-ph	osphate via the gala	ctose moiety	of lactose (%)						
Mid         87.10         87.40         97.20         78.30         16.30         0.583         0.972         0.571           Late         56.00         42.70         58.90         58.30         11.20         0.553         0.515         0.588           Metabolism of glucose 6-phosphate via Embden-Meyerhof pathway (µmol/min)         Early         -115.80         -103.40         -197.50         -216.40         111.40         0.977         0.748         0.892           Mid         -213.60         91.10         -345.90         -226.60         168.20         0.243         0.351         0.597           Late         124.30         465.70         321.3         111.60         139.30         0.649         0.713         0.083           Metabolism of glucose 6-phosphate via Embden-Meyerhof pathway (%)         Early         -22.50         -42.10         -43.00         -25.90         11.40         0.914         0.949         0.146           Mid         -25.60         -12.40         -61.10         -61.40         21.60         0.774         0.370         0.762           Late         19.60         26.60         25.00         13.00         4.10         0.565         0.582         0.050		Early	81.60	90.10	67.70	74.90	10.10	0.459	0.317	0.951
Late         56.00         42.70         58.90         58.30         11.20         0.553         0.515         0.588           Metabolism of glucose 6-phosphate via Embden-Meyerhof pathway (µmol/mi) <th< td=""><td></td><td>Mid</td><td>87.10</td><td>87.40</td><td>97.20</td><td>78.30</td><td>16.30</td><td>0.583</td><td>0.972</td><td>0.571</td></th<>		Mid	87.10	87.40	97.20	78.30	16.30	0.583	0.972	0.571
Metabolism of glucose 6-phosphate via Embden-Meyerhof pathway (µmol/min)           Early         -115.80         -103.40         -197.50         -216.40         111.40         0.977         0.748         0.892           Mid         -213.60         91.10         -345.90         -226.60         168.20         0.243         0.351         0.597           Late         124.30         465.70         321.3         111.60         139.30         0.649         0.713         0.083           Metabolism of glucose 6-phosphate via Embden-Meyerhof pathway (%)         Early         -22.50         -42.10         -43.00         -25.90         11.40         0.914         0.949         0.146           Mid         -25.60         -12.40         -61.10         -61.40         21.60         0.774         0.370         0.762           Late         19.60         26.60         25.00         13.00         4.10         0.565         0.582         0.050		Late	56.00	42.70	58.90	58.30	11.20	0.553	0.515	0.588
Early         -115.80         -103.40         -197.50         -216.40         111.40         0.977         0.748         0.892           Mid         -213.60         91.10         -345.90         -226.60         168.20         0.243         0.351         0.597           Late         124.30         465.70         321.3         111.60         139.30         0.649         0.713         0.083           Metabolism of glucose 6-phosphate via Embden-Meyerhof pathway (%)         Early         -22.50         -42.10         -43.00         -25.90         11.40         0.914         0.949         0.146           Mid         -25.60         -12.40         -61.10         -61.40         21.60         0.774         0.370         0.762           Late         19.60         26.60         25.00         13.00         4.10         0.565         0.582         0.050	Metabolism of glucose 6-ph	osphate via Embder	-Meyerhof p	athway (µmol/n	1in)					
Mid         -213.60         91.10         -345.90         -226.60         168.20         0.243         0.351         0.597           Late         124.30         465.70         321.3         111.60         139.30         0.649         0.713         0.083           Metabolism of glucose 6-phosphate via Embden-Meyerhof pathway (%)         Early         -22.50         -42.10         -43.00         -25.90         11.40         0.914         0.949         0.146           Mid         -25.60         -12.40         -61.10         -61.40         21.60         0.774         0.370         0.762           Late         19.60         26.60         25.00         13.00         4.10         0.565         0.582         0.050		Early	-115.80	-103.40	-197.50	-216.40	111.40	0.977	0.748	0.892
Late         124.30         465.70         321.3         111.60         139.30         0.649         0.713         0.083           Metabolism of glucose 6-phosphate via Embden-Meyerhof pathway (%)         -         -         -         -         -         -         -         -         -         -         -         -         -         0.649         0.713         0.083           Metabolism of glucose 6-phosphate via Embden-Meyerhof pathway (%)         Early         -22.50         -42.10         -43.00         -25.90         11.40         0.914         0.949         0.146           Mid         -25.60         -12.40         -61.10         -61.40         21.60         0.774         0.370         0.762           Late         19.60         26.60         25.00         13.00         4.10         0.565         0.582         0.050		Mid	-213.60	91.10	-345.90	-226.60	168.20	0.243	0.351	0.597
Metabolism of glucose 6-phosphate via Embden-Meyerhof pathway (%)         Early         -22.50         -42.10         -43.00         -25.90         11.40         0.914         0.949         0.146           Mid         -25.60         -12.40         -61.10         -61.40         21.60         0.774         0.370         0.762           Late         19.60         26.60         25.00         13.00         4.10         0.565         0.582         0.050		Late	124.30	465.70	321.3	111.60	139.30	0.649	0.713	0.083
Early-22.50-42.10-43.00-25.9011.400.9140.9490.146Mid-25.60-12.40-61.10-61.4021.600.7740.3700.762Late19.6026.6025.0013.004.100.5650.5820.050	Metabolism of glucose 6-ph	osphate via Embder	-Meyerhof p	athway (%)						
Mid-25.60-12.40-61.10-61.4021.600.7740.3700.762Late19.6026.6025.0013.004.100.5650.5820.050		Early	-22.50	-42.10	-43.00	-25.90	11.40	0.914	0.949	0.146
Late 19.60 26.60 25.00 13.00 4.10 0.565 0.582 0.050		Mid	-25.60	-12.40	-61.10	-61.40	21.60	0.774	0.370	0.762
		Late	19.60	26.60	25.00	13.00	4.10	0.565	0.582	0.050

SEM = Standard Error of the Mean; <sup>1</sup>p-values for the effects; MFC = Misty-Fan Cooling effect; rbST = rbST effect; MFCxrbST = Interaction effect of MFC and rbST

Table 9: Fatty acid composition of milk fat during rbST administration at different stages of lactation of Holstein cows housing in Normal Shade (NS) and shade plus Misty-Fan Cooling (MFC)

		NS	,	MFC			<sup>1</sup> Effect		
Stages of lactation	Fatty acid chain length	Pre	rbST	Pre	rbST	SEM	rbST	MFC	rbSTxMFC
Early lactation: (µmol mL <sup>-1</sup>	)								
	C6	0.97	1.30	0.53	1.79	0.20	0.004	0.937	0.051
	C8	0.37	0.68	0.95	0.84	0.11	0.373	0.060	0.080
	C10	0.81	1.40	2.06	1.67	0.19	0.582	0.041	0.030
	C12	0.88	1.47	2.20	1.75	0.15	0.670	0.021	0.009
	C14	4.16	5.59	6.74	6.47	0.45	0.231	0.040	0.094
	C16:0	16.67	21.64	22.38	25.35	1.30	0.016	0.131	0.464
	C16:1	0.17	0.67	0.66	0.82	0.08	0.004	0.065	0.079
	C18:0	5.72	4.92	4.00	5.68	0.40	0.304	0.586	0.014
	C18:1	11.94	12.05	8.91	16.89	1.58	0.037	0.605	0.033
	C18:2 trans	0.67	0.86	0.71	0.69	0.15	0.577	0.728	0.509
	C18:2 cis	1.64	0.91	0.80	0.86	0.38	0.408	0.325	0.327
	Total	45.25	50.26	49.93	62.81	2.93	0.016	0.195	0.217
Mid lactation: (µmol mL <sup>-1</sup> )									
	C6	1.43	1.50	1.49	1.80	0.15	0.240	0.757	0.467
	C8	0.69	0.17	0.69	0.85	0.09	0.314	0.78	0.458
	C10	1.56	1.81	1.42	1.85	0.20	0.121	0.946	0.661
	C12	1.61	1.97	1.51	2.00	0.24	0.108	0.956	0.778
	C14	5.54	6.53	5.40	6.69	0.55	0.074	0.993	0.793
	C16:0	19.69	22.54	19.17	23.45	1.34	0.029	0.963	0.608
	C16:1	0.61	0.77	0.81	1.00	0.07	0.029	0.203	0.854
	C18:0	4.86	5.24	3.39	4.29	0.60	0.314	0.264	0.675
	C18:1	9.07	10.91	10.43	13.77	1.00	0.033	0.048	0.478
	C18:2 trans	0.16	0.14	0.13	0.20	0.02	0.195	0.679	0.016
	C18:2 cis	0.91	1.02	0.58	0.82	0.08	0.083	0.038	0.477
	Total	46.11	53.14	45.04	56.74	2.83	0.011	0.870	0.433
Late lactation: (µmol mL <sup>-1</sup> )									
	C6	1.54	1.69	1.50	1.91	0.11	0.029	0.876	0.255
	C8	0.68	0.66	0.72	0.96	0.05	0.048	0.268	0.027
	C10	1.51	1.47	1.47	2.10	0.08	0.008	0.335	0.004
	C12	1.65	1.72	1.53	2.19	0.07	0.001	0.438	0.003
	C14	6.08	5.89	5.37	6.75	0.33	0.113	0.912	0.047
	C16:0	21.60	23.05	20.56	24.43	1.78	0.174	0.954	0.515
	C16:1	0.84	1.01	0.89	1.10	0.15	0.252	0.749	0.888
	C18:0	5.29	4.82	4.50	6.36	0.67	0.326	0.747	0.118
	C18:1	12.30	14.67	13.37	19.63	1.17	0.006	0.289	0.134
	C18:2 trans	0.21	0.20	0.23	0.27	0.02	0.478	0.371	0.135
	C18:2 cis	1.03	0.95	0.86	1.24	0.12	0.229	0.686	0.081
	Total	53.10	55.76	51.89	66.05	4.03	0.070	0.553	0.191

SEM = Standard Error of the Mean;  $^{1}$ p-values for the effects; MF = Misty-Fan Cooling effect; rbST = rbST effect; MF x rbST = Interaction effect of MF and rbST

According to this equation, complete metabolism of one molecule of glucose 6-phosphate would require three cycles of the pentose phosphate pathway. Therefore, the flux through the pathway would be three times for the net rate of glucose metabolized in the pentose phosphate pathway. The present results showed that the intracellular glucose phosphorylated by the mammary gland were metabolized more via the pentose phosphate pathway by rbST either in terms of absolute values or the percentage values in early and mid lactation. These values were declined in late lactation. Values of metabolism of glucose 6-phosphate via the galactose moiety of lactose were decreased as lactation advanced to late lactation neither cooling nor rbST in both groups. Metabolism of glucose 6-phosphate via the Embden-Meyerhof pathway was calculated in term of the proportion of metabolized glucose, which was considerable variation throughout stages of lactation in

cooled and non-cooled cows. However, the absolute rate of metabolism of glucose via the Embden-Meyerhof pathway appeared to increase in the late lactation in both cooled and non-cooled cows whether supplemental rbST or not, but the inclement was not statistical significant.

**Milk fatty acid concentration:** The data in Table 9 showed the marked increases in the total milk fatty acids concentrations during supplemental rbST in both cooled and non-cooled cows. The statistical significant effects of rbST on total milk fatty acids concentrations were apparent in early and mid lactation in both groups. The long chain length fatty acids ( $C_{16}$ - $C_{18}$ ) concentrations in milk were significantly increased by supplemental rbST in both cooled and non-cooled cows in each stage of lactation.

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		NS		MFC			<sup>1</sup> Effect		
	Stages of								
Parameter	lactation	Pre	rbST	Pre	rbST	SEM	rbST	MFC	rbSTxMFC
Requirement of all NADPH for	Early	1134.0	1761.0	1920.0	2067.0	134.00	0.020	0.203	0.112
fatty acid synthesis (µmol/min)	Mid	1235.0	1698.0	1428.0	2150.0	181.00	0.011	0.539	0.495
	Late	1240.0	1460.0	1154.0	2062.0	150.00	0.006	0.481	0.052
Requirement of all NADPH	Early	16.7	18.5	19.5	24.9	3.14	0.286	0.482	0.594
formation from glucose via the	Mid	25.6	26.5	31.6	25.6	7.36	0.744	0.667	0.651
pentose phosphate pathway (%)	Late	35.4	25.6	26.5	17.8	5.16	0.112	0.255	0.925

Table 10: NADPH production for fatty acid synthesis in the udder during rbST administration at different stages of lactation of Holstein cows housing in Normal Shade (NS) and shade plus Misty-Fan Cooling (MFC)

SEM = Standard Error of the Mean; <sup>1</sup>p-values for the effects; MFC =Misty-Fan Cooling effect;rbST = rbST effect; MFC x rbST = Interaction effect of MFC and rbST

Table 11: <sup>3</sup>H/<sup>14</sup>C ratios in plasma glucose and related products during rbST administration at different stages of lactation of Holstein cows housing in Normal Shade (NS) and shade plus Misty-Fan Cooling (MFC)

_		NS		MFC	MFC			<sup>1</sup> Effect			
Parameter	Stage of lactation	Pre	rbST	Pre	rbST	SEM	rbST	MFC	rbSTxMFC		
Plasma glucose											
	Early	0.76	0.81	0.86	0.82	0.08	0.973	0.393	0.589		
	Mid	0.80	0.82	0.74	0.68	0.04	0.653	0.217	0.317		
	Late	0.80	0.76	0.84	0.68	0.04	0.029	0.781	0.176		
Milk lactose											
	Early	0.83	0.73	0.65	0.74	0.06	0.930	0.104	0.173		
	Mid	0.88	0.88	0.73	0.70	0.08	0.831	0.152	0.869		
	Late	0.72	0.60	0.71	0.71	0.08	0.493	0.539	0.471		
Milk galactose											
8	Early	0.86	0.87	0.60	0.66	0.07	0.576	0.047	0.715		
	Mid	0.83	0.93	0.72	0.73	0.11	0.638	0.231	0.677		
	Late	0.64	0.64	0.67	0.74	0.11	0.732	0.472	0.759		
Milk triacylglyce	erol										
	Early	1.43	2.14	2.74	1.64	0.60	0.852	0.176	0.076		
	Mid	3.51	2.93	3.70	1.79	0.72	0.124	0.721	0.381		
	Late	2.45	2.61	3.12	1.90	0.50	0.327	0.976	0.206		
Milk citrate											
	Early	0.86	0.74	0.78	0.81	0.05	0.410	0.964	0.189		
	Mid	0.98	0.86	0.87	0.86	0.06	0.291	0.463	0.367		
	Late	0.81	0.86	0.87	0.78	0.04	0.640	0.899	0.135		
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SEM = Standard Error of the Mean;  $^{1}$ p-values for the effects; MFC = Misty-Fan Cooling effect; rbST = rbST effect; MFC x rbST = Interaction effect of MFC and rbST

**NADPH production from glucose:** The requirement of NADPH for fatty acid synthesis which was calculated from milk fatty compositions and output is shown in Table 10. The NADPH productions for fatty acid synthesis were significantly increased by supplemental rbST in each stage of lactation in both cooled and non-cooled cows. The percentage of NADPH production from glucose via the pentose phosphate pathway was considerable variation throughout stages of lactation in cooled and non-cooled cows.

The 3H/14C ratio in glucose and related products: The  ${}^{3}$ H/ ${}^{14}$ C ratios in plasma glucose and related products are shown in Table 11. The  ${}^{3}$ H/ ${}^{14}$ C ratio in arterial plasma glucose was lower than that of the infusion in both groups. These values were not different among cooled and non-cooled cows supplemental rbST in different stages of lactation, indicating some recycling of glucose-C in the whole animals during periods of study. A further decrease in the <sup>3</sup>H/<sup>14</sup>C ratio was seen in milk lactose. As the glucose moiety of lactose arises directly from plasma glucose, this decrease in the ratio was due to metabolism of glucose 6-phosphate within the udder before incorporation into lactose as galactose. The <sup>3</sup>H/<sup>14</sup>C ratio of milk triacylglycerol was shown to be high whether with or without rbST in both cooled and non-cooled cows, indicating <sup>3</sup>H-glucose was removed and detected in milk triacylglycerol. The <sup>3</sup>H and <sup>14</sup>C from glucose were shown to be incorporated into milk citrate. The<sup>3</sup>H/<sup>14</sup>C ratio of milk citrate was not affected by cooling or rbST supplementation.

# DISCUSSION

In the present study, the values of THI in NS and MF barns in either morning or afternoon, were always higher than critical value (THI 72) for lactating dairy cows housing in both barns (Smith *et al.*, 2006).

Crossbred cows in the present study were therefore always subjected to moderate heat stress throughout experimental periods (i.e., THI = 78.3-85.4). The effect of misters and fans cooling in the present study was not sufficient to completely eliminate heat stress in cows because THI measured under the cooling system remained high. The THI values might not accurately reflect heat stress when using a mister and fan system for evaporative cooling that result in higher humidity but also cause cooling. Although the cooling effect using the misty-fan system was not sufficient to adequately reduce THI in the barn, but there is a beneficial effect as indicated by a lower RR and RT in cooled cows and also higher milk yield throughout lactation. These results support the study of Fike et al. (2002) that housing cows during the day with fans and sprinklers effectively reduced heat stress as indicated by lower body temperature and respiration rate. In the present study in both groups, milk yield was increased by rbST which accompanied with increases in both RT and RR throughout the experimental periods. The observation for an increase in heat production during rbST supplementation agrees with the reports of West et al. (1991) and West (1994) that rbST-treated cows can increase heat production in a hot environment either high or lower milk producing cows.

It is known that dairy cattle adapt to high temperatures with variety of hormonal and metabolic responses, which may involve changes in the process of milk synthesis in the mammary gland. Milk yield initially showed significant increases in early lactation of cooled and non-cooled cows either supplemental rbST or not and it decreased as lactation advances. These findings confirm that an increase in milk yield in response to rbST administration will not be sustained indefinitely (Bauman, 1992) and it is influenced by the stage of lactation (Phipps et al., 1991). The low potential for extended persistency of lactation in rbST treated cows appears similar to that which occurs in higher yielding cows (Chase, 1993). However, it has been reported that the response to bovine somatotropin for whole lactation might be reduced if treatment begins very early in lactation (Burton et al., 1994; Bauman and Vernon, 1993). It is known that milk production requires glucose for synthesis of lactose which is essential for milk secretion and glucose moiety of lactose arises directly from plasma glucose (Ebner and Schanbacher, 1974). An increase in milk yield without an alteration of the plasma glucose concentration during supplemental rbST in both cooled and non-cooled cows indicates that a substantial increase in supply of glucose to the mammary gland would depend mainly on the capacity for transmembrane transport and/or

intracellular metabolism of glucose. An increase in mammary blood flow is a factor for glucose uptake by the mammary gland (Linzell, 1973), which the rate of mammary plasma flow of cows was significantly increased by supplemental rbST. However, an increase in mammary plasma flow coinciding with the high level of endogenouse IGF-I, which was inferred during rbST supplementation (Chaiyabutr et al., 2005), or increase in circulating concentrations of IGF-I during prolonged exposure to a long daily photoperiod (Spicer et al., 2007), would not be a major determinant in the mediation of nutrient delivery and uptake by the mammary gland for increase in milk production throughout lactation. Local changes for biosynthetic capacity within the mammary gland would be a factor in identification of the utilization of substrates in the rate of decline in milk yield with advancing lactation (Chaiyabutr et al., 2005).

Effects of supplemental rbST and cooling on glucose kinetics are shown in Table 5 and 6. The maintenance of the plasma glucose concentrations over a wide range at different stages of lactation in both cooled and non-cooled cows indicates that steady state conditions between the rate of gluconeogenesis and the rate of utilization of glucose existed in the body pool of glucose in both groups. However, it has been reported that the plasma glucose concentration would increase during injection of bovine somatotropin in cows with low milk yield but not in cows with high milk yields (Bines et al., 1980). The reversible turnover rate of [3-<sup>3</sup>H] glucose (the total glucose entry rate) and the irreversible turnover rate of [U-14C] glucose (the utilization rate of glucose) of cooled cows without rbST were slightly higher than those of non-cooled cows in all stages of lactation. It is probably that the turnover rate of glucose correlated positively with a higher milk vield in cool cows. However, both glucose entry and utilization rates were not affected by supplementation of rbST in both cooled and non-cooled cows throughout lactation. It is possible that both cooled and non-cooled cows with or without supplemental rbST were fed TMR diet to satisfy requirements for metabolizable energy and the body weights increased stepwise throughout periods of study. It indicates that both cooled and noncooled cows were in positive energy balance. These findings would not affect the irreversible loss of glucose, which has been shown to increase in cows with negative energy balance (McDowell et al., 1987). The reversible turnover rate of [3-<sup>3</sup>H] glucose represents the total glucose turnover rate as the <sup>3</sup>H is not recycled from products of partial glucose degradation (Katz et al., 1965). Thus, recycling of glucose-C was estimated by simultaneous infusion of [3-<sup>3</sup>H] glucose

and [U-<sup>14</sup>C] glucose in both cooled and non-cooled cows, which was not affected by rbST in each stage of lactation. These findings suggest that a constant level of tricarbon units originally derived from glucose being again reincorporated into glucose.

The utilization of glucose across the mammary gland during supplemental rbST in both cooled and non-cooled cows at different stages of lactation are complex regulatory mechanisms. It would depend both on the partitioning of blood flow between extramammary tissues and local regulation. The present results for the mammary uptake of plasma glucose in both groups were not based on changes in A-V concentration differences and extraction ratio of glucose. An increase in the rate of blood flow to the mammary gland during supplemental rbST in both cooled and non-cooled cows would be a major determinant of the rate of glucose uptake by the mammary gland. In all stages of lactation, the net mammary glucose uptake were increased approximately 8-48% by supplemental rbST in both cooled and non-cooled cows. Glucose extracted by the mammary gland has several possible metabolic fates in mammary epithelial cells that may occur at another level than transmembrane transport (Xiao and Cant, 2003). The glucose uptake by the mammary gland during supplemental rbST and cooling would be rate limiting for the transport of glucose to the mammary cell. The high blood flow to the mammary gland during supplemental rbST would decrease the transit time of glucose, thereby reduction for prolonging the contact time between glucose in blood and glucose transporter in mammary epithelial cell (Chaiyabutr et al., 2007).

It is known that glucose is an important intermediary of metabolism for the biosynthesis of lactose, triacylglycerol and citrate by the mammary gland. The bovine mammary gland cannot synthesize its own glucose because of lacking of glucose-6phosphatase (Scott et al., 1976). Glucose plays a crucial role in their metabolism and lactose synthesis, which is formed in Golgi vesicles from a combination of glucose either directly or after phosphorylation to glucose 6phosphate and conversion to UDP-galactose (Ebner and Schanbacher, 1974). The calculated amount of metabolism of glucose 6-phosphate to the galactose moiety of lactose during supplemental rbST in both cooled and non-cooled cows in each stage of lactation would be sufficient to account for the cytosolic lactose synthesis. The utilization of glucose carbon incorporation to lactose in the udder were increased by supplemental rbST in early and mid lactation but not for late lactation in both cooled and non-cooled cows (Table 7). The decrease in the metabolism of glucose 6phosphate to the galactose moiety of lactose as lactation advanced to late lactation in both cooled and noncooled cows would affect to the lactose synthesis and milk production. A low enzymatic activity for lactose synthesis might be expected to appear as lactation advances. According to Davis and Bauman (1974), 50-60% of the glucose in the glucose-6-phosphate pool is converted into galactose. Major part of the galactose has been shown to derive from mammary extracted glucose, as well as from glycerol and other metabolic pathways. However, glucose is not the sole carbon source for lactose synthesis but remains the main one. An increase in the glucose concentration in milk representing an increase in glucose concentration in the mammary epithelial cell during prolonged treatment of rbST has been noted (Chaiyabutr et al., 2008c).

It is known that 80-85% of lactose carbon atoms arise from glucose (Faulkner and Peaker, 1987). The quantitative utilization of the glucose taken up by the mammary gland is used directly in the synthesis of lactose, while the remaining of extracted glucose can participate in the supply of ATP (Embden-Meyerhof pathway and the tricarboxylic acid cycle), other portions would be metabolized via the pentose phosphate pathway, In the present studies, glucose 6phosphate was metabolized via the pentose phosphate pathway by average 11.0-16.5% in both cooled and non-cooled cows without rbST, while it was increased by supplemental rbST from 13.2-16.8% in early and mid-lactation but these values were decreased in the late lactation (Table 8). These results agree with prolonged treatment of rbST in crossbred HF cows, which percentage of glucose 6-phosphate metabolized via the pentose phosphate pathway were variable in different stages of lactation (Chaiyabutr et al., 2008b). However, these findings differed to those studies in the isolated perfused udder of cow by Wood et al. (1965), in which about 23- 30% of the glucose was metabolized via the pentose phosphate pathway. It is probable that no consideration of the recycling of glucose 6phosphate metabolized via the pentose cycle in the udder with the consequent loss of <sup>3</sup>H from glucose 6phosphate (Davis and Bauman, 1974). However, the net proportion of the metabolism of glucose 6-phosphate via the pentose cycle pathway was increased by supplemental rbST at early stage of lactation of cooled and non-cooled cows. Metabolism of glucose via the pentose phosphate pathway yields 2 molecules of NADPH per molecule of glucose, only one of which could be labelled with <sup>3</sup>H in the present experiments. In the present studies, estimation of the contribution of the pentose phosphate pathway in providing NADPH for fatty acid synthesis in vivo have been estimated by

based on the assumption that all the glucose that was oxidized to  $CO_2$ , which was metabolized via the pentose phosphate pathway. High metabolism of glucose 6-phosphate in early and mid lactation of rbST treated cows appeared to be due primarily to a high flux through the lactose synthesis and to pentose phosphate pathway, probably reflecting the high milk production during rbST supplementation.

The utilization of glucose carbon by the mammary epithelial cell for the synthesis of milk compositions of lactose, citrate and triacylglycerol (Table 7) show that absolute amount of glucose carbon incorporation to milk lactose were increased by supplemental rbST in early and mid lactation in both cooled and non-cooled cows but it decreased in late lactation. These findings would parallel to its effects on milk yield. It indicates that during supplemental rbST in late lactation, the metabolism of glucose-6-phosphate declines the flux towards the pentose phosphate pathway and in lactose synthesis. In parallel, a higher proportion of glucose-6phosphate would be metabolized via the Embden-Meyerhof pathway and was oxidized in the tricarboxylic acid cycle.

Both the proportion and absolute amount of glucose carbon incorporation to milk triacylglycerol were increased, while glucose carbon incorporation to milk citrate was slightly decreased by supplemental rbST. It is known that, citrate would not be used directly as a carbon source for lipogenesis in ruminant; it does appear to be directly involved in the provision of reducing equivalent (NADPH). These changes can be interpreted in terms of metabolic shifts occurring within the mammary epithelial cell. It might speculate that such changes reflect more flux of the utilization of glucose carbon by the mammary epithelial cell through the synthesis of lactose and milk triacylglycerol during supplemental rbST. In addition to the use of glucose carbon for milk triacylglycerol synthesis, the hydrogen from glucose has shown to be incorporated more into milk fatty acid in early and mid lactation in both cooled and non-cooled cows supplemental rbST (Table 8), although studies in vitro have shown that fatty acid synthesis could occur from the utilization of acetate in the perfused goat udder (Hardwick et al., 1963). It has been known that milk fat is synthesized from fatty acids of both blood lipids and from de novo synthesis within the mammary epithelial cells. However, an increase in milk fat concentration after rbST supplementation was associated with the increased yield of long-chain fatty acids characteristic of plasma free fatty acids and body fat. Significant increases in plasma free fatty acids in rbST-treated cows have been published elsewhere (Chaiyabutr et al., 2007). Thus, the lipolytic activity

would be a function of rbST treatment per se instead of the associated changes in energy balance.

Glucose can also participate in the milk fat formation, by supplying the glycerol (triose phosphate pathway) and the NADPH essential to elongating milk fatty acids (pentose phosphate and isocitrate dehydrogenase pathways). However, very marginally, less than 11% of glucose could supply carbon atoms for the synthesis of milk triacylglycerol in either supplemental rbST or cooling. The method used to estimate NADPH requirement for fatty acids synthesis in the present study is calculated from the incorporation of <sup>3</sup>H from [3-<sup>3</sup>H] glucose in fatty acids, which assume that the NADPH is used exclusively for biosynthesis of fatty acids (Katz et al., 1974). This technique has been used to study the in vitro metabolism of rat mammary and adipose tissue (Katz and Wals, 1970; Katz and Wals, 1972; Katz et al., 1966) and it has also been used for the study of the in vivo metabolism of goat mammary tissue (Chaiyabutr et al., 1980). Data from the present study (Table 10) provide evidences that the requirement of NADPH for fatty acid synthesis de novo in the udder were ranging 16.7-35.4% in cooled and non-cooled cows without rbST and 17.8-26.5% during rbST supplementation. If there is a common pool of glucose 6-phosphate which is available for both lactose synthesis and pentose phosphate metabolism; then the recycling of glucose 6-phosphate within the udder could show a low value for NADPH production from glucose metabolism in the present results.

Metabolism of glucose 6-phosphate via the pentose phosphate pathway usually loss of all <sup>3</sup>H from [3-<sup>3</sup>H] glucose in lactating cows. During lactation, a higher level of <sup>3</sup>H/<sup>14</sup>C ratio in milk triacyglycerol (Table 11) would be due to an increase in disequilibrium of the triose phosphate isomerase reaction occurring in the udder of crossbred animals, which needs to be further investigated. Tritium and carbon-14 in glucose molecule were also shown to be incorporated into milk citrate which provided by averaged 22  $\mu$ mol min<sup>-1</sup> (16.5-0.25.5) in cooled and non-cooled cows without rbST and provided by averaged 16 µmol min<sup>-1</sup> (8.1-21.2) for the carbon skeleton of citrate during rbST supplementation in both groups. Milk citrate could be synthesized from 2-oxoglutarate via the NADPdependent isocitrate dehydrogenase reaction (Hardwick, 1965). In addition <sup>3</sup>H is lost to NADPH or water in metabolism via the pentose phosphate pathway or glycolytic pathway, so it is likely that <sup>3</sup>H incorporation into milk citrate was also via NADP<sup>3</sup>H. It is possible that the incorporation of <sup>3</sup>H into milk citrate may occur in different manners in the exchange reaction of the cytosolic NADP-dependent isocitrate

dehydrogenase. Both fatty acid synthesis and the NADP-dependent isocitrate dehydrogenase reaction may have different mechanisms with a common pool of cytosolic NADPH between cows without rbST and cows supplemental rbST. The concentrations of FFA in milk were significantly increased by supplemental rbST in cooled and non-cooled cows (Table 7). A similar result for an increase in milk fat content due to prolonged administration of rbST has also been observed previously (West *et al.*, 1991; Chaiyabutr *et al.*, 2000b). It has been known that milk fat is synthesized from fatty acids of both blood lipids and from de novo synthesis within the mammary epithelial cells.

### CONCLUSION

The data presented here represent the estimation in vivo of glucose metabolism in the mammary gland and its distribution to lactose synthesis, the pentose phosphate pathway and the Embden-Meyerhof pathway by the effects of supplemental rbST and cooling in 87.5% HF animals. The rbST exerts its galactopoietic action, in part, association with an increase in mammary blood flow, which partitions the distribution of glucose to the mammary gland. The stimulant effect for milk yield by supplemental rbST was transiently and the glucose turnover rate was not significantly increased as compared with pre-treatment period in all stages of lactation. It indicates that rbST induced enhancement of milk yield in all stages of lactation, which would be compensated by mobilization of body energy reserves (i.e. plasma free fatty acids) to the extent of the elevated energy requirements for supporting the increased milk production. In early and mid lactation, the glucose taken up by the udder of both cooled and non-cooled cows with rbST and without supplemental rbST, were metabolized in the pentose phosphate pathway and contributed to NADPH production by mean average 14 and 24%, respectively. An increased flux of the sufficient pool of intracellular glucose 6-phosphate during early and mid lactation came across through the lactose synthesis and pentose cycle pathway. On late lactation of both cooled and non-cooled cows, the reductions of the metabolism of glucose taken up by the udder via the pentose phosphate pathway and the contribution to NADPH production were apparent by mean averaged 9% and 22%, respectively. It would appear that a larger proportion of the glucose 6-phosphate was metabolized via Embden-Meyerhof pathway in late lactation. The present results suggest that the regulation of biosynthetic capacity within the mammary gland would be influenced more by local than by systemic factors in identification of the utilization of substrates in the rate of decline in milk yield with advanced lactation.

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