

# Bulk milk quality traits differ according to the ClassyFarm welfare assessment score

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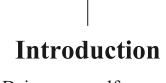








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Dairy cow welfare and ClassyFarm protocol

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Focus on lactose, SCS and TBC

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## 01

## Introduction

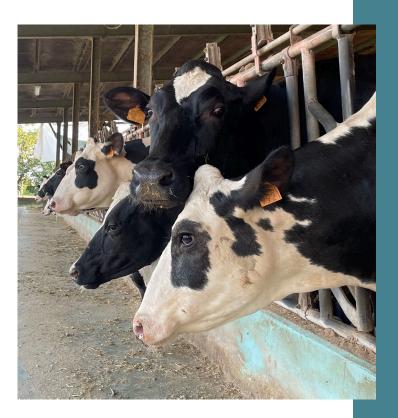
### Dairy cow welfare

Dairy cow welfare can significantly influence productive performance, behaviour, fertility and milk quality.

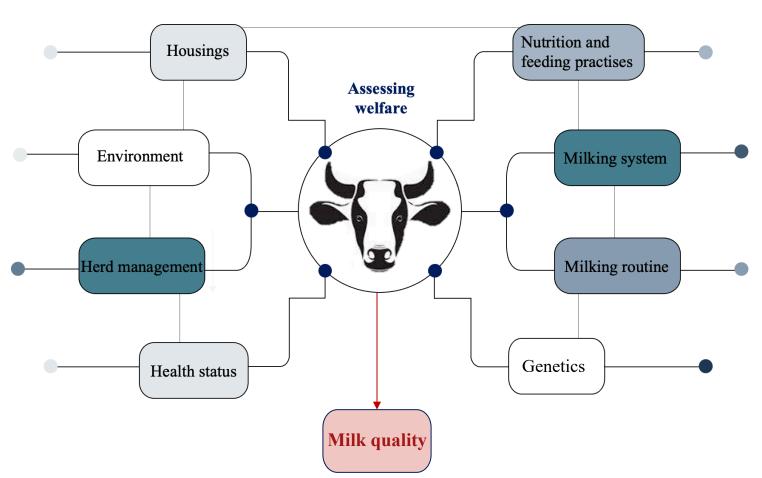
Protocols for evaluating welfare standards have been published and some of them are constantly evolving.



https://www.classyfarm.it/index.php/en/what-en

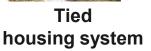


### Dairy cow welfare and milk quality



#### The ClassyFarm protocols







Loose housing system



Biosecurity

Management

Structures

Animal-based measures (ABM)

Major Hazards

15 items (1<sup>st</sup>-15<sup>th</sup>)

25 items (16<sup>th</sup>- 40<sup>th</sup>)

28 items (41<sup>st</sup>- 68<sup>th</sup>)

22 items (69th- 90th)

9 items (91st- 99th)

15 items (1<sup>st</sup>-15<sup>th</sup>)

 $30 \text{ items } (16^{th} - 45^{th})$ 

32 items (46<sup>th</sup>-77<sup>th</sup>)

19 items (78<sup>th</sup>- 96<sup>th</sup>)

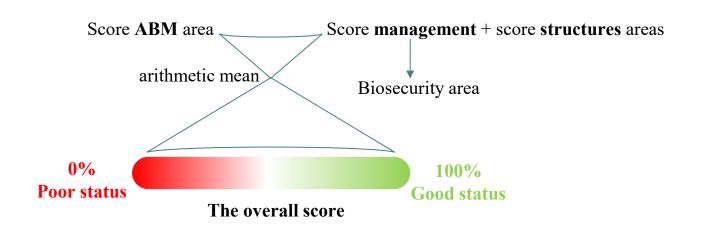
9 items (97<sup>th</sup>- 105<sup>th</sup>)

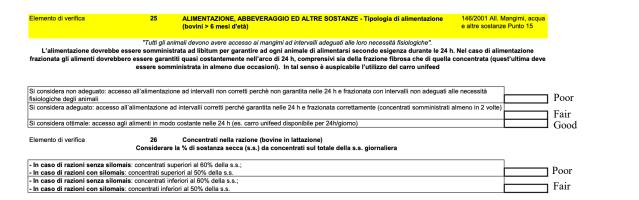
(Bertocchi et al., 2018)

99 items

105 items

#### The ClassyFarm protocols





## AIM OF THE STUDY

Investigation of the effect of the animal welfare level on bulk milk quality traits comparing loose-housing system (LHS) and tied-housing system (THS).

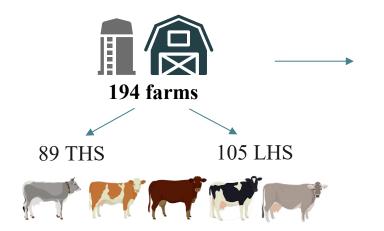
## 02

# Materials and Methods

#### Animal welfare and milk data







#### 194 welfare assessments

The welfare assessment protocols (year 2022) were applied in all the farms by the same Lattebusche veterinarian expert.



#### 6,971 milk tests

Bulk milk samples were collected on average 2.99 times (SD =0.17) per month.

#### Milk quality traits

Bulk tank milk sampled in tubes with H<sub>2</sub>O<sub>2</sub>



- Fat, protein, casein, and **lactose** contents (%);
- urea concentration (mg/dL);
- somatic cell count (SCC, cells/mL)  $\rightarrow$  SCS = 3 + log2(SCC/100,000) (Ali and Shook, 1980).



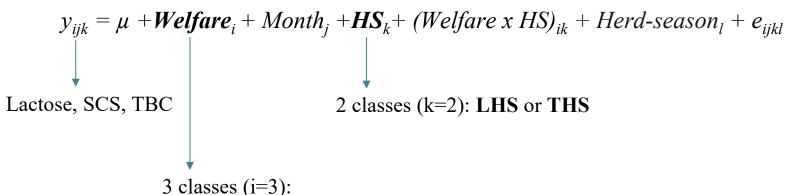
CombiFossTM7 (FOSS Electric A/S, Hillerød, Denmark)

• The **total bacterial count** (TBC, CFU/mL), log10-trasformed.



BactoScanTM FC+ (FOSS Electric, Hillerød, Denmark)

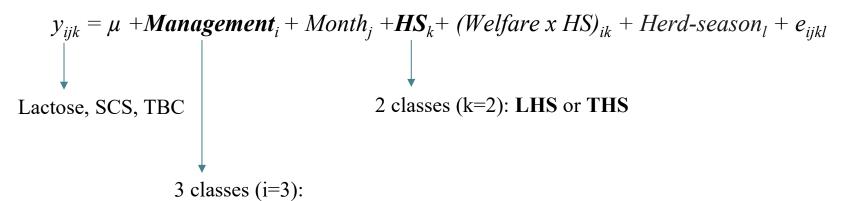
The following mixed linear model was used for variance analysis:



- insufficient (INS, score <65%);
- intermediate (INT, score between 65% and 84%);
- excellent (EXC, score  $\geq 85\%$ ).



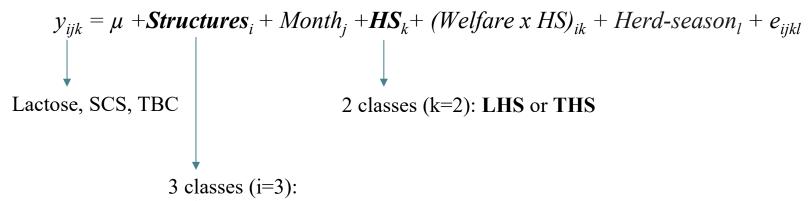
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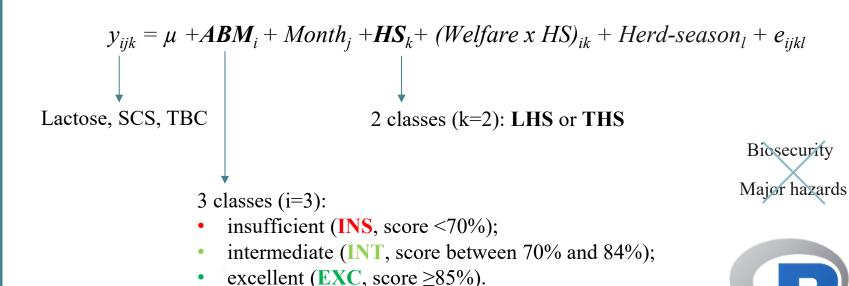
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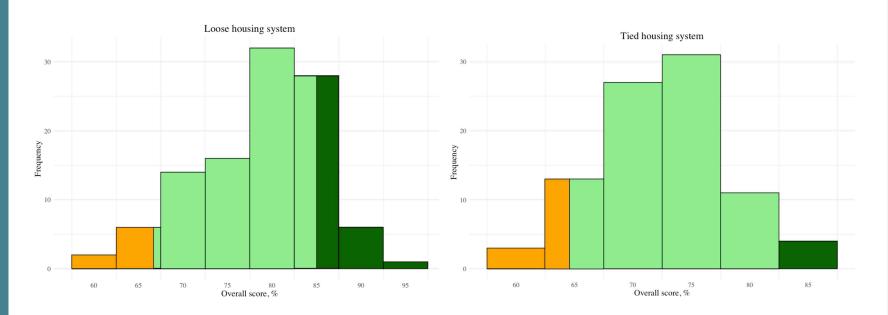


## 03

# Results and Discussion

#### Frequencies distributions

For THS, 90% of farms were classified as intermediate, 4% as excellent, 6% as insufficient, while for LHS, the corresponding percentages were 75%, 20%, and 5%, respectively.



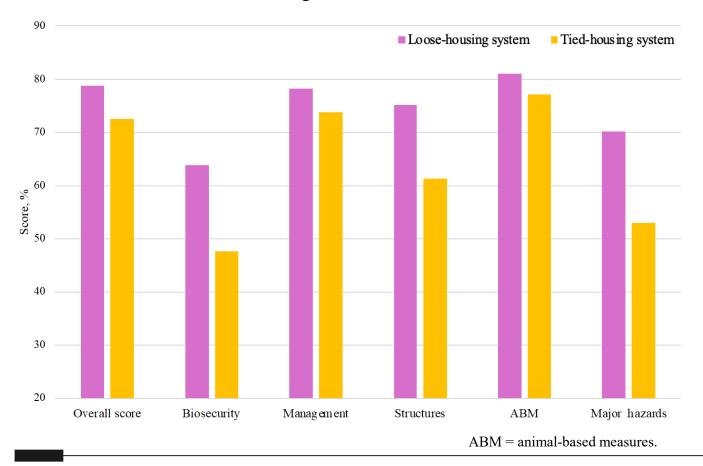
#### **Descriptive statistics**

T	Loose housi	ng system	Tied housing system		
Traits —	Mean (SD)	Range	Mean (SD)	Range	
Fat, %	$3.96 \pm 0.24$	3.18 -4.83	$4.01 \pm 0.28$	3.14 -4.85	
Protein, %	$3.42\pm0.16$	2.80 -3.99	$3.34 \pm 0.22$	2.78 -3.99	
Casein, %	$2.71 \pm 0.14$	2.16 -2.71	$2.65 \pm 0.19$	2.16 -3.20	
Lactose, %	$4.77 \pm 0.07$	4.45 -4.96	$4.73 \pm 0.10$	4.45 -4.95	
Urea, mg/dL	25.11 ±4.85	8.40 -42.60	22.91 ±5.99	8.40 -42.60	
SCS	$3.91 \pm 0.73$	1.16 -6.27	$4.05 \pm 0.98$	0.60 -6.97	
TBC, CFU/mL	14.24 ±25.77	10.00-722.00	$24.24 \pm 183.74$	10.00 -10'000.00	
$\log_{10}$ TBC	$1.07 \pm 0.19$	1.00 -2.86	$1.19 \pm 0.27$	1.00 -4.00	
Fat-to-protein	$1.16 \pm 0.06$	0.92 -1.45	$1.21 \pm 0.07$	0.86 -1.54	

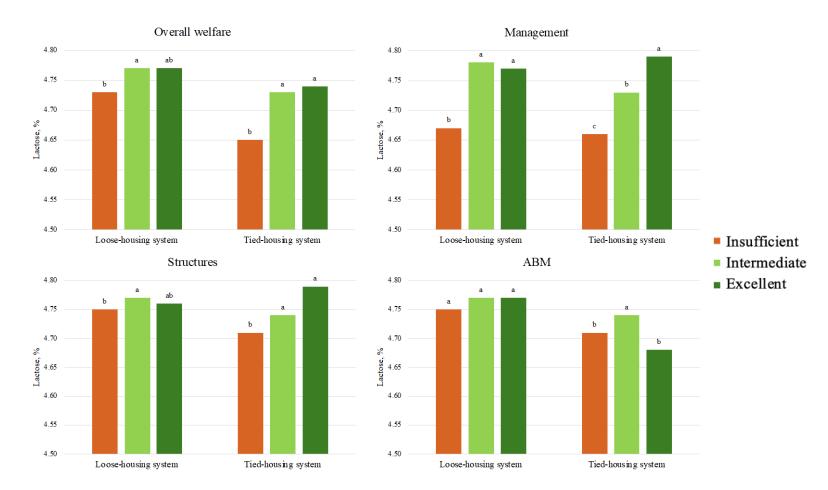
TBC= total bacterial count.

SCS= somatic cells score calculated as SCS=3+log2(SCC/100).

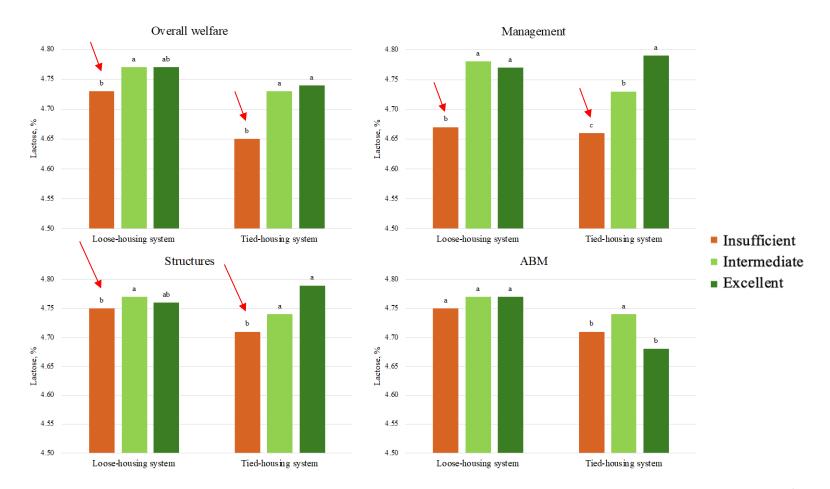
#### **ClassyFarm scores**



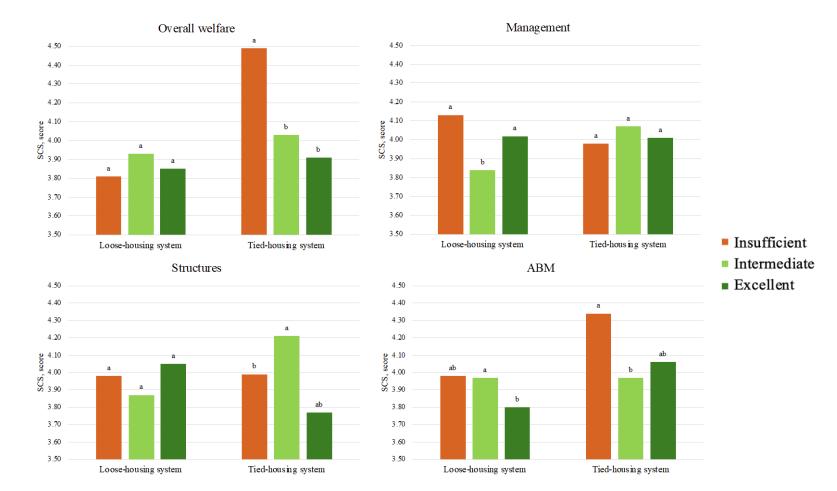
#### Lactose



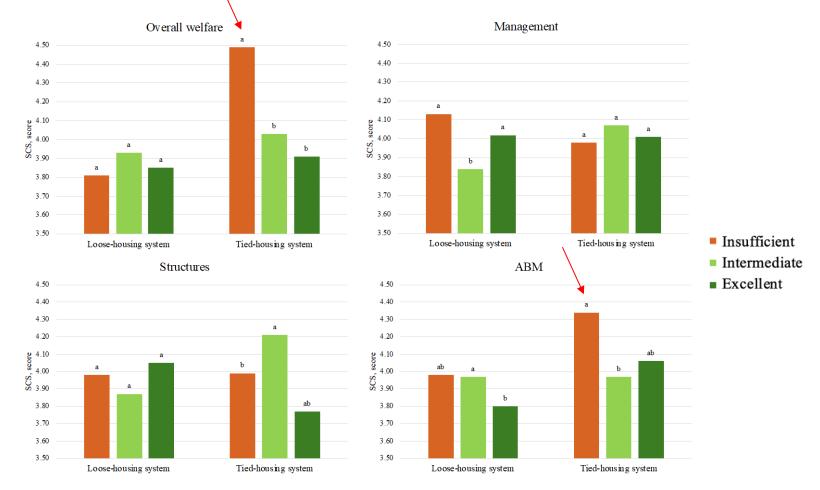
#### Lactose



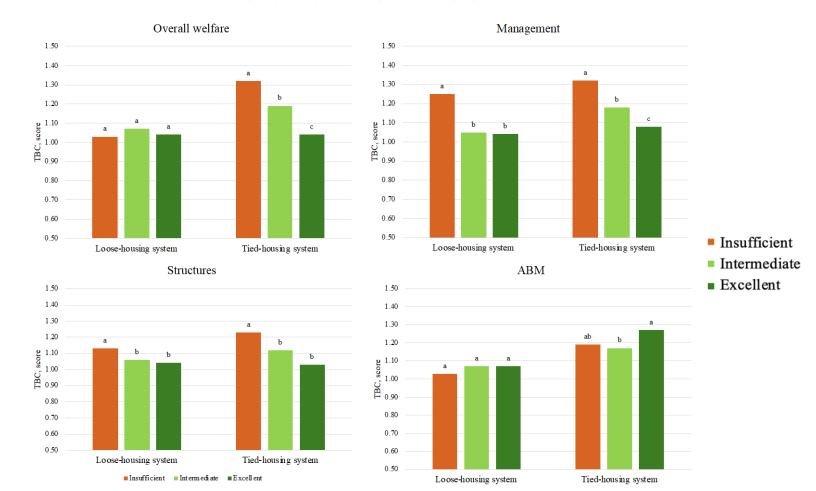
#### Somatic cell score



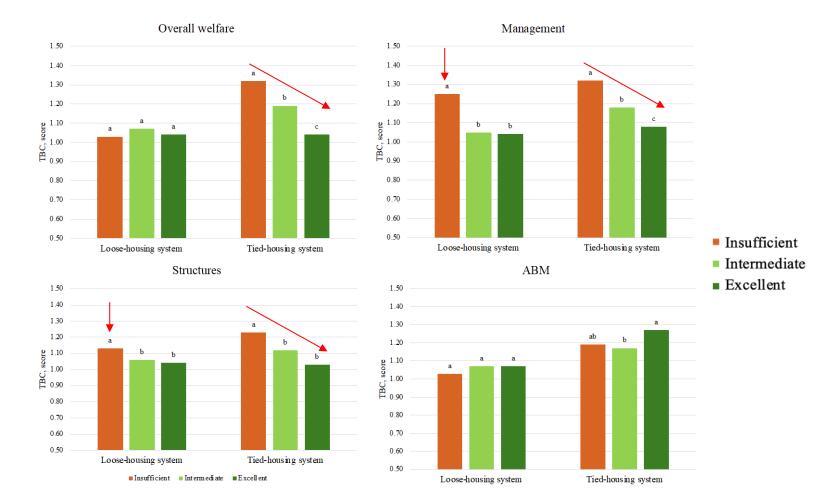
#### Somatic cell score



#### **Total bacterial count**



#### **Total bacterial count**





### Conclusions

Farms in the worst welfare class exhibited the highest TBC and SCS and the lowest lactose content.

ClassyFarm as a tool to be implemented in the milk quality payment system to stimulate the farmers both to enhance animals welfare conditions and milk quality traits.

To better understand causal relationship of welfare scores and milk traits... more research!



#### Thanks for the kind attention

Do you have any questions?

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#### **Additional materials**

	Ec	ı. [1]	]	Eq. [2]	Eq. [3] Eq. [4]			
Trait <sup>3</sup>								
	Welfare	Welfare * HS	Management	Management * HS	Structures	Structures * HS	ABM	ABM * HS
Fat, %	4.66**	9.76***	8.76 ***	9.23 ***	5.64 **	5.09 **	1.10	5.23 **
Protein, %	2.61 †	4.27 *	1.12	0.30	15.43 ***	1.87	1.40	0.41
Casein, %	2.51 †	5.31 **	1.54	0.64	13.85 ***	2.32 †	1.79	0.95
Lactose, %	11.62 ***	1.26	77.75 ***	12.77 ***	12.39 ***	3.76 *	10.95 ***	10.16 ***
Urea, mg/dL	1.43	2.33 †	0.30	13.75 ***	4.31 *	6.40 **	3.58 *	5.58 **
SCS, score	1.79	3.58 *	1.08	3.87 *	0.89	6.22 **	4.03 *	4.73 **
TBC, score	8.18 ***	7.92 ***	52.90 ***	3.70 *	24.85 ***	1.50	4.86 **	5.89 **
Fat-to-protein	17.07 ***	1.47	20.96 ***	7.86 ***	33.04 ***	1.64	6.36 **	4.93 **
FA, $g/100 g$								
De novo FA	1.26	12.28 ***	3.08 *	12.29 ***	6.77 **	3.13 *	0.81	1.10
Mixed FA	0.68	7.29 ***	1.28	10.10 ***	1.92	0.34	0.29	0.89
Preformed FA	17.36 ***	0.30	28.54 ***	3.18 *	10.99 ***	11.44 ***	6.09 **	4.62 *
MUFA	13.06 ***	1.63	24.97 ***	3.69 *	6.47 **	11.30 ***	5.34 **	5.49 **
PUFA	9.52 ***	6.43 **	1.64	11.63 ***	13.87 ***	13.55 ***	8.44 ***	7.33***
SFA	0.73	11.52 ***	2.71 †	11.30 ***	3.17 *	1.57	0.04	3.39 *

 $<sup>^{1}***</sup>P < 0.001$ ; \*\*P < 0.01; \*P < 0.05, †P < 0.10.

 $<sup>^{2}</sup>$  HS = housing system; ABM = animal-based measures.

 $<sup>^3</sup>$  TBC =  $\log_{10}$  of total bacterial count (CFU/mL); FA = Fatty acids.

**Table 6.** Least squares means (standard error) of bulk milk traits<sup>1</sup> for the fixed effect of housing system<sup>2</sup>. Means with different superscript letters within row are significantly different ( $P \le 0.05$ ).

Trait	LHS	THS
Fat, %	4.02 <sup>a</sup> (0.01)	4.01 <sup>a</sup> (0.02)
Protein, %	$3.42^{a}(0.01)$	$3.34^{b}(0.02)$
Casein, %	$2.71^{a}(0.01)$	$2.64^{b}(0.02)$
Lactose, %	$4.76^{a}(0.01)$	4.71 <sup>b</sup> (0.01)
Urea, mg/dL	$24.40^{a}(0.40)$	$23.60^{a}(0.50)$
SCS, score	$3.86^{b} (0.06)$	$4.14^{a}(0.08)$
TBC, score	$1.05^{b} (0.02)$	$1.18^a (0.02)$
Fat-to-protein	$1.18^{b} (0.005)$	1.21 <sup>a</sup> (0.006)
FA, g/100 g milk		
De novo	$0.89^{a}(0.01)$	$0.84^{b}(0.01)$
Mixed	$1.40^{a}(0.01)$	$1.37^{b}(0.01)$
Preformed	$1.19^{b} (0.01)$	$1.27^{a}(0.01)$
MUFA	$1.00^{b} (0.01)$	$1.04^{a}(0.01)$
PUFA	$0.085^{a}(0.001)$	$0.084^{a}(0.001)$
SFA	$2.73^{a}(0.01)$	$2.68^{b}(0.01)$

<sup>&</sup>lt;sup>1</sup> TBC = log<sub>10</sub> of total bacterial count (CFU/mL); FA = Fatty acids;

<sup>&</sup>lt;sup>2</sup> LHS = loose housing system; THS = tied housing system.