

## Supplemental Material

### The effect of the genotypes of the *CSN2* gene on test-day milk yields in the Slovak Holstein cow

#### Supplemental material S1 – Genetics structure of Slovak Holstein cattle

**Table S1.** Genotype and allelic frequencies for *CSN2* gene in a population of Slovak Holstein cattle per breed.

Herd	Genotype frequencies			Allelic frequencies		$\chi^2$	p
	A1A1	A1A2	A2A2	A1	A2		
Herd 1	0.1496	0.4567	0.3937	0.3780	0.6220	0.104	0.9491
Herd 2	0.1379	0.4598	0.4023	0.3678	0.6322	0.011	0.9945
Herd 3	0.0159	0.6984	0.2857	0.3651	0.6349	16.163	0.003
Herd 4	0.0684	0.3632	0.5684	0.25	0.75	0.230	0.8914
Herd 5	0.0310	0.3320	0.6370	0.1970	0.8030	2.337	0.3108
<b>Total</b>	<b>0.0528</b>	<b>0.3708</b>	<b>0.5764</b>	<b>0.2382</b>	<b>0.7618</b>	<b>0.694</b>	<b>0.7069</b>

**Table S2.** The population genetic indices for *CSN2* gene in a population of Slovak Holstein cattle per breed.

Herd	He <sub>(obs)</sub>	He <sub>(exp)</sub>	PIC	E	ENA	V %	F <sub>IS</sub>
Herd 1	0.4567	0.4702	0.3596	0.5298	1.8875	47.39	0.0287
Herd 2	0.4598	0.4650	0.3570	0.535	1.8692	47.04	0.0111
Herd 3	0.6984	0.4636	0.3561	0.5364	1.8463	47.26	-0.5065
Herd 4	0.3632	0.3750	0.3047	0.625	1.6	37.66	0.0315
Herd 5	0.3320	0.3164	0.2664	0.6836	1.4628	31.67	-0.0493
<b>Total</b>	<b>0.3708</b>	<b>0.3629</b>	<b>0.2971</b>	<b>0.6371</b>	<b>1.5696</b>	<b>36.31</b>	<b>-0.0218</b>

## Supplemental material S2 - Multiple comparisons between CSN2 genotype groups

**Table S3.** CSN2 genotype Least Squares Means comparisons - milk kg

CSN 2	Milk kg LSMEAN	LSMEAN Number
11	32.6246680	1
12	32.8660110	2
22	33.0439772	3

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*Least Squares Means for effect CSN2*  
 $Pr > |t|$  for  $H_0: LS\text{Mean}(i)=LS\text{Mean}(j)$

*Dependent Variable: Milk kg*

<i>i/j</i>	1	2	3
1		0.4280	0.0732
2	0.4280		0.1126
3	0.0732	0.1126	

**Table S4.** CSN2 genotype Least Squares Means comparisons - fat kg

CSN 2	Fat kg LSMEAN	LSMEAN Number
11	1.18586792	1
12	1.18383171	2
22	1.18252520	3

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*Least Squares Means for effect CSN2*  
 $Pr > |t|$  for  $H_0: LS\text{Mean}(i)=LS\text{Mean}(j)$

*Dependent Variable: Fat kg*

<i>i/j</i>	1	2	3
1		0.9646	0.9056
2	0.9646		0.9322
3	0.9056	0.9322	

**Table S5.** CSN2 genotype Least Squares Means comparisons – fat %

CSN 2	Fat % LSMEAN	LSMEAN Number
11	3.75156951	1
12	3.70272478	2
22	3.68289739	3

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*Least Squares Means for effect CSN2*  
 $Pr > |t|$  for  $H_0: LS\text{Mean}(i)=LS\text{Mean}(j)$

*Dependent Variable: Fat %*

<i>i/j</i>	1	2	3
1		<b>0.0242</b>	<b>0.0006</b>
2	0.0242		0.0549
3	0.0006	0.0549	

**Table S6.** CSN2 genotype Least Squares Means comparisons - protein kg

CSN 2	Protein kg LSMEAN	LSMEAN Number
11	1.05678001	1
12	1.06083980	2
22	1.07193063	3

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*Least Squares Means for effect CSN2*  
*Pr > |t| for H0: LSMean(i)=LSMean(j)*

*Dependent Variable: Protein kg*

i/j	1	2	3
1		0.7946	<b>0.0381</b>
2	0.7946		<b>0.0003</b>
3	0.0381	0.0003	

**Table S7.** CSN2 genotype Least Squares Means comparisons – protein %

CSN2	Protein % LSMEAN	LSMEAN Number
11	3.30155631	1
12	3.29569765	2
22	3.31511985	3

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*Least Squares Means for effect CSN2*  
*Pr > |t| for H0: LSMean(i)=LSMean(j)*

*Dependent Variable: Protein %%*

i/j	1	2	3
1		0.7653	0.2314
2	0.7653		<b>&lt;.0001</b>
3	0.2314	<.0001	

**p < 0.05** significant differences, **p < 0.01** highly significant differences

# Supplemental material S3 - Multiple comparisons between herds

**Table S8.** Herd Least Squares Means comparisons - milk kg

<b>Herd</b>	<b>Milk kg</b>	
	<b>LSMEAN</b>	<b>LSMEAN Number</b>
<i>Herd1</i>	36.8824538	1
<i>Herd2</i>	33.9344846	2
<i>Herd3</i>	26.8113516	3
<i>Herd4</i>	34.0743805	4
<i>Herd5</i>	32.5217566	5

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*Least Squares Means for effect Herd*  
 $Pr > |t| \text{ for } H_0: LS\text{Mean}(i)=LS\text{Mean}(j)$

*Dependent Variable: Milk kg*

<i>i/j</i>	1	2	3	4	5
1		<.0001	<.0001	<.0001	<.0001
2	<.0001		<.0001	0.9831	<.0001
3	<.0001	<.0001		<.0001	<.0001
4	<.0001	0.9831	<.0001		<.0001
5	<.0001	<.0001	<.0001	<.0001	

**Table S9.** Herd Least Squares Means comparisons - fat kg

<b>Herd</b>	<b>Fat kg</b>	
	<b>LSMEAN</b>	<b>LSMEAN Number</b>
<i>Herd1</i>	1.29921928	1
<i>Herd2</i>	1.19686961	2
<i>Herd3</i>	0.98821177	3
<i>Herd4</i>	1.22339508	4
<i>Herd5</i>	1.21267898	5

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*Least Squares Means for effect Herd*  
 $Pr > |t| \text{ for } H_0: LS\text{Mean}(i)=LS\text{Mean}(j)$

*Dependent Variable: Fat kg*

<i>i/j</i>	1	2	3	4	5
1		<.0001	<.0001	<.0001	<.0001
2	<.0001		<.0001	0.0771	0.2406
3	<.0001	<.0001		<.0001	<.0001
4	<.0001	0.0771	<.0001		0.6790
5	<.0001	0.2406	<.0001	0.6790	

**Table S10.** Herd Least Squares Means comparisons – fat %

<b>Herd</b>	<b>Fat %</b>	
	<b>LSMEAN</b>	<b>LSMEAN Number</b>
<i>Herd1</i>	3.61123238	1
<i>Herd2</i>	3.64520011	2
<i>Herd3</i>	3.84887859	3
<i>Herd4</i>	3.64731172	4
<i>Herd5</i>	3.80936335	5

Least Squares Means for effect Herd <i>Pr</i> >   <i>t</i>   for <i>H</i> <sub>0</sub> : <i>LS</i> Mean( <i>i</i> )= <i>LS</i> Mean( <i>j</i> )					
Dependent Variable: Fat %					
<i>i/j</i>	1	2	3	4	5
1		0.5266	<.0001	0.4976	<.0001
2	0.5266		<.0001	1.0000	<.0001
3	<.0001	<.0001		<.0001	0.4475
4	0.4976	1.0000	<.0001		<.0001
5	<.0001	<.0001	0.4475	<.0001	

**Table S11.** Herd Least Squares Means comparisons - protein kg

<i>Herd</i>	<i>Protein kg</i>	<i>LS</i> MEAN
	<i>LS</i> MEAN	<i>Number</i>
<i>Herd1</i>	1.15070013	1
<i>Herd2</i>	1.09367916	2
<i>Herd3</i>	0.93432596	3
<i>Herd4</i>	1.06880460	4
<i>Herd5</i>	1.06840756	5

  

Least Squares Means for effect Herd <i>Pr</i> >   <i>t</i>   for <i>H</i> <sub>0</sub> : <i>LS</i> Mean( <i>i</i> )= <i>LS</i> Mean( <i>j</i> )					
Dependent Variable: Protein kg					
<i>i/j</i>	1	2	3	4	5
1		<.0001	<.0001	<.0001	<.0001
2	<.0001		<.0001	0.0182	0.0002
3	<.0001	<.0001		<.0001	<.0001
4	<.0001	0.0182	<.0001		1.0000
5	<.0001	0.0002	<.0001	1.0000	

**Table S12.** Herd Least Squares Means comparisons - protein %

<i>Herd</i>	<i>Protein %</i>	<i>LS</i> MEAN
	<i>LS</i> MEAN	<i>Number</i>
<i>Herd1</i>	3.17357861	1
<i>Herd2</i>	3.27121905	2
<i>Herd3</i>	3.57144039	3
<i>Herd4</i>	3.17548508	4
<i>Herd5</i>	3.32889988	5

  

Least Squares Means for effect Herd <i>Pr</i> >   <i>t</i>   for <i>H</i> <sub>0</sub> : <i>LS</i> Mean( <i>i</i> )= <i>LS</i> Mean( <i>j</i> )					
Dependent Variable: Protein %%					
<i>i/j</i>	1	2	3	4	5
1		<.0001	<.0001	0.9997	<.0001
2	<.0001		<.0001	<.0001	<.0001
3	<.0001	<.0001		<.0001	<.0001
4	0.9997	<.0001	<.0001		<.0001
5	<.0001	<.0001	<.0001	<.0001	

**p** < 0.05 significant differences, **p** < 0.01 highly significant differences