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1	Haptoglobin and serum amyloid A in bulk tank milk in relation to raw milk
2	quality
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## 25 Summary

The aim of the present study was to evaluate relationships between the presence of the two major bovine acute phase proteins haptoglobin (Hp) and serum amyloid A (SAA) and raw milk quality parameters in bulk tank milk samples. Hp and SAA have been suggested as specific markers of mastitis but recently also as markers for raw milk quality. Since mastitis has detrimental effects on milk quality, it is important to investigate if the presence of Hp or SAA indicates such changes in the composition and properties of the milk. Bulk tank milk samples (n=91) were analysed for Hp, SAA, total protein, casein, whey protein, proteolysis, fat, lactose, somatic cell count and coagulating properties. Samples with detectable levels of Hp had lower casein content, casein number and lactose content, but higher proteolysis than samples without Hp. Samples with detectable levels of SAA had lower casein number and lactose content, but higher whey protein content than samples without SAA. The presence of APP in bulk tank milk is suggested as an indicator for unfavourable changes in the milk composition, e.g. protein quality, due to udder health disturbances, with economical implications for the dairy industry. 

50 The bulk tank milk composition has a decisive influence on the production of safe, 51 high quality dairy products. One major factor responsible for the deterioration of the 52 quality of the raw milk is mastitis, and its negative effect on the milk composition is 53 well established (Kitchen, 1981; Munro et al. 1984). It is the sub-clinical form of 54 mastitis that constitutes problems to the dairies, since these cases often go undetected 55 and the milk is delivered to the dairy (Leitner et al. 2008). Studies show that during 56 mastitis the casein content, valuable for the cheese making industry, will decrease 57 while the whey protein content will increase (Barbano et al. 1991; Auldist et al. 1996; 58 Urech et al. 1999). In addition, increased proteolysis is often observed in milk from 59 cows with mastitis (Schaar, 1985; Auldist et al. 1996). Proteolysis in milk is one of 60 the major product deteriorating factors with negative impact on the quality and 61 stability of milk and dairy products (Mara et al. 1998; Kelly et al. 2006). Saeman et 62 al. (1988) found that after an udder infection the proteolytic activity may sustain even 63 though the SCC has returned to normal levels. Larsen et al. (2004) established that 64 casein degradation not only occurred in the infected quarter but also in the 65 neighbouring quarters, even though there was no effect on the SCC. In the bulk tank, 66 milk from healthy udder quarters will be commingled with milk from infected 67 quarters, and thus, the entire bulk tank may be affected by protein degradation. This is 68 a problem, especially for the cheese making industry, since curd formation properties 69 will be impaired and yield reduced (Mara et al. 1998, Leitner et al. 2008). Likewise, 70 proteolytic activity in UHT-milk may cause off flavours and gelation, and 71 consequently reduced shelf-life of the products (Ma et al. 2000; Santos et al. 2003; Barbano et al. 2006; De Noni et al. 2007). 72

73

74	Milk somatic cell count (SCC) has been used extensively since the 1960s in the
75	diagnosis of mastitis, and the BTMSCC (bulk tank milk somatic cell count) is widely
76	used in the assessment of raw milk quality. In many EU countries milk payment
77	systems favours a low BTMSCC. There is, however, no clear scientific data defining
78	the level of BTMSCC that is associated with additional benefits in terms of milk
79	quality. Several authors have reported that SCC is not a suitable indicator of
80	proteolysis in quarter milk samples (Le Roux et al. 1995; Urech et al. 1999) and
81	recent studies have also shown that the BTMSCC gives a poor prediction of raw milk
82	quality for cheese production (Leitner et al. 2008). Research to find new sensitive and
83	specific markers for disadvantageous changes in raw milk composition due to udder
84	health disturbances is therefore warranted.

86 The acute phase proteins (APP) have become important diagnostic markers of disease 87 in human medicine and are also being evaluated in veterinary diagnostics (Eckersall, 88 2004). The major bovine APP are haptoglobin (Hp) and serum amyloid A (SAA), 89 which both increase dramatically upon infection, inflammation or trauma. Hp and 90 SAA are mainly produced by the liver but are also produced locally in the mammary 91 gland (McDonald et al. 2001; Hiss et al. 2004). Some studies have found that Hp and 92 SAA have antibacterial effects (Eaton et al. 1982; Hari-Dass et al. 2005; Larson et al. 93 2005) and considering that they are locally produced, their role in the inflammatory 94 defence is interesting. In several studies Hp and SAA in milk have been evaluated as 95 markers for mastitis (Horadagoda et al. 1999; Eckersall et al. 2001; Grönlund et al. 96 2003; Nielsen et al. 2004; Grönlund et al. 2005; Eckersall et al. 2006; Hiss et al. 97 2007), but so far little attention has been paid regarding their potential in predicting 98 changes in milk composition and technological properties of the raw milk. In a

99	previous study we reported that detectable levels of Hp and SAA could be found in
100	bulk tank milk samples (Åkerstedt et al. 2007). In a more recent study we also
101	investigated APP in relation to raw milk quality parameters in cow composite milk
102	samples (Åkerstedt et al. 2008). To our knowledge, these papers are the only studies
103	examining APP in bovine bulk tank milk or applying APP research in the field of
104	product quality. However, no studies have been reported on Hp and SAA in relation
105	to the quality of the raw bulk tank milk. Quality programs for milk payment and
106	advisory measures to improve the raw milk quality are mostly based on analysis of
107	bulk tank milk samples. For APP to be a potential candidate as indicator for
108	unfavorable changes in milk composition due to udder health disturbances in the herd
109	it is important that levels of APP in the bulk tank milk are related to important quality
110	traits of the raw milk.

The aim of this study was to investigate relationships between the presence of Hp and SAA, and different raw milk quality parameters, i.e. total protein, casein, whey protein, proteolysis, fat, lactose and SCC in bulk tank milk samples. In addition, APP in relation to the coagulating properties of the bulk tank milk samples were evaluated.

117 Materials and methods

118 Bulk tank milk samples

The study included 91 bulk tank milk samples collected from different dairy farms in cooperation with the Milko dairy cooperative (Grådö, Hedemora, Sweden). One representative sample from each farm was taken by the tanker driver just before

- 122 emptying the bulk tank, in connection with the ordinary milk collection, which
- 123 occurred every second day. At the sampling occasion the farms delivered 90-13,025

124 kg milk (average 1,610 kg), indicating that the herd size of the participating farms

125 varied markedly. The average BTMSCC for the samples was 195,000 cells/ml,

ranging from 33,000 to 1,365,000 cells/ml (median 146,000 cells/ml). The bulk tank

127 milk samples were collected at the dairy plant for further transportation to the

128 university laboratory the same day. Sample aliquots for the analyses of Hp, SAA and

129 proteolysis were frozen and stored at -70°C until analysis, whereas the other

130 parameters were analysed using fresh milk samples.

131

132 Assay of haptoglobin and serum amyloid A

133 Hp was analysed by an earlier described optical biosensor assay (Åkerstedt *et al.* 

134 2006; Åkerstedt et al. 2008) with some additional modifications. In this study, the Hp

135 surface was prepared by using a solution of 20 mg/l instead of 500 mg/l Hp in 0.01 M

136 acetate buffer, and the activation of the surface during immobilisation was reduced

137 from 7 minutes to 3 minutes. For regeneration of the sensor surface the concentration

138 of sodium dodecyl sulphate (SDS) was increased from 2 mM to 3 mM. An extra

139 reconditioning step was added, in which 50 mM glycine pH 9.5 was injected over the

140 sensor surface for 30 seconds, after the ordinary regeneration step. Bovine Hp (Life

141 diagnostics, Clarkston, GA, USA) was used for immobilization and standards, and the

142 limit of detection (LOD) of the modified assay was 0.3 mg/l.

143

144 SAA was determined using a commercial ELISA with a LOD of 0.3 mg/l (Phase<sup>TM</sup>

145 Serum Amyloid A Assay, Tridelta Development Ltd, Wicklow, Ireland).

146

147 Measurement of the somatic cell count, total protein, whey protein and casein content,

148 casein number, fat and lactose content

149	SCC in the bulk tank milk samples was measured by an electronic fluorescence based
150	cell counting technique (Fossomatic 5000, Foss, Hillerød, Denmark). Total protein,
151	fat and lactose contents were measured on fresh milk using mid infrared spectroscopy
152	(Fourier Transform Instrument, FT 120, Foss). The casein content was determined by
153	an indirect method which was described earlier (Åkerstedt et al. 2008).
154	
155	Measurement of proteolysis
156	The extent of proteolysis in the milk sample was measured according to a
157	fluorescamine method as previously described (Wiking et al. 2002).
158	
159	Measurement of coagulating properties
160	The coagulating properties of the milk was measured with a Bohlin VOR Rheometer
161	(Malvern Instruments Nordic AB, Uppsala, Sweden) according to Hallén et al. (2007)
162	with one minor modification, i.e. Chymax Plus, strength 200 IMCU per gram
163	(Christian Hansen A/S, DK-2970, Hørsholm, Denmark), was used instead of pure
164	chymosin. The coagulation time was measured, i.e. the time (s) elapsed from
165	chymosin addition until a weak coagulum corresponding to 5 Pa was formed. In
166	addition, curd firmness (Pa) was measured 25 min after chymosin addition.
167	
168	Statistical analyses
169	Parametric t-test using SAS (Version 9.1, SAS Institute Inc., Cary, NC, USA) was
170	used to evaluate the relationships between APP and the different raw milk quality
171	parameters analysed. The bulk tank milk samples were categorised into two groups;

- 172 detectable or non-detectable levels of Hp or SAA, based on the detection limits of the
- assays used to determine the proteins in milk. SCC and curd firmness were

174 logarithmically transformed before statistical analyses to obtain normally distributed

175 data. Differences between groups were considered significant if  $p \le 0.05$ .

176

## 177 **Results**

178 Descriptive statistics for the milk quality parameters analysed in the study are

179 presented in Table 1. Detectable levels of Hp were found in 19 (21%) of the 91 bulk

tank milk samples. The average Hp concentration in these 19 samples was 1.02±0.99

181 mg/l and the average SCC in these samples was 387,000±266,000 cells/ml. Detectable

182 levels of SAA were found in 68 (75%) of the 91 bulk tank milk samples. The average

183 SAA concentration in these 68 samples was 1.12±1.16 mg/l and the average SCC in

184 these samples was 218,000±179,000 cells/ml. Samples not containing detectable

185 levels of Hp had the average SCC of 144,000±78,000 cells/ml while samples not

186 containing detectable levels of SAA had the average SCC of 127,000±116,000

187 cells/ml.

188

189 Table 1 near here.

190

Table 2 and 3 present differences in milk composition between samples with and without detectable levels of Hp and SAA, respectively. Bulk tank milk samples in which Hp could be detected had a lower casein number, contained less casein, had increased proteolysis, a lower lactose content and higher SCC compared to samples without detectable levels of Hp. Bulk tank milk samples in which SAA could be detected had a lower casein number, increased whey protein, lower lactose content and higher SCC compared to samples not containing detectable levels of SAA.

198

199 Table 2, 3 and 4 near here.

200



## 209 **Discussion**

The present study is, to our knowledge, the first investigating relationships betweenthe quality of raw bulk tank milk and the presence of APP as indicator of mastitis.

212 When studying markers for udder health disturbances and effects on milk composition

it is important to have in mind that the type of milk sample, i.e. quarter, cow

214 composite or bulk tank milk, may affect the results. Due to the dilution effect from

215 quarter milk to composite and bulk tank milk, significant relationships found at

216 quarter level may not be present at cow or bulk tank level and vice versa. However,

the main findings of the present study, i.e. unfavourable changes in protein

218 composition in bulk tank milk samples with detectable levels of APP, are largely in

agreement with the results of our earlier study in cow composite milk (Åkerstedt *et al.* 

220 2008).

221

In this study, bulk tank milk samples with detectable levels of Hp had lower casein

223 content as well as lower casein number. This was most likely due to increased

224 proteolytic activity, since these samples also had significantly higher proteolysis than 225 samples without detectable levels of Hp. Bulk tank milk samples containing 226 detectable levels of SAA also had a lower casein number compared to samples 227 without detectable levels of SAA. In this case, however, the effect was likely due to 228 the observed increased whey protein content in the samples. It is thus likely that 229 decreased synthesis, proteolysis and influx of components from the blood will occur 230 simultaneously but to evaluate to what extent was outside the scope of this paper. 231 In this study, detectable levels of Hp in bulk tank milk was related to increased 232 proteolytic activity, which is in contrast to the results in our previous study on cow 233 composite milk samples (Åkerstedt et al. 2008). One possible explanation for not 234 observing proteolysis in cow composite milk but in bulk tank milk may be related to 235 the fact that bulk tank milk consists of commingled milk from different milkings, as 236 well as milk from non-infected and sub-clinically infected glands (Leitner et al. 237 2008). Moreover, the storage time before collection and freezing was longer for bulk 238 tank, than for cow composite milk samples, allowing proteolysis to proceed for a 239 longer time. The bulk tank milk samples were collected at the dairy plant and were 240 kept at +4°C during the entire chain from farm to laboratory. Since the milk is 241 collected every second day, the oldest batch of milk in the tank was stored for 242 approximately 2.5 days before it was frozen. Refrigerated storage of raw milk is 243 known to favor the growth of psychrotropic bacteria, which may produce heat-244 resistant extra-cellular proteases and lipases. Proteases are mainly secreted at the end of the log phase of the bacterial growth, at numbers in the order of  $10^7$  cfu/ml, 245 246 indicating that very high numbers of bacteria are required to result in proteolysis 247 (Sørhaug & Stepaniak, 1997). In a study by Haryani et al. (2003), such high numbers 248 were reached first after 7 days at 4°C, and proteolysis, as measured by the

249	fluorescamine method, was observed on day 6. Considering the very high numbers of
250	bacteria needed for proteolysis to become a problem, it is unlikely that microbial
251	contamination of milk samples would explain the observed relationships between the
252	presence of Hp and increased proteolysis. An influence of microbial proteases cannot,
253	however, be excluded at this stage.
254	
255	
256	Bulk tank milk samples with detectable levels of SAA contained less lactose in
257	agreement with earlier studies on udder quarter and cow composite milk samples
258	(Lindmark-Månsson et al. 2006; Åkerstedt et al. 2008). In the present study, bulk tank
259	milk samples with detectable levels of Hp also contained reduced levels of lactose.
260	The most common explanation to decreased lactose content is reduced synthesis due
261	to damaged epithelial cells. Another explanation, suggested by Silanikove et al.
262	(2000), might be that proteolysis of $\beta$ -case n will result in release of peptides with a
263	regulatory effect on lactose secretion.
264	
265	Bulk tank milk samples with detectable levels of Hp and SAA had significantly higher
266	SCC compared to samples without Hp and SAA, respectively. In our previous study,
267	bulk tank milk samples with detectable levels of SAA, but not samples with
268	detectable levels of Hp, had higher SCC (Åkerstedt et al. 2007). The discrepancy
269	between the studies might be explained by the use of different statistical methods,
270	differences in the categorisation of the samples and the selection of milk samples. To
271	assess the potential of APP as markers for milk quality in comparison with the
272	commonly applied SCC, we also investigated relationships between SCC and the
273	different quality traits in the same samples. In this study, bulk tank milk samples with

274 elevated SCC (>195,000 cells/ml), had a lower casein number than samples with a 275 lower SCC (<195,000 cells/ml). In our previous study (Åkerstedt *et al.* 2008) with 276 cow composite milk, samples with elevated SCC (>83,000 cells/ml) had reduced 277 lactose and increased whey protein content compared to samples with a lower SCC 278 (<83,000 cells/ml). The threshold values used in those studies (195 000 cells/ml and 279 83,000 cells/ml) are median and mean values, respectively. These threshold values are 280 relevant since several studies demonstrate that the composition is deteriorated on 281 quarter and cow level between 50,000-100,000 cells/ml, while the Swedish milk 282 payment system give additional bonus payment when the bulk tank contains less than 283 175,000-200,000 cells/ml. Our studies suggest that the presence of APP in milk is 284 related to disadvantageous changes in several milk quality parameters and that these 285 relationships are valid in both cow composite and bulk tank milk. SCC, on the other 286 hand, is not related to milk quality traits to the same extent as APP and the type of 287 relationships observed differs between cow composite and bulk tank milk.

288

289 No significant relationships between APPs and coagulation properties, i.e. coagulation 290 time and curd firmness, were found in this study. Since there are no other studies 291 published investigating APP in relation to the coagulating properties of milk, it is 292 difficult to evaluate the results obtained. In earlier studies, significant correlations 293 between mastitis and impaired coagulating properties were observed (for review see 294 Munro et al. 1984). In contrast, Leitner et al. (2008) found no correlation between 295 coagulation time and SCC at bulk tank or silo level. In general, most published 296 research on coagulating properties is based on studies with milk from a small number 297 of animals (Barbano et al. 1991; Mazal et al. 2007). It is also common that batches of 298 milk with a specific SCC are constructed by pooling milk with very high SCC, often

originating from cows with clinical mastitis, and milk with a low SCC. Such a
procedure is not ideal, as the composition of milk from cases of clinical mastitis is
very deviant from milk originating from cows without clinical signs. Consequently,
this type of constructed milk samples are not representative for real bulk tank milk
samples, representing commingled milk from a large number of clinically healthy
cows.

305

306 The casein content is an important quality parameter in cheese production and 307 decreased casein content implies large losses for the dairy industry. At present, the 308 total protein content is used as a major quality parameter, largely affecting the milk 309 price to the producer. The total protein content, however, also includes the whey 310 proteins, of which those originating from blood, with no interest for the dairies, will 311 increase during mastitis. The extent of proteolysis is another important factor 312 influencing milk quality, although not presently assessed. There are thus no 313 techniques in place allowing reliable, large-scale analyses of the protein quality of the 314 milk, although research and development in this field is ongoing. In a future 315 perspective, the dairies might have the possibility to differentiate raw milk based on 316 quality to be used for different purposes. SCC is a good marker for udder health 317 disturbances at udder quarter level but several studies have demonstrated that SCC is 318 not a strong candidate for predicting the processing quality of the bulk tank milk (Le 319 Roux et al. 1995; Urech et al. 1999; Leitner et al. 2006; Leitner et al. 2008). Since 320 many dairy products like cheese and fermented products require high protein quality it 321 should be of great importance to have a sensitive and specific marker for 322 disadvantageous changes in milk composition, e.g. those associated to poor udder 323 health. Changes in levels of such a marker should preferably be associated to the

324	protein composition of the raw milk. In this study, Hp and SAA have shown to be
325	potential candidates for predicting the raw bulk tank milk quality, specifically in
326	relation to protein quality. This study and our previous studies therefore suggest that
327	APP may be used as indicators for changes in milk composition as consequence of
328	udder health disturbances, in quarter or cow composite milk samples at the farm, as
329	well as in bulk tank milk at the dairy plant or milk grading laboratory.
330	
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333	Department of Food Science for their technical assistance, the dairy cooperative
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335	sampling.
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**Table 1.** Milk composition, including contents of haptoglobin (Hp) and serum

509	amyloid A (SAA), a	nd coagulating propert	ies of 91 bulk tank milk	samples.
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Parameter	Unit	Mean (SD)	Minimum	Maximum
Нр	mg/l	$\mathrm{ND}^\dagger$	<0.3	4.70
SAA	mg/l	ND	<0.3	8.79
Total protein	%	3.54 (0.18)	3.12	4.21
Casein	%	2.58 (0.14)	2.20	3.07
Casein number		0.73 (0.01)	0.71	0.75
Whey protein	%	0.91 (0.05)	0.82	1.09
Proteolysis	eq leu <sup>‡</sup>	1.11 (0.11)	0.89	1.71
Fat	%	4.48 (0.43)	3.54	6.25
Lactose	%	4.61 (0.11)	4.30	4.91
Somatic cell count	cells/ml	195,000 (169,600)	33,000	1,365,000
Coagulation time	S	120 (42)	32	278
Curd firmness	Pa	350 (118)	162	798

<sup>†</sup>ND= not determined. Since many of the samples did not contain detectable levels of

514 Hp or SAA, i.e levels were below 0.3 mg/l, it was not considered relevant to calculate

515 a mean value.

 $\ddagger$  eq leu = equivalent mM leucine.

**Table 2.** Differences in milk composition between bulk tank milk samples with (Hp+)522and without (Hp-) detectable levels of Hp. Differences between Hp+ and Hp- samples523were evaluated by parametric t-test and were considered significant if  $p \le 0.05$ .

	Hp+ (n=19)	SE <sup>†</sup>	Hp- (n=72)	SE	p-value
Total protein (%)	3.493	0.025	3.549	0.023	NS <sup>‡</sup>
Casein (%)	2.532	0.018	2.593	0.017	0.016
Casein number	0.725	0.001	0.731	0.001	< 0.001
Whey protein (%)	0.913	0.008	0.908	0.006	NS
Proteolysis (eq leu) <sup>§</sup>	1.159	0.019	1.100	0.014	0.038
Fat (%)	4.455	0.049	4.491	0.055	NS
Lactose (%)	4.579	0.014	4.618	0.014	0.05
Log SCC (cells/ml)	5.528	0.049	5.104	0.027	< 0.001
Coagulation time (s)	110	9.862	123	4.963	NS
Log curd firmness (Pa)	2.547	0.039	2.517	0.015	NS

 $^{\dagger}$  SE = Standard Error

 $^{\ddagger}$  NS = not significant

528 <sup>§</sup> eq leu = equivalent mM leucine.

**Table 3.** Differences in milk composition between bulk tank milk samples with

545 (SAA+) and without (SAA-) detectable levels of SAA. Differences between SAA+

and SAA- samples were evaluated by parametric t-test and were considered

547 significant if  $p \le 0.05$ .

	SAA+ (n=68)	SE <sup>†</sup>	SAA- (n=23)	SE	p-value
Total protein (%)	3.553	0.021	3.488	0.042	NS <sup>‡</sup>
Casein (%)	2.588	0.016	2.559	0.032	NS
Casein number	0.728	0.001	0.734	0.001	< 0.001
Whey protein (%)	0.918	0.006	0.883	0.011	0.004
Proteolysis (eq leu) <sup>§</sup>	1.122	0.014	1.082	0.023	NS
Fat (%)	4.477	0.049	4.504	0.102	NS
Lactose (%)	4.583	0.012	4.692	0.018	< 0.001
Log SCC (cells/ml)	5.261	0.029	4.989	0.063	< 0.001
Coagulation time (s)	122	5.055	113	9.276	NS
Log curd firmness (Pa)	2.521	0.016	2.533	0.030	NS

 $^{\dagger}$  SE = Standard Error

 $^{\ddagger}$  NS = not significant

552 <sup>§</sup> eq leu = equivalent mM leucine.

**Table 4.** Differences in milk composition between bulk tank milk samples with high568SCC (>195,000 cells/ml) and low SCC (<195,000 cells/ml). Differences between high</td>569SCC and low SCC samples were evaluated by parametric t-test and were considered570significant if  $p \le 0.05$ .

	high SCC (n=29)	SE <sup>†</sup>	low SCC (n=62)	SE	p-value
Total protein (%)	3.498	0.027	3.555	0.025	NS
Casein (%)	2.541	0.021	2.599	0.018	NS
Casein number	0.726	0.001	0.731	0.001	0.002
Whey protein (%)	0.909	0.008	0.909	0.007	NS
Proteolysis (eq leu) <sup>§</sup>	1.126	0.020	1.105	0.014	NS
Fat (%)	4.453	0.059	4.498	0.060	NS
Lactose (%)	4.585	0.014	4.622	0.015	NS
Coagulation time (s)	118	8.053	121	5.359	NS
Log curd firmness (Pa)	2.507	0.030	2.531	0.015	NS

 $^{\dagger}$  SE = Standard Error

 $^{\ddagger}$  NS = not significant

575 <sup>§</sup> eq leu = equivalent mM leucine.