

Antimicrobial treatment of bovine subclinical mastitis during lactation



Bart van den Borne¹, Gerdien van Schaik², Theo Lam^{2,3} and Mirjam Nielen¹

1 Utrecht University, Faculty of Veterinary Medicine, Department of Farm Animal Health, Utrecht, The Netherlands 2 GD Animal Health Service, Deventer, The Netherlands

3 Dutch Udder Health Centre, Deventer, The Netherlands



Email: b.vandenborne@uu.nl

Introduction

- Bacteriological cure rates after lactational antimicrobial treatment of bovine subclinical mastitis differ between pathogens, being low for chronic Staphylococcus aureus mastitis.
- Cure rates are influenced by several cow and quarter factors such as parity, number of quarters infected, lactation stage and location of quarter.
- It is unknown whether early treatment of subclinical mastitis will improve cure rates.

Objective

 To determine bacteriological cure rates after antimicrobial treatment of recently acquired subclinical mastitis during lactation in a randomized field trial.

Materials and Methods

 39 herds participated in the trial and were selected based on participation in the fourweekly test day interval and on a high incidence of new elevated cow somatic cell count (CSCC) cases.

- Cows with a first elevated CSCC (CSCC>250,000 cells/ml) after two consecutive low CSCC were aseptically sampled twice for bacteriological culture.
- Pathogen-specific blocks within a herd were filled to randomly allocate antimicrobial treatment to bacteriological culture positive cows. All bacteriological-positive quarters were treated within a cow.
- A quarter was defined to be cured if the prior identified pathogen was not cultured in two consecutive milk samples three and four weeks post treatment.
- Statistics: multivariable hierarchical logistic regression model at the quarter level with corrections for random cow- and herd-effects.
- Intra-class correlation coefficients (ICC) were calculated for random effects.

Results

 Treatment was associated with higher bacteriological cure and differed between pathogens (Table 1).

- Cure of penicillin-sensitive Staphylococcus aureus after treatment was remarkably higher than 34% as observed earlier in chronic cases in the Netherlands.
- Bacteriological cure was associated with treatment, two pathogens factors and with quarter SCC in the statistical model (Table 2).
- Previously established cow factors were not related to cure of recently acquired subclinical mastitis.
- The random herd effect was insignificant and was left out of the model.
- The ICC for the random cow effect was 0.27.

Conclusions

- Treatment of recently acquired subclinical mastitis results in higher cure rates than natural cure and had no cow factors associated with bacteriological cure.
- Previously established cow factors are therefore deemed to be related to chronic infections and do not apply for treatment of recently acquired subclinical mastitis.
- Whether treatment of recently acquired subclinical mastitis is economically attractive remains to be analyzed and should take into account transmission of pathogens within the herd.



Table 1. Crude bacteriological cure rates.

	Control			T	Treatment			
Pathogen	All	Cure	%	All	Cure	%		
Penicillin-sensitive Staph. aureus	114	25	21.9	34	22	64.7		
Penicillin-resistant Staph. aureus	35	3	8.6	12	6	50.0		
Coagulase-negative staphylococci	226	66	29.2	66	52	78.8		
Streptococcus uberis	67	27	40.3	23	15	65.2		
Streptococcus dysgalactiae	25	8	32.0	19	15	79.0		
Streptococcus agalactiae	16	1	6.3	7	7	100		
Other streptococci	21	11	52.4	20	18	90.0		
Escherichia coli	13	8	61.5	6	6	100		

Table 2. Factors associated with bacteriological cure of recently acquired subclinical mastitis.

Variable	Category		P-Wald	OR	95%-CI	
Treatment	Yes	2.09	0.00	8.11	5.14	12.82
	No	-	-	ref.	-	-
Pathogen	Penicillin-sensitive Staphylococcus aureus	0.33	0.28	1.39	0.76	2.54
Penicillin-resistant Staphylococcus aureus Coagulase-negative staphylococci Streptococcus dysgalactiae Streptococcus uberis and other streptococc	Penicillin-resistant Staphylococcus aureus	-0.71	0.14	0.49	0.19	1.26
	Coagulase-negative staphylococci	0.54	0.04	1.72	1.02	2.88
	Streptococcus dysgalactiae	0.97	0.02	2.65	1.16	6.08
	Streptococcus uberis and other streptococci	-	-	ref.	-	-
Quarter SCC	Continuous (on a log scale)	-0.23	0.01	0.79	0.66	0.95
Infectious status pretreatment	Bacteriological culture-negative or other pathogen present in second sample	1.32	0.00	3.74	2.53	5.53
	Same pathogen present in second sample	-	-	ref.	-	-

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