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# The effect of the dam-calf relationship on serostatus to *Neospora caninum* on 20 Costa Rican dairy farms

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#### Abstract

An epidemiological study was conduced on 20 dairy herds previously diagnosed as seropositive for Neospora caninum. The number of females per farm varies from 41 to 296. All females present on the farms were bled once in the period of July and August 2000. A total of 3002 females were bled. An indirect ELISA was used to determine the serostatus of the animals. The analysis of the data was performed in four steps: (1) descriptive statistics about the serological status and general characteristics of the cattle; (2) calculation of vertical and horizontal transmission; (3) an univariate analysis and, (4) a multivariate logistic regression analysis with herd as random effect. The within-herd seroprevalence varied between 25.0% (34/136) and 70.5% (203/288). Seven hundred and forty-seven dam-daughter pairs were available, involving daughters of any age. Daughters in the specific age-class of 2- and 3-years old had a higher seroprevalence (P < 0.01) compared with younger and older age-classes. The risk of being seropositive when being born to a mother that tested seropositive (prevalence ratio (PR)) was 2.8-fold increased which coincides with a 5.3-fold increased odds. The probability of horizontal infection amounts to 0.22. The probability of a seropositive offspring due to vertical transmission was 0.64 (attributable fraction among exposed  $(AF_{exp})$ ). The multivariate logistic regression showed a significant 6.0-fold increased odds for being seropositive when born from a seropositive mother. Also the within-herd seroprevalence level was significantly associated with the serostatus of the daughters. In the specific conditions of the dairy herds involved in this study, the serostatus of the cows should be not used as a predictor of the serostatus of daughters due to the increased probability of horizontal transmission. © 2003 Elsevier Science B.V. All rights reserved.

Keywords: Neospora caninum; Vertical and horizontal transmission; Serostatus; Dairy cattle; Epidemiology

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#### 1. Introduction

Neosporosis is a recently recognised parasitic (protozoan) disease caused by *Neospora caninum*. This disease affects a wide range of animals and has effects on cattle reproduction. It causes foetal losses by resorption, mummification or abortion between the 4th and the 9th month of pregnancy (Dubey, 1999). Since the first report of *Neospora*-associated abortion in cattle (Thilsted and Dubey, 1989), many studies have reported *N. caninum* as a causative agent of abortion around the world (Anderson et al., 1995; Otter et al., 1995; Boulton et al., 1995; Dubey et al., 1997; Thurmond et al., 1997; Venturini et al., 1999; Wouda et al., 1999a,b; Jenkins et al., 2000). In Costa Rica neosporosis has been diagnosed and associated with bovine abortion since 1996 (Pérez et al., 1998; Romero et al., 2000).

The life cycle of Neospora was elucidated in 1998, when the dog was recognised as the definitive host (McAllister et al., 1998). Therefore, dogs may play an important role in the introduction and maintenance of the infection in herds (Dubey, 1999). Two means of transmission among cattle are assumed: vertical and horizontal. The vertical mode (transplacentally from the mother to her calf) is the most prevalent means of transmission (Paré et al., 1996; Thurmond et al., 1997; Schares et al., 1998; Venturini et al., 1999; Davison et al., 1999a). Horizontal transmission has not been documented until recently. Some authors report findings that suggest that horizontal transmission is possible by means of ingestion of feed or water contaminated with Neospora oocysts (McAllister et al., 1998; De Marez et al., 1999; Wouda et al., 1999b; Dubey, 1999; Anderson et al., 2000; Bergeron et al., 2000). Another suggested means of horizontal transmission in newborn calves is by ingestion of colostrum contaminated with tachyzoites, especially in the first hours of life (Uggla et al., 1998). Ingestion of infected amniotic fluid or infected placenta have been proposed in other studies as potential routes of infection (Ho et al., 1997; Piergili-Fioretti et al., 2000; Bergeron et al., 2001); however, these suggested means of infection in cows remain to be proven, despite evidence for this in dogs (Dijkstra et al., 2001).

It is widely accepted that abortion due to neosporosis is the result of a lethal infection of the foetus following a maternal parasitaemia. However, an intra-uterine infection with *N. caninum* will not always induce foetal loss. Factors such as the timing of the parasitaemia during gestation, the quantity and duration of the parasitaemia, the effectiveness of the maternal immune response and the ability of the foetus to mount an immune response have been recognised as crucial factors that determine the fate of the foetus (Hemphill et al., 2000). The acquisition of infection in the third trimester of gestation allows the foetus to develop a sufficient grade of immunocompetency to limit the parasite multiplication, resulting in a calf that will be born clinically healthy (Williams et al., 2000).

Several studies in which *N. caninum* antibodies were measured before colostrum intake by calves report that between 81 and 95% of seropositive cows gave birth to seropositive calves (Paré et al., 1996; Wouda et al., 1998; Davison et al., 1999a). On the other hand, in a cross-sectional study there was not a clear relationship between the serologic status of dam and offspring, resulting in proportions of seropositive daughters born from seropositive dams ranging from 0 to 85% (Bergeron et al., 2000).

Other cross-sectional studies have focused on the proportions of seropositive daughters born to seronegative dams in an effort to document postnatal acquisition of *Neospora*. Thus, Dyer et al. (2000) using IFAT to detect antibodies for *Neospora* reported that 77.9%

(120/154) of seronegative dams had seronegative daughters, while the rest of seronegative dams (22.1%) had seropositive daughters. In the same way, Dijkstra et al. (2001), using an ELISA based on a detergent lysate of whole sonicated tachyzoite, reported that 83.5% (207/248) of seronegative dams had seronegative daughters and 16.5% (41/248) had seropositive daughters.

The objective of our study was to assess the mother—calf relationship with regard to serological reactivity to *Neospora* in 20 Costa Rican dairy herds. We investigated whether the serological status of the dam, could be used as a predictor for the serological status of the calf. If so, this prediction might be used as potential criterion for selective culling.

## 2. Materials and methods

## 2.1. Study population

In this study, 20 dairy farms previously diagnosed as seropositive to *Neospora* were used. The farms were located between 1500 and 2200 m above sea level, the temperature range were between 12.9 and 28 °C and the relative humidity oscillated between 77 and 93%. There were two clearly defined seasons, which were dry and rainy. The total rain precipitation was approximately 500 mm in the dry season and 3500 mm in the rainy season. The farms involved in this study were selected intentionally. Since 1986 they participated in a research project focused on dairy production and herd health, and all of them used VAMPP 5.1 (Veterinary Automated Management and Production control Programme, [Noordhuizen and Buurman, 1984]) as their management information system. The most frequent breeds in these herds were Holstein–Friesian, Jersey and crosses between these. The number of females (cows and calves) per farm varied from 41 to 296.

#### 2.2. Data collection

All females present in the selected farms were bled once in the period of July/August 2000. Sera were stored at  $-20\,^{\circ}$ C until assay. An indirect ELISA (Paré et al., 1995) with 87.8% sensitivity (CI 95%: 81.3–94.3) and 96.4% specificity (CI 95%: 93.3–99.5) for detection of antibodies against *Neospora* was used to determine the serologic status of the animals (Romero et al., 2002).

VAMPP 5.1 was used as the source of basic data (animals per farm, identification of the cow, age, breed, date of birth, days of gestation at the moment of sampling and genealogy of the cows). Dam-daughter pairs were extracted from VAMPP and merged with a file containing the ELISA results.

## 2.3. Data analysis

Descriptive statistics about the general characteristics of the cattle, regarding the serological status, especially for the dam–daughter pairs, were performed using STATA® (StataCorp., 1999).

Status of daughters	Status of dam		
	Seropositive	Seronegative	
Seropositive	a	b	
Seronegative	c	d	
-	a + c	b+d	

Table 1 Contingency  $(2 \times 2)$  table to explain the way of calculation of vertical and horizontal transmission of the disease as was performed in this study

## 2.3.1. Calculation of vertical and horizontal transmission

The overall probability of horizontal transmission, calculated as the proportion of seropositive daughters born from seronegative dams with its exact binomial 95% confidence interval (95% CI) was assessed using EpiInfo 6.0 (Center for Disease Control and Prevention, USA).

The probability of vertical transmission was calculated in several steps. In order to carry out this calculation we used a contingency  $(2 \times 2)$  table in which the seropositivity of the dams was defined as exposure (Table 1). As it is a cross-sectional study, with plausible evidence of horizontal transmission, we assumed that a proportion of seropositive dams (a+c) were horizontally infected after the date the calf was born. Therefore the "crude" frequencies were adjusted in the following way. First, the proportion of daughters that was horizontally infected (b/b+d) was calculated. Next, the frequencies of a, c and (a+c) were reduced according to this proportion while these subtracted numbers were added to b, d and (b+d), respectively. For both the crude and adjusted frequencies the probabilities of vertical transmission using the attributable fraction among exposed  $(AF_{exp})$  were calculated.  $AF_{exp}$  represents the proportion of daughters that were seropositive due to being born from seropositive mothers. Confidence intervals for these proportions were calculated with EpiInfo 6.0 (Center for Disease Control and Prevention, USA).

## 2.3.2. Univariate analyses

Age-specific seroprevalences of daughters and mothers were calculated using age categories of 1 year each, except for calves under 1-year old which were divided in two categories: under 150 days old and between 151 and 365 days old. Cows of more than 7-years old were grouped in the same age-class. Differences between proportions of seropositives by age-class were tested by means of the  $\chi^2$  statistic using EpiInfo 6.0 (Center for Disease Control and Prevention, USA).

In order to eliminate the effect of possible false positives by maternal antibodies in daughters (Davison et al., 1999b; Tizard, 2000), the age of daughters was categorised in two levels, 150 days or less, and older than 150 days. A new data set keeping pairs with daughters older than 150 days was constructed. Furthermore, to assess the effect of the within-herd seroprevalence on the means of transmission of the disease, herds were divided in three categories. Herds with seroprevalence between 25 and 35% were classified as 'low', herds with seroprevalence between 35.1 and 45% were classified as 'medium' and herds with seroprevalence higher than 45.1% were classified as 'high'. A contingency table for pairs in each subset was constructed to calculate the probability of horizontal transmission.

The prevalence ratio (PR) as measure of association and the  $AF_{exp}$  as measure of effect of the exposure to seropositive dams were calculated using the epidemiological software WinEpiscope 2.0 (Thrusfield et al., 2001).

## 2.3.3. Multivariate analysis

Finally, a multivariate logistic regression model with herd as random effect was used to assess the relationship between the serostatus of the daughter and her age, the serostatus of the mother and the within-herd seroprevalence (STATA® release 6.0, StataCorp., 1999). The multivariate analysis can only be carried out on unadjusted frequencies, because we do not know which mothers were horizontally infected after the calf was born.

#### 3. Results

## 3.1. Seroprevalence

A total of 3002 females were bled on the 20 farms studied. The overall seroprevalence for *Neospora* was 39.7% (1191/3002), whereas the within-herd seroprevalence varied between 25.0% (34/136) and 70.5% (203/288). Taking the sensitivity and specificity of the test into account, the overall prevalence is 42.8%.

Seven hundred and forty-seven dam—daughter pairs were available for analysis. In the group of daughters, 614 (82.2%) were less than 2 years old, whereas 711 (95.2%) were younger than 3 years old. In the group of dams, 684 (91.5%) were more than 3 years old. Apparent seroprevalences in daughters and dams were 33.3% (249/747) and 38.2% (285/747), respectively. For the age categories of dams the seroprevalence varied between 33.8 and 44.3%, with no statistical differences between age categories (Fig. 1). Daughters in

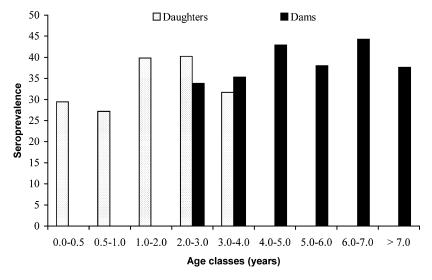


Fig. 1. Seroprevalence in dams and daughters by age groups.

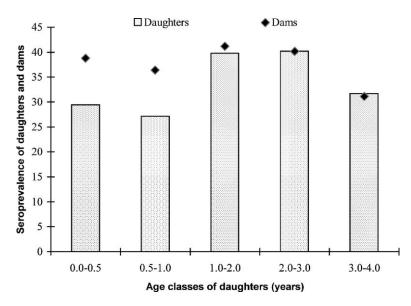


Fig. 2. Seroprevalence for daughters and their dams, by age-groups of daughters, at the date of sampling. Seropositive daughters born from seropositive or negative dams and negative daughters born from seropositive or negative dams were present in each age-group.

the specific age-class of 2 and 3-years old had a higher seroprevalence (P < 0.01) compared with the other age-classes, but not among them. Furthermore, there were no differences in seroprevalence between the other age-classes (Fig. 1). In all age-classes of daughters there were seropositive and seronegative daughters that were born from seropositive or seronegative dams. Furthermore, the percentage of seropositive dams (around 38.0%) within the age categories of daughters was not significantly different (P > 0.5) (Fig. 2).

## 3.2. Association between serostatus of daughters and dams

Results of the 747 pairs are shown in Table 2. The risk of being seropositive is 2.8-fold increased when being born from a mother that is tested seropositive (prevalence ratio) which coincides with a 5.3-fold increased odds, both measures are based on the adjusted numbers.

The probability of horizontal transmission is 0.22 (110/512). Of seropositive daughters born from seropositive mothers, 63.7% is due to the fact that their mothers were seropositive (AF<sub>exp</sub>), which is the probability of seropositive offspring due to vertical transmission.

From the general data set containing 747 pairs, a subset with 542 pairs was selected excluding daughters younger than 150 days old at the time of bleeding. In this data set, 93.2% of daughters were younger than 3 years old, while 92.3% of dams were more than 3 years old.

The seroprevalences among daughters and dams in this subset were 34.9% (189/542) and 37.8% (205/542), respectively. An increased risk of being seropositive was observed in daughters that were born from seropositive dams (PR = 2.6, 95% CI: 2.1-3.3). The proportion of daughters horizontally infected was 0.23 (95% CI: 0.19-0.28). In this subset,

Table 2 Probabilities of vertical (AF $_{\rm exp}$ ) and horizontal transmission of N. caninum

	Daughters	Dams		Proportion of vertical	Proportion of horizontal	Prevalence ratio
		Positive	Negative	transmission (95% CI)	transmission (95% CI)	(95% CI)
General (747 pairs)	Positive	168 [139]	81 [110]	0.70 (0.69, 0.71)	0.18 (0.14, 0.21)	3.4 (2.7, 4.2)
	Negative	117 [96]	381[402]	[0.64 (0.62, 0.66)]	[0.22 (0.18, 0.25)]	[2.8 (2.3, 3.3)]
Older than 150 days	Positive	122 [105]	67 [84]	0.67 (0.65, 0.68)	0.20 (0.16, 0.25)	3.0 (2.3, 3.8)
(542 pairs)	Negative	83 [71]	270 [282]	[0.62 (0.60, 0.63)]	[0.23 (0.19, 0.28)]	[2.6 (2.1, 3.3)]
High (85 pairs)	Positive	40 [37]	13 [16]	0.43 (0.41, 0.46)	0.42 (0.25, 0.61)	1.8 (1.1, 2.8)
	Negative	14 [13]	18 [19]	[0.38 (0.36, 0.41)	[0.46 (0.29, 0.63)]	[1.6 (1.1, 2.4)]
Medium (154 pairs)	Positive	37 [35]	24 [26]	0.60 (0.58, 0.62)	0.25 (0.17, 0.35)	2.5 (1.7, 3.7)
	Negative	22 [21]	71 [72]	[0.58 (0.56, 0.60)]	[0.27 (0.18, 0.36)]	[2.4 (1.6, 3.5)]
Low (303 pairs)	Positive	45 [43]	30 [32]	0.71 (0.69, 0.72)	0.14 (0.09, 0.19)	3.4 (2.3, 5.1)
. 1	Negative	47 [45]	181 [183]	[0.70 (0.68, 0.71)]	[0.15 (0.10, 0.20)]	[3.3 (2.2, 4.8)]

This table includes the overall probabilities using all pairs, the probabilities when daughters younger than 150 days were excluded and the probabilities of the latter data-set stratified to high, medium and low within-herd seroprevalence. PR is included as measure of association. Numbers and output between square brackets correspond to values adjusted for the proportion of dams infected horizontally.

Factor	Level	Frequency (%)	Prevalence (daughters)	β (95% CI)	OR (95% CI)	P
Serostatus of	Seronegative	61.8	0.18	_	1.0	_
the dam at sampling	Seropositive	38.2	0.59	1.79 (1.62, 1.97)	6.0 (5.1, 7.2)	0.000
Age of the	>150 days	72.6	0.35	_	1.0	_
daughter at sampling	≤150 days	27.4	0.28	-0.72 (-1.34, -0.10)	0.5 (0.3, 0.9)	0.023
Within-herd seroprevalence	High	15.7	0.62	_	1.0	_
	Medium	28.4	0.40	0.49 (0.09, 0.89)	1.6 (1.1, 2.4)	0.021
	Low	55.9	0.25	1.21 (0.75, 1.52)	3.4 (2.1, 4.6)	0.000

Table 3 Multivariate analysis of the relation between seropositivity to *N. caninum* for daughters with serostatus of the dam at sampling, age of the daughter and the within-herd seroprevalence

The proportion of the total variance contributed by the herd-level variance component ( $\rho$ , rho) was 17.8, where  $\rho = \sigma_{\nu}^2/(\sigma_{\nu}^2 + 1)$ , log of standard deviation = -1.52, standard deviation ( $\sigma_{\nu}$ ) = 0.46.

the probability of vertical transmission resulting in antibodies was 0.62 (95% CI: 0.60–0.63) (Table 2).

In herds with high, medium and low within-herd seroprevalence, the probability of horizontal transmission increases as the within-herd seroprevalence increases while the probability of vertical transmission decreases as the within-herd seroprevalence increases. (Tables 2 and 3).

The multivariate logistic regression with herd as random effect showed a significant 6.0-fold increased odds for being seropositive when born from a seropositive mother. Also the within-herd seroprevalence level was significantly associated with the sero-status of the daughters (Table 3). The inclusion of the herd effect showed that 18% of the variation is due to herd related factors that were not included in the analysis.

## 4. Discussion

## 4.1. General

In this study, we were able to estimate both vertical and horizontal transmission of *N. caninum*, because of the presence of seropositive daughters born from seropositive dams, as well as seropositive daughters born from seronegative dams. We are aware that the results of this study can be partially biased because the study design used is not an optimal one for making accurate and high quality estimates about the impact of vertical and horizontal transmission. For example, a dam that tested seropositive at the moment of bleeding might have been negative at the moment the calf was born. Indeed, she might have acquired the infection after parturition in a horizontal way. In a cross-sectional study this bias will lead to an underestimation of the horizontal transmission and subsequently in an overestimation of the vertical transmission. We adjusted for this at least partially by assuming that the level of horizontal transmission was equal in dams and daughters. A cohort study, as has been

carried out by Davison et al. (1999a,b), in which both dam and daughter are bled at the moment of parturition would give better information about the sero-status at the moment of birth, and when sero-conversion occurred.

Other sources of misclassification can be: variation in antibody titers throughout pregnancy (Conrad et al., 1993; Hemphill et al., 2000; Stenlund et al., 1999; Maley et al., 2001), a transient rise of antibody titers in daughters without a true *Neospora* infection (Hietala and Thurmond, 1999) and the limitation of an imperfect diagnostic test with a relatively low sensitivity (0.88) compared to the specificity (0.97). However, it is very unlikely that these potential sources of bias affect the results to a large extent (e.g. true prevalence in both daughters and dams hardly change after correction for imperfect SE and SP). Also, there is still a lack of information about the behavior of the infection and the course of antibody titers over time in naturally infected cattle.

### 4.2. Vertical and horizontal transmission

Under the assumption that horizontal transmission in dams and daughters is equal, we adjusted for the horizontal infection in dams. Then, the proportion of animals infected vertically reduces from 0.70 to 0.64, while the proportion of horizontal infection increases from 0.18 to 0.22 (Table 2, all pairs). When excluding animals that might still have colostral antibodies, the proportion of vertical transmission reduces from 0.64 (all pairs) to 0.62 (542 pairs). As both proportions are quite similar it can be concluded that presence of colostral antibodies has little influence on the outcome of the proportion of animals vertically or horizontally infected. Hietala and Thurmond (1999) determined that colostral anti-Neospora antibodies decayed at 128 days after parturition. In our study, seropositive daughters fewer than 150 days could be positive due to intra-uterine infection and/or horizontal infection and/or colostral antibodies. It is possible that in those daughters under 150 day of age, antibodies titers were not increased much by colostral uptake, and the presence of antibodies was mainly due to intra-uterine infection and/or postnatal acquired infection, as in the group older than 150 days. The probabilities of vertical infection that were found in this study were lower than in other studies that used blood samples before ingestion of colostrum. Those studies reported probabilities of vertical infection over 0.80 (Paré et al., 1996; Schares et al., 1998; Davison et al., 1999a; Hietala and Thurmond, 1999). However, in two other cross-sectional studies, carried out in Spain (Mainar-Jaime et al., 1999) and Canada (Bergeron et al., 2000), the rates of vertical transmission, calculated as AF<sub>exp</sub>, were lower than 60.0%. The authors of those studies conclude that the low vertical transmission rate could be due to differences related to environmental conditions resulting in an increased efficiency of horizontal transmission. Secondly, the low probability of vertical transmission found in our study may be due to a continuous stimulation of the maternal immune system in conditions of endemicity. Under these conditions, dams might be frequently exposed to environmental sources of the parasite, leading to a continuous stimulation of the immune system sufficient to prevent the exposure of the foetus to an infective dose of tachyzoites (Paré et al., 1996). However, there is no conclusive evidence about the role of the maternal immune system in prevention of vertical transmission of neosporosis. Thirdly, antibody concentrations may decline to a level below the detection limit when the infection occurred a long time ago. This might explain some cases in which mothers were negative while daughters were positive. This fluctuation in antibody titres for *Neospora* was demonstrated by Maley et al. (2001) in experimentally infected cattle. We do not have firm evidence that indicates a different behaviour in naturally infected cows.

In our study, we observed an increased probability of horizontal transmission as the within-herd seroprevalence increased, whereas the probability of vertical infection decreased at the same time. This seroprevalence related probability of horizontal transmission might be explained by an increased exposure to environmental sources of infection. These could include infected placentas or amniotic fluid (Ho et al., 1996, 1997; Bergeron et al., 2001) or food and water contaminated with *Neospora* oocysts from naturally infected dogs (Dubey, 1999).

## 4.3. Age-specific results

A higher seroprevalence for daughters in the age-class between 1 and 3 years old was observed (Fig. 1). This was also observed by Dijkstra et al. (2001). The differences in seroprevalence by age-classes in daughters might be due to a point source of infection as suggested by others (McAllister et al., 1996, 2000; Dijkstra et al., 2001). Furthermore, there might have been unrecognised host and/or environmental factors just previously to the reproductive age, which promotes the horizontal acquisition of the infection. This strongly suggests that horizontal transmission is present in the studied dairy herds, but that the level of it may not be constant over time. Other studies reported no difference in seroprevalence by age-classes, suggesting that the horizontal transmission plays a minimal role in the transmission of the disease (Davison et al., 1999a,b). However, by means of mathematical models, it was estimated that in the presence of imperfect vertical transmission, that without horizontal transmission prevalence of the disease has to decline over time (French et al., 1998, 1999). Our study supports this finding. The seroprevalences for the dams in all age-classes of the daughters were not different but seroprevalences do differ between age categories of daughters (Fig. 2), which might indicate that the serostatus of the daughters was not solely due to vertical infection.

In Costa Rica, veterinary practitioners have used the serostatus of an animal to predict the serostatus of its mother or its off-spring. This attitude was based on studies showing vertical transmission rates of over 80% (Paré et al., 1996; Schares et al., 1998; Davison et al., 1999a; Hietala and Thurmond, 1999). However, under the specific conditions of the dairy herds involved in this study, the serostatus of the mother should be not used as a predictor of the serostatus of her daughter(s), because of the much lower probability of vertical transmission.

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