

Table S1. Species and genera of the *Mollicutes* class isolated from the analyzed nasal swabs.

Isolates	# positives (%)	# pure cultures (%)
<i>M. bovirhinis</i>	283 (39.8)	130 (18.3)
<i>M. bovis</i>	136 (19.1)	47 (6.6)
<i>M. dispar</i>	86 (12.1)	51 (7.2)
<i>M. arginini</i>	40 (5.6)	7 (1)
<i>M. alkalescens</i>	26 (3.7)	9 (1.3)
<i>M. ovipneumoniae</i>	5 (0.7)	4 (0.6)
<i>M. fermentans</i>	3 (0.4)	2 (0.3)
<i>Ureaplasma spp.</i>	66 (9.3)	4 (0.6)
<i>A. laidlawii</i>	16 (2.2)	10 (1.4)

Table S2: Parameter estimates of the logistic mixed effects model analyzing the isolation frequency of organisms of the *Mollicutes* class.

Fixed Effects			
<i>Predictors</i>	<i>Log-Odds</i>	<i>95% CI</i>	<i>p</i> *
(Intercept)	-0.09	-0.65 – 0.61	0.830
time [day 15]	1.53	0.80 – 2.35	0.001
time [day 60]	1.80	0.66 – 2.73	0.001
Random Effects			
<i>Groups</i>	<i>Predictors</i>	<i>Std.Dev</i>	<i>Corr.</i>
farm	(Intercept)	1.27	
	time [day 15]	1.30	-0.46
	time [day 60]	1.56	-0.49

* Wald's test *p* value

Table S3: Analysis of deviance table (type II likelihood ratio tests) of the full model relating the isolation frequency of organisms of the *Mollicutes* class to the variables time and season.

Fixed Effects				
<i>Predictors</i>	<i>npar</i>	<i>AIC</i>	X^2	<i>p</i>
time	2	759.68	9.50	0.009
season	1	752.52	0.34	0.56
Random Effects				
<i>Predictors</i>	<i>npar</i>	<i>AIC</i>	X^2	<i>p</i>
time farm	6	820.55	78.37	7.75x10 ⁻¹⁵

Legend: *npar* = number of parameters associated to the relative predictor; *AIC* = Akaike's information criterion value observed upon dropping of the relative predictor; X^2 = likelihood ratio test statistic; *p* = likelihood ratio test *p* value.

Table S4: Parameter estimates of the logistic mixed effects model analyzing the frequency of *M. bovis* isolation

Fixed Effects			
<i>Predictors</i>	<i>Log-Odds</i>	<i>95% CI</i>	<i>p</i> *
(Intercept)	-4.72	-6.68 – -2.58	0.002
time [day 15]	4.30	1.90 – 6.62	0.005
time [day 60]	2.79	0.79 – 4.65	0.049
Random Effects			
<i>Groups</i>	<i>Predictors</i>	<i>Std.Dev</i>	<i>Corr.</i>
farm	(Intercept)	2.94	
	time [day 15]	3.20	-0.94
	time [day 60]	2.40	-0.97

* Wald's test *p* value

Table S5: Analysis of deviance table (type II likelihood ratio tests) of the full model relating the frequency of isolation of *M. bovis* to the variables time and season.

Fixed Effects				
<i>Predictors</i>	<i>npar</i>	<i>AIC</i>	X^2	<i>p</i>
time	2	600.69	13.24	0.001
season	1	589.69	0.25	0.62
Random Effects				
<i>Predictors</i>	<i>npar</i>	<i>AIC</i>	X^2	<i>p</i>
time farm	6	619.79	40.35	3.89x10 ⁻⁷

Legend: *npar* = number of parameters associated to the relative predictor; *AIC* = Akaike's information criterion value observed upon dropping of the relative predictor; X^2 = likelihood ratio test statistic; *p* = likelihood ratio test *p* value.

Table S6: Parameter estimates of the logistic mixed effects model analyzing the frequency of *M. bovis*-specific PCR positives.

Fixed Effects			
<i>Predictors</i>	<i>Log-Odds</i>	<i>95% CI</i>	<i>p</i> *
(Intercept)	-3.89	-6.19 – -2.05	0.005
time [day 15]	5.36	3.55 – 7.55	<0.001
time [day 60]	3.65	1.90 – 6.27	0.006
Random Effects			
<i>Groups</i>	<i>Predictors</i>	<i>Std.Dev</i>	<i>Corr.</i>
farm	(Intercept)	3.53	
	time [day 15]	2.79	-0.91
	time [day 60]	3.31	-0.99

* Wald's test *p* value

Table S7: Analysis of deviance table (type II likelihood ratio tests) of the full model relating the frequency of *M. bovis*-specific PCR positives to the variables time and season.

Fixed Effects				
<i>Predictors</i>	<i>npar</i>	<i>AIC</i>	X^2	<i>p</i>
time	2	708.25	20.5	3.60x10 ⁻⁵
season	1	689.85	0.061	0.81
Random Effects				
<i>Predictors</i>	<i>npar</i>	<i>AIC</i>	X^2	<i>p</i>
time farm	6	780.14	100.35	2.12x10 ⁻¹⁹

Legend: *npar* = number of parameters associated to the relative predictor; *AIC* = Akaike's information criterion value observed upon dropping of the relative predictor; X^2 = likelihood ratio test statistic; *p* = likelihood ratio test *p* value.

Table S8: Parameter estimates of the logistic mixed effects model analyzing the frequency of *M. dispar* isolation

Fixed Effects

<i>Predictors</i>	<i>Log-Odds</i>	<i>95% CI</i>	<i>p</i> *
(Intercept)	-2.27	-2.63 – -1.91	> 0.001

Random Effects

<i>Groups</i>	<i>Predictors</i>	<i>Std.Dev</i>	<i>Corr.</i>
farm	(Intercept)	1.28	
	time [day 15]	1.55	-0.89
	time [day 60]	1.79	-0.93

* Wald's test *p* value

Table S9: Analysis of deviance table (type II likelihood ratio tests) of the full model relating the frequency of isolation of *M. dispar* to the variables time and season.

Fixed Effects				
<i>Predictors</i>	<i>npar</i>	<i>AIC</i>	X^2	<i>p</i>
time	2	514.84	1.93	0.38
season	1	517.36	2.44	0.12
Random Effects				
<i>Predictors</i>	<i>npar</i>	<i>AIC</i>	X^2	<i>p</i>
time farm	6	517.77	12.86	0.04

Legend: *npar* = number of parameters associated to the relative predictor; *AIC* = Akaike's information criterion value observed upon dropping of the relative predictor; X^2 = likelihood ratio test statistic; *p* = likelihood ratio test *p* value.

Table S10: Parameter estimates of the logistic mixed effects model analyzing the frequency of *M. bovirhinis* isolation

Fixed Effects

<i>Predictors</i>	<i>Log-Odds</i>	<i>95% CI</i>	<i>p</i> *
(Intercept)	-1.29	-1.69 – 0.73	> 0.001
season [warm]	0.59	0.08 – 1.02	0.014

Random Effects

<i>Groups</i>	<i>Predictors</i>	<i>Std.Dev</i>	<i>Corr.</i>
farm	(Intercept)	0.90	
	time [day 15]	1.55	-0.51
	time [day 60]	2.50	-0.04

* Wald's test *p* value

Table S11: Analysis of deviance table (type II likelihood ratio tests) of the full model relating the frequency of isolation of *M. bovirhinis* to the variables time and season.

Fixed Effects				
<i>Predictors</i>	<i>npar</i>	<i>AIC</i>	X^2	<i>p</i>
time	2	858.51	5.84	0.054
season	1	860.64	5.96	0.014
Random Effects				
<i>Predictors</i>	<i>npar</i>	<i>AIC</i>	X^2	<i>p</i>
time farm	6	921.60	76.93	1.54x10 ⁻¹⁴

Legend: *npar* = number of parameters associated to the relative predictor; *AIC* = Akaike's information criterion value observed upon dropping of the relative predictor; X^2 = likelihood ratio test statistic; *p* = likelihood ratio test *p* value.

Table S12: Data structure description

Farm	batches	season*	Bulls analyzed at		
			0 days p.a.	15 days p.a.	60 days p.a.
I	1	warm	10	10	10
	2	warm	10	10	10
	3	warm	10	10	10
	4	warm	10	10	10
	5	warm	10	9	9
	6	warm	10	10	10
	7	cold	10	10	10
	8	cold	10	10	9
	9	cold	10	10	9
II	1	cold	10	10	10
III	1	cold	10	10	10
IV	1	cold	10	10	10
V	1	cold	10	10	10
VI	1	cold	10	10	10
VII	1	warm	10	10	10
	2	cold	10	10	10
VIII	1	warm	10	10	10
IX	1	cold	10	10	9
	2	cold	10	10	9
X	1	cold	10	10	10
XI	1	cold	10	10	10
XII	1	cold	10	10	10
XIII	1	warm	10	9	8
	2	warm	10	10	10

p.a.: *post* arrival

* Season describes the environmental conditions at the arrival: “cold” if the bulls were stabled between November and March, “warm” otherwise