

# Additional file 1: Description of the used within herd model of Map Spread

## A Equations for the within-herd dynamic

The description of the model we used corresponds to the within-herd model described in [2] and based on [11]. The set of parameters used are described in tables 1, 2 and 3.

### Notations

In the following equations,  $X_{(t,a)}$  is the number of animals in health state  $X$  at time  $t$  and age  $a$ . Age is given in weeks until the entry in the adult stage (from 1 to  $cal$ , with intermediary stages  $w$  for weaning age,  $y$  for young heifers age,  $h$  for heifers age and  $u$  for maximal age in the susceptible compartments), then by age group ( $P_1$  to  $P_{5+}$ ). Some variables can have a prefix: "b" for births in health states  $X$  ( $bX$ ), "N" for animals transiting between two health states ( $NX$ ) and "s" for exits (mortality and culling) ( $sX$ ). After entering the adult stage, flows corresponding to aging are noted using a superscript  $ng$ , whereas those remaining in the same age group are noted using a superscript  $sg$ .  $N_{(t,a)}$  is the number of animals of age  $a$  at time  $t$ . Average duration in health states are noted by  $v_X$ . The remaining terms used are defined when introduced.

### Equations for the updating of variables describing health states

In this section we introduce the equations for the updating of variables corresponding to the health states, for a given herd  $i$ .

#### Susceptible ( $S$ ) and No more Susceptible ( $R$ )

$$S_{(t+1,a=1)} = bS_{(t)} - NT_{(t+1,a)}$$

$$S_{(t+1,a \in [2;52])} = [S_{(t,a-1)} - sS_{(t,a-1)}] - NT_{(t+1,a)}$$

$$R_{(t+1,53)} = S_{(t,52)} - sS_{(t,52)}$$

$$R_{(t+1,a \in [54;cal])} = R_{(t,a-1)} - sR_{(t,a-1)}$$

—

$$R_{(t+1,P_1)} = R_{(t,P_1)}^{sg} - sR_{(t,P_1)}^{sg} + R_{(t,cal)} - sR_{(t,cal)}$$

$$R_{(t+1,P_i \in [P_2;P_4])} = R_{(t,P_i)}^{sg} - sR_{(t,P_i)}^{sg} + R_{(t,P_{i-1})}^{ng} - sR_{(t,P_{i-1})}^{ng}$$

$$R_{(t+1,P_{5+})} = R_{(t,P_{5+})} - sR_{(t,P_{5+})} + R_{(t,P_4)}^{ng} - sR_{(t,P_4)}^{ng}$$

### Transiently infected ( $T$ )

$$T_{(t+1,a=1)} = bT_{(t)} + NT_{(t+1,a)}$$

$$T_{(t+1,a \in [2;52])} = [T_{(t,a-1)} - sT_{(t,a-1)}] - NL_{(t+1,a)} + NT_{(t+1,a)}$$

$$T_{(t+1,a \in [53;cal])} = [T_{(t,a-1)} - sT_{(t,a-1)}] - NL_{(t+1,a)}$$

### Latently infected ( $L$ )

$$L_{(t+1,a=2)} = NL_{(t+1,a)}$$

$$L_{(t+1,a \in [3;h])} = [L_{(t,a-1)} - sL_{(t,a-1)}] + NL_{(t+1,a)}$$

$$L_{(t+1,a \in [h+1;cal])} = [L_{(t,a-1)} - sL_{(t,a-1)}] - NIs_{(t+1,a)} + NL_{(t+1,a)}$$

—

$$L_{(t+1,P_1)} = [L_{(t,P_1)}^{sg} - sL_{(t,P_1)}^{sg} + L_{(t,cal)} - sL_{(t,cal)} + T_{(t,cal)} - sT_{(t,cal)}] - NIs_{(t+1,P_1)}$$

$$L_{(t+1,P_i \in [P_2;P_4])} = [L_{(t,P_i)}^{sg} - sL_{(t,P_i)}^{sg} + L_{(t,P_{i-1})}^{ng} - sL_{(t,P_{i-1})}^{ng}] - NIs_{(t+1,P_i)}$$

$$L_{(t+1,P_{5+})} = [L_{(t,P_{5+})} - sL_{(t,P_{5+})} + L_{(t,P_4)}^{ng} - sL_{(t,P_4)}^{ng}] - NIs_{(t+1,P_{5+})}$$

### Moderate shedding ( $Is$ )

$$Is_{(t+1,a=h+1)} = NIs_{(t+1,a)}$$

$$Is_{(t+1,a \in [h+2;cal])} = [Is_{(t,a-1)} - sIs_{(t,a-1)}] - NIs_{(t+1,a)} + NIs_{(t+1,a)}$$

—

$$Is_{(t+1,P_1)} = [Is_{(t,P_1)}^{sg} - sIs_{(t,P_1)}^{sg} + Is_{(t,cal)} - sIs_{(t,cal)}] - NIs_{(t+1,P_1)} + NIs_{(t+1,P_1)}$$

$$Is_{(t+1,P_i \in [P_2;P_4])} = [Is_{(t,P_i)}^{sg} - sIs_{(t,P_i)}^{sg} + Is_{(t,P_{i-1})}^{ng} - sIs_{(t,P_{i-1})}^{ng}]$$

$$- NIs_{(t+1,P_i)} + NIs_{(t+1,P_i)}$$

$$Is_{(t+1,P_{5+})} = [Is_{(t,P_{5+})} - sIs_{(t,P_{5+})} + Is_{(t,P_4)}^{ng} - sIs_{(t,P_4)}^{ng}] - NIs_{(t+1,P_{5+})} + NIs_{(t+1,P_{5+})}$$

### High shedding and clinically affected ( $Ic$ )

$$Ic_{(t+1,a=h+2)} = NIs_{(t+1,a)}$$

$$Ic_{(t+1,a \in [h+3;cal])} = [Ic_{(t,a-1)} - sIc_{(t,a-1)}] + NIs_{(t+1,a)}$$

—

$$Ic_{(t+1,P_1)} = [Ic_{(t,P_1)}^{sg} - sIc_{(t,P_1)}^{sg} + Ic_{(t,cal)} - sIc_{(t,cal)}] + NIs_{(t+1,P_1)}$$

$$Ic_{(t+1,P_i \in [P_2;P_4])} = [Ic_{(t,P_i)}^{sg} - sIc_{(t,P_i)}^{sg} + Ic_{(t,P_{i-1})}^{ng} - sIc_{(t,P_{i-1})}^{ng}] + NIs_{(t+1,P_i)}$$

$$Ic_{(t+1,P_{5+})} = [Ic_{(t,P_{5+})} - sIc_{(t,P_{5+})} + Ic_{(t,P_4)}^{ng} - sIc_{(t,P_4)}^{ng}] + NIs_{(t+1,P_{5+})}$$

## Equations describing flows

New incoming (and outgoing) flows in each health states are mainly drawn using binomial laws.

### Births ( $bX$ )

At each time step  $t$ , births are calculated with regards to the health state of the dam. These births are then distributed into  $S$  and  $T$  states

$$bS_{(t)} = bS_{(t)}^R + bS_{(t)}^L + bS_{(t)}^{Is} + bS_{(t)}^{Ic} \quad \text{et} \quad bT_{(t)} = bT_{(t)}^L + bT_{(t)}^{Is} + bT_{(t)}^{Ic},$$

where  $bS_{(t)}^X$  et  $bT_{(t)}^X$  ( $X \in R, L, Is, Ic$ ) represent the number of births at time  $t$  from cows in health state  $X$ :

$$\begin{aligned} bS_{(t)}^R &\sim Bin \left( R_{(t,cal)} + \sum_{i=1}^{i=5} [R_{(t,P_i)}^{ng}]; 1 - \sigma_B \right) \\ bS_{(t)}^L &= nbV_{(t)}^L - bT_{(t)}^L, \quad bT_{(t)}^L \sim Bin \left( nbV_{(t)}^L; p_L * \varphi_{pLIs} \right) \\ bS_{(t)}^{Is} &= nbV_{(t)}^{Is} - bT_{(t)}^{Is}, \quad bT_{(t)}^{Is} \sim Bin \left( nbV_{(t)}^{Is}; p_{Is} * \varphi_{pLIs} \right) \\ bS_{(t)}^{Ic} &= nbV_{(t)}^{Ic} - bT_{(t)}^{Ic}, \quad bT_{(t)}^{Ic} \sim Bin \left( nbV_{(t)}^{Ic}; p_{Ic} * \varphi_{pIc} \right) \end{aligned}$$

In equations above,  $nbC_{(t)}^X$  is the number of female calves alive born at time  $t$ . It is obtained from  $nbV_{(t)}^X$ , the number of female calves born at time  $t$ , from cows in the health state  $X$ :

$$\begin{aligned} nbV_{(t)}^L &\sim Bin \left( L_{(t,cal)} + T_{(t,cal)} + \sum_{i=1}^{i=5} [L_{(t,P_i)}^{ng}]; 1 - \sigma_B \right) \\ nbV_{(t)}^{Is} &\sim Bin \left( Is_{(t,cal)} + \sum_{i=1}^{i=5} [Is_{(t,P_i)}^{ng}]; 1 - \sigma_B \right) \\ nbV_{(t)}^{Ic} &\sim Bin \left( Ic_{(t,cal)} + \sum_{i=1}^{i=5} [Ic_{(t,P_i)}^{ng}]; 1 - \sigma_B \right) \end{aligned}$$

We note that at the age of moving in the adult group ( $cal$ ), all the animals in the health state  $T$  enter the health state  $L$  ( $L_{(t,cal)} = L_{(t,cal)} + T_{(t,cal)}$ ).  $\varphi_{p_x}$  is a factor, varying from 100% to 0%, of decrease in probability of in utero transmission where  $x$  correspond to the dam giving birth in health states  $L$  and  $Is$  ( $\varphi_{pLIs}$ ) or  $Ic$  ( $\varphi_{pIc}$ ).

### Change in age group ( $X^x$ )

$$\begin{aligned} X_{(t,P_i)} &= X_{(t,P_i)}^{sg} + X_{(t,P_i)}^{ng} \\ X_{(t,P_i)}^{ng} &\sim Bin \left( X_{(t,P_i)}; \frac{1}{\tau_{aa}} \right) \quad ; \quad \text{where } X = \{R, L, Is, Ic\}, \end{aligned}$$

and  $\tau_{aa}$  is the average time spent in each of age group  $P_1$  to  $P_4$ .

## Exits ( $sX$ )

The mortality of calves during the first week of life is applied at birth and defined above in the section concerning births.

From age 1 to  $cal$ , mortality and culling rates  $\sigma_x$  are defined as:

- $a \in \{1; 2\} \rightarrow \sigma_x = \sigma_{c1}$
- $a \in [3; w] \rightarrow \sigma_x = \sigma_{c2}$
- $a \in [w + 1; cal] \rightarrow \sigma_x = \sigma_{c3}$

Then, exits following death and culling write as:

for  $a = 1$  :

$$sX_{(t,a)} \sim Bin\left(X_{(t,a)}; \sigma_x\right), \quad \text{where } X = \{S, T\},$$

for  $a \in [2; 4]$  :

$$sX_{(t,a)} = sale + death, \quad \text{where } X = \{S, T, L\}$$

$$sale \sim Bin\left(X_{(t,a)}; \sigma_m\right)$$

$$death \sim Bin\left(X_{(t,a)} - sale; \sigma_x\right)$$

for  $a \in [5; cal - 11]$  :

$$sX_{(t,a)} \sim Bin\left(X_{(t,a)}; \sigma_x\right) \quad ; \quad X = \{S, R, T, L\}$$

for  $a = cal - 10$ , we consider management by heifers (safe management, keep all female calves):

$$Cows_{(t)} = \sum_{i=1}^5 R_{(t,P_i)} + \sum_{i=1}^5 L_{(t,P_i)} + \sum_{i=1}^5 Is_{(t,P_i)} + \sum_{i=1}^5 Ic_{(t,P_i)}$$

$$Heifers_{(t)} = \sum_{a=1}^u S_{(t,a)} + \sum_{a=u+1}^{cal} R_{(t,a)} + \sum_{a=1}^{cal} T_{(t,a)} + \sum_{a=1}^{cal} L_{(t,a)} + \sum_{a=1}^{cal} Is_{(t,a)} + \sum_{a=1}^{cal} Ic_{(t,a)}$$

If the number of heifers,  $Heifers_{(t)}$ , is greater than  $K_g$  or that the number of cows,  $Cows_{(t)}$ , is greater than  $K_v$ , we consider the sale of heifers:

$$sX_{(t,a)} = sale + death \quad ; \quad X = \{R, T, L, Is, Ic\}$$

$$sale \sim Bin\left(X_{(t,a)}; \exp\left(-\sigma_h \cdot (Cows_{(t)}/K_v)^6\right) \cdot ((Heifers_{(t)}/K_g)^6)\right)$$

$$death \sim Bin\left(X_{(t,a)} - sale; \sigma_x\right)$$

where  $K_v$  is the capacity of the holding in number of cows (see Table 1) and  $K_g = \sigma_P * K_v * (cal - h)$  is the capacity of the holding in number of heifers.

Otherwise, we do not consider the sale of heifers:

$$sX_{(t,a)} \sim Bin\left(X_{(t,a)}; \sigma_x\right)$$

for  $a \in [cal - 9; cal]$ :

$$sX_{(t,a)} = Bin\left(X_{(t,a)}; \sigma_x\right)$$

for  $a \in [P_1; P_{5+}]$ :

$$sX_{(t,P_i)}^x \sim Bin\left(X_{(t,P_i)}^x; \sigma_{P_i}\right), \quad \text{where } X = \{R, L, Is\},$$

$$sIc_{(t,P_i)}^x \sim Bin\left(Ic_{(t,P_i)}^x; \frac{1}{v_{Ic}}\right)$$

where  $v_{Ic}$  corresponds to the average time spent in the health state  $Ic$ .

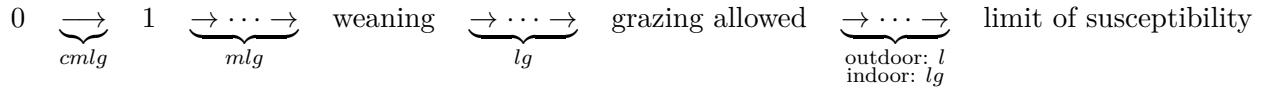
### New infections ( $S \rightarrow T$ , except for *in-utero* transmission)

We have:

$$nT_{(t+1,a)} = inf_{(t+1,a)}^c + inf_{(t+1,a)}^m + inf_{(t+1,a)}^l + inf_{(t+1,a)}^g$$

Superscripts correspond to different possible routes of transmission, respectively colostrum ( $c$ ), milk ( $m$ ), local environment ( $l$ ) and global environment ( $g$ ). New infections by *in-utero* transmission are accounted for through births.

By age, the possible routes of infection are:



### Transmission through colostrum

It is considered that calves drink colostrum from their mothers during the first three days before drinking milk:

$$inf_{(t+1,a=1)}^c = \sum_1^{bS_{(t)}^{Is}} \left[ Bern\left(1 - \exp\left(-\frac{\beta_l q_c^{Is}}{\alpha}\right)\right) \right] + \sum_1^{bS_{(t)}^{Ic}} \left[ Bern\left(1 - \exp\left(-\frac{\beta_l q_c^{Ic}}{\alpha}\right)\right) \right],$$

with

$$q_c^{Is} \sim Bern(sh_{Is}) \times \left[ 3 \times b \left( 10^5 \cdot Beta(8; 8) + 1 + 10^3 \cdot Beta(1; 25) \right) \right] * \varphi Milk_{Is},$$

$$q_c^{Ic} \sim Bern(sh_{Ic}) \times \left[ 3 \times b \left( 10^5 \cdot Beta(8; 8) + 10^{(3+10 \cdot Beta(50; 200))} \right) \right] * \varphi Milk_{Ic}.$$

Where  $\varphi Milk_x$  is a factor, varying from 100% to 0%, of decrease in quantity of bacteria shed in colostrum and milk by an animal in health state  $x$  (here  $x$  could be  $Is$  or  $Ic$ )

## Transmission through milk

Regarding the age, we have:

$$a = 1 : \quad inf_{(t+1,a)}^m \sim Bin \left( bS; 1 - \exp \left( -\frac{\beta_l q_l \frac{4}{7}}{\alpha} \right) \right),$$

$$a \in \{2; w\} : \quad inf_{(t+1,a)}^m \sim Bin \left( [S_{(t,a-1)} - sS_{(t,a-1)}]; \left[ 1 - \exp \left( -e^{(-\gamma(a-1))} \cdot \frac{\beta_l q_l}{\alpha} \right) \right] \right),$$

with

$$q_l = \frac{7 \times d \times (Q_{I_s(t)}^{milk} + Q_{I_c(t)}^{milk})}{MilkTot_{(t)}},$$

where

$$Q_{I_s(t)}^{milk} = 7 \times \varepsilon \times g_{I_s} \times \left( \sum_1^{nbExcr_{(t)}^{I_s}} (10^5 \cdot Beta(8; 8)) + \sum_1^{nbLac_{(t)}^{I_s}} (1 + 10^3 \cdot Beta(1; 25)) \right) * \varphi Milk_{I_s},$$

$$Q_{I_c(t)}^{milk} = 7 \times \varepsilon \times g_{I_c} \times \left( \sum_1^{nbExcr_{(t)}^{I_c}} (10^5 \cdot Beta(8; 8)) + \sum_1^{nbLac_{(t)}^{I_c}} 10^{(3+10 \cdot Beta(50; 200))} \right) * \varphi Milk_{I_c},$$

$$MilkTot_{(t)} = 7 \times \varepsilon \times (nbLac_{(t)}^R + g_L \cdot nbLac_{(t)}^L + g_{I_s} \cdot nbLac_{(t)}^{I_s} + g_{I_c} \cdot nbLac_{(t)}^{I_c}).$$

with

$$nbExcr_{(t)}^{I_s} \sim Bin (nbLac_{(t)}^{I_s}; sh_{I_s}),$$

$$nbExcr_{(t)}^{I_c} \sim Bin (nbLac_{(t)}^{I_c}; sh_{I_c}),$$

$$nbLac_{(t)}^R \sim Bin \left( \sum_{i=1}^{i=5} R_{(t,P_i)}, prop \right),$$

$$nbLac_{(t)}^L \sim Bin \left( \sum_{i=1}^{i=5} L_{(t,P_i)}, prop \right),$$

$$nbLac_{(t)}^{I_s} \sim Bin \left( \sum_{i=1}^{i=5} Is_{(t,P_i)}, prop \right),$$

$$nbLac_{(t)}^{I_c} \sim Bin \left( \sum_{i=1}^{i=5} Ic_{(t,P_i)}, prop \right).$$

## Local transmission (in collective pens, $a \in [1; u]$ )

$$inf_{(t+1,a)}^l \sim Bin (S_{(t,a-1)} - sS_{(t,a-1)}; p_{inf_{(t+1,a)}^i}),$$

where

during housing period:

$$p_{inf(t+1,a)}^i = 1 - \exp\left(-e^{(-\gamma(a-1))} \cdot \frac{\beta_c E_{(t+1)}^i}{\alpha N_{(t+1)}^i}\right), \text{ and}$$

during grazing period, regarding the age,

$$a \in [1; 26] : p_{inf(t+1,a)}^i = 1 - \exp\left(-e^{(-\gamma(a-1))} \cdot \frac{\beta_c E_{(t+1)}^i}{\alpha N_{(t+1)}^i}\right),$$

$$a \in [27; u] : p_{inf(t+1,a)}^i = 1 - \exp\left(-e^{(-\gamma(a-1))} \cdot \frac{\beta_o E_{(t+1)}^i}{\alpha N_{(t+1)}^i}\right).$$

$N_{(t+1)}^i$  is the total number of animals in environment  $i$ , across all health states, and  $E^i$  represents the quantity of bacteria in the environment, with  $i$  corresponding to the specific area.

### Global transmission

This occurs up to the age allowing to go to the pasture (26 weeks - 6 months) during the grazing period, and up to the age limit for sensitivity ( $u$ ) during the housing period:

$$inf_{(t+1,a)}^g \sim Bin\left(S_{(t,a)} - sS_{(t,a)}; p_{inf(t+1,a)}^g\right),$$

where

$$p_{inf(t+1,a)}^g = 1 - \exp\left(-\exp[-h(a-1)] \cdot \frac{\beta_g E_{(t+1)}^g}{\alpha N_{(t+1)}^g}\right), \text{ with } E_{(t+1)}^g = \sum_{i=1}^{i=5} E_{(t+1)}^{INT^i}$$

$E^{INT^i}$  represents the quantity of bacteria in the environment where  $INT^i$  corresponds to a specific area (see the section below about the dynamics of bacteria in the environments).

### New latently infected ( $T \rightarrow L$ )

For  $a < cal$ :

$$NL_{(t+1,a \in [2; cal-1])} \sim Bin\left(T_{(t,a)} - sT_{(t,a)}; \frac{1}{v_T}\right).$$

After age  $cal$ , there are no more animals in  $T$  state:

$$NL_{(t+1,P_1)} = T_{(t,cal)} - sT_{(t,cal)}.$$

### New subclinically infected ( $\mathbf{L} \rightarrow \mathbf{Is}$ )

For heifers:

$$NIS_{(t+1, a \in [h+1; cal])} \sim Bin \left( L_{(t, a)} - sL_{(t, a)}; \frac{1}{v_L} \right).$$

For cows:

$$NIS_{(t+1, P_i)} \sim Bin \left( n; \frac{1}{v_L} \right),$$

with, regarding the age,

$$\begin{aligned} P_1 \rightarrow n &= \left[ L_{(t, P_1)}^{sg} - sL_{(t, P_1)}^{sg} + L_{(t, cal)} - sL_{(t, cal)} + T_{(t, cal)} - sT_{(t, cal)} \right], \\ \{P_2; P_4\} \rightarrow n &= \left[ L_{(t, P_i)}^{sg} - sL_{(t, P_i)}^{sg} + L_{(t, P_{i-1})}^{ng} - sL_{(t, P_{i-1})}^{ng} \right], \\ P_{5+} \rightarrow n &= \left[ L_{(t, P_{5+})} - sL_{(t, P_{5+})} + L_{(t, P_4)}^{ng} - sL_{(t, P_4)}^{ng} \right]. \end{aligned}$$

### New clinically infected ( $\mathbf{Is} \rightarrow \mathbf{Ic}$ )

For heifers:

$$NIC_{(t+1, a \in [h+1; cal])} \sim Bin \left( IS_{(t, a)} - sIS_{(t, a)}; \frac{1}{v_{Is}} \right).$$

For cows:

$$NIC_{(t+1, P_i)} \sim Bin \left( n; \frac{1}{v_{Is}} \right),$$

with

$$\begin{aligned} P_1 \rightarrow n &= \left[ IS_{(t, P_1)}^{sg} - sIS_{(t, P_1)}^{sg} + IS_{(t, cal)} - sIS_{(t, cal)} \right], \\ \{P_2; P_4\} \rightarrow n &= \left[ IS_{(t, P_i)}^{sg} - sIS_{(t, P_i)}^{sg} + IS_{(t, P_{i-1})}^{ng} - sIS_{(t, P_{i-1})}^{ng} \right], \\ P_{5+} \rightarrow n &= \left[ IS_{(t, P_{5+})} - sIS_{(t, P_{5+})} + IS_{(t, P_4)}^{ng} - sIS_{(t, P_4)}^{ng} \right]. \end{aligned}$$

### New resistant ( $\mathbf{S} \rightarrow \mathbf{R}$ )

At age  $u$ , the transition from compartment S to compartment R is done in a deterministic way.

### Dynamics of bacteria in the environments ( $\mathbf{E}$ )

The composition of the environments according to the season is the following:

$$\text{Grazing : } a = \overbrace{1 \dots \dots \dots w}^{\text{INT1}} \overbrace{\dots \dots \dots 26}^{\text{INT2}} \overbrace{\dots \dots \dots h}^{\text{EXT1+EXT2}} \overbrace{\dots \dots \dots cal}^{\text{EXT3}}$$



$$\text{Housing : } a = \overbrace{1 \dots \dots \dots w}^{\text{INT1}} \overbrace{\dots \dots \dots y}^{\text{INT2}} \overbrace{\dots \dots \dots h}^{\text{INT3}} \overbrace{\dots \dots \dots cal}^{\text{INT4}}$$

Dynamics of bacteria in the environments ( $E$ ) are defined below:

$E_{(t+1)}^{\text{INT1}} = E_{(t)}^{\text{INT1}} \cdot (1 - \mu_g^{\text{int}}) + Q_{(t)}^{\text{TNS}}$ . If pens are empty, it becomes  $E_{(t+1)}^{\text{INT1}} = E_{(t+1)}^{\text{INT1}} \cdot (1 - \mu_{cp})$ , where  $Q$  represents the quantity of bacteria shed.

During grazing period, we have:

$$E_{(t+1)}^{\text{INT2}} = E_{(t)}^{\text{INT2}} \cdot (1 - \mu_g^{\text{int}}) + Q_{(t)}^{\text{Ts1}}$$

$$\text{If } \sum_{a=w+1}^{a=26} SR_{t,a} + \sum_{a=w+1}^{a=26} T_{t,a} + \sum_{a=w+1}^{a=26} L_{t,a} == 0 \text{ then } E_{(t+1)}^{\text{INT2}} = E_{(t+1)}^{\text{INT2}} \cdot (1 - \mu_{cp})$$

$$E_{(t+1)}^{\text{INT3}} = E_{(t)}^{\text{INT3}} \cdot (1 - \mu_g^{\text{int}})$$

$$E_{(t+1)}^{\text{INT4}} = E_{(t)}^{\text{INT4}} \cdot (1 - \mu_g^{\text{int}})$$

$$E_{(t+1)}^{\text{INT5}} = E_{(t)}^{\text{INT5}} \cdot (1 - \mu_g^{\text{int}})$$

$$E_{(t+1)}^{\text{EXT1}} = E_{(t)}^{\text{EXT1}} \cdot (1 - \mu_g^{\text{ext}}) + Q_{(t)}^{\text{Ts2}}$$

$$E_{(t+1)}^{\text{EXT2}} = E_{(t)}^{\text{EXT2}} \cdot (1 - \mu_g^{\text{ext}}) + Q_{(t)}^{\text{TY}}$$

$$E_{(t+1)}^{\text{EXT3}} = E_{(t)}^{\text{EXT3}} \cdot (1 - \mu_g^{\text{ext}}) + Q_{(t)}^{\text{TH}} + Q_{(t)}^{\text{ISH}} + Q_{(t)}^{\text{ICH}}.$$

During housing period, we have:

$$E_{(t+1)}^{\text{INT2}} = E_{(t)}^{\text{INT2}} \cdot (1 - \mu_g^{\text{int}}) + Q_{(t)}^{\text{Ts1}} + Q_{(t)}^{\text{Ts2}}$$

$$\text{If } \sum_{a=w+1}^{a=52} SR_{t,a} + \sum_{a=w+1}^{a=52} T_{t,a} + \sum_{a=w+1}^{a=52} L_{t,a} == 0 \text{ then } E_{(t+1)}^{\text{INT2}} = E_{(t+1)}^{\text{INT2}} \cdot (1 - \mu_{cp})$$

$$E_{(t+1)}^{\text{INT3}} = E_{(t)}^{\text{INT3}} \cdot (1 - \mu_g^{\text{int}}) + Q_{(t)}^{\text{TY}}$$

$$E_{(t+1)}^{\text{INT4}} = E_{(t)}^{\text{INT4}} \cdot (1 - \mu_g^{\text{int}}) + Q_{(t)}^{\text{TH}} + Q_{(t)}^{\text{ISH}} + Q_{(t)}^{\text{ICH}}$$

$$E_{(t+1)}^{\text{INT5}} = E_{(t)}^{\text{INT5}} \cdot (1 - \mu_g^{\text{int}}) + Q_{(t)}^{\text{IS}} + Q_{(t)}^{\text{IC}}$$

$$E_{(t+1)}^{\text{EXT1}} = 0, \quad E_{(t+1)}^{\text{EXT2}} = 0, \quad E_{(t+1)}^{\text{EXT3}} = 0.$$

Shed quantities of bacteria are defined, regarding the health states and the age, by:

unweaned calves T:

$$Q_{(t)}^{\text{TNS}} = \sum_{a=1}^{a=w} \left[ 7 \cdot f_1 \cdot 10^6 \cdot \sum_{T_{t,a}} \text{Beta}(8.8; 19) \right] * \varphi faeces_T,$$

weaned calves T, without access to grazing:

$$Q_{(t)}^{\text{Ts1}} = \sum_{a=w+1}^{a=26} \left[ 7 \cdot f_2 \cdot 10^6 \cdot \sum_{T_{t,a}} \text{Beta}(8.8; 19) \right] * \varphi faeces_T,$$

weaned calves T, with access to grazing:

$$Q_{(t)}^{Ts2} = \sum_{a=27}^{a=y} \left[ 7.f_2.10^6 \cdot \sum^{T_{(t,a)}} \text{Beta}(8.8; 19) \right] * \varphi faeces_T,$$

young heifers T:

$$Q_{(t)}^{TY} = \sum_{a=y+1}^{a=h} \left[ 7.f_Y.10^6 \cdot \sum^{T_{(t,a)}} \text{Beta}(8.8; 19) \right] * \varphi faeces_T,$$

heifers T:

$$Q_{(t)}^{TH} = \sum_{a=h+1}^{a=cal} \left[ 7.f_A.10^6 \cdot \sum^{T_{(t,a)}} \text{Beta}(8.8; 19) \right] * \varphi faeces_T,$$

heifers Is:

$$\text{if } \sum_{a=h+1}^{a=cal} (Is_{(t,a)}) > 0 : Q_{(t)}^{ISH} = \sum_{a=h+1}^{a=cal} \left[ 7.f_A.10^{(4+10 \times \sum^{Is_{(t,a)}} \text{Beta}(2.65; 17))} \right] * \varphi faeces_{Is}$$

$$\text{else } Q_{(t)}^{ISH} = 0,$$

cows Is:

$$\text{if } \sum_{i=1}^{i=5} (Is_{(t,P_i)}) > 0 : Q_{(t)}^{IS} = \sum_{i=1}^{i=5} \left[ 7.f_A.10^{(4+10 \times \sum^{Is_{(t,P_i)}} \text{Beta}(2.65; 17))} \right] * \varphi faeces_{Is}$$

$$\text{else } Q_{(t)}^{IS} = 0,$$

heifers Ic:

$$\text{if } \sum_{a=h+1}^{a=cal} (Ic_{(t,a)}) > 0 : Q_{(t)}^{ICH} = \sum_{a=h+1}^{a=cal} \left[ 7.f_A.10^{(8+10 \times \sum^{Ic_{(t,a)}} \text{Beta}(2; 17))} \right] * \varphi faeces_{Ic}$$

$$\text{else } Q_{(t)}^{ICH} = 0,$$

cows Ic:

$$\text{if } \sum_{i=1}^{i=5} (Ic_{(t,P_i)}) > 0 : Q_{(t)}^{IC} = \sum_{i=1}^{i=5} \left[ 7.f_A.10^{(8+10 \times \sum^{Ic_{(t,P_i)}} \text{Beta}(2; 17))} \right] * \varphi faeces_{Ic}$$

$$\text{else } Q_{(t)}^{IC} = 0.$$

$\varphi faeces_x$ , in the equation above is a factor of decrease in quantity of bacteria shed by an animal in health state  $x$ .  $x$  could be  $T$ ,  $Is$  or  $Ic$ .

## B Parameters related to population dynamics

**Table 1:** Parameters for management and population dynamics used in the *Mycobacterium avium subsp. paratuberculosis* (*Map*) infection dynamics model within a structured dairy herd (reproduced from [2], Table 1).

Notation	Value	Definition	Source
$\sigma_B$	0.07	Mortality rate of calves at birth	*, [17]
$\sigma_m$	0.206	Exit rate of male calves, weeks 2 to 4 (per week)	
$\sigma_{c1}$	0.015	Death rate of female calves, weeks 1 and 2 (per week)	[17]
$\sigma_{c2}$	0.0035	Death rate of female calves, weeks 3 to weaning (per week)	[9]
$\sigma_{c3}$	0.00019	Death rate of heifers from weaning to entry in adult group (per week)	†
$\sigma_h$	0.011	Sale rate of blackcol heifers 10 weeks before 1st calving	†
$\sigma_{Ai}$	0.27,0.25,0.31, 0.31,0.62	Yearly culling rate of cows in adult group $i$ : 1, 2, 3, 4 and above 5 respectively (%)	*, [1]
$w$	10	Weaning age (weeks)	[12]
$y$	52	Age when entering the young heifer group (weeks)	
$h$	91	Age when entering the heifer group (weeks)	*
$cal$	130	Age when entering the adult group (weeks)	*, †
$\tau_{aa}$	56.3	Mean time spent in adult age groups 1 to 4 (weeks)	*, †
$b$	5	Quantity of colostrum fed to calves (L/day for 3 days)	†
$d$	7	Quantity of milk fed to calves after 3 days (L/day/calf)	†
$prop$	0.85	Proportion of lactating cows	*
$\varepsilon$	25	Quantity of milk or colostrum produced (L/day/cow)	*
$f_1$	0.5	Quantity of faeces produced by a non-weaned calf (kg/day)	†
$f_2$	5.5	Quantity of faeces produced by a weaned calf (kg/day)	†
$f_Y$	10	Quantity of faeces produced by a heifer (kg/day)	†
$f_A$	30	Quantity of faeces produced by a cow (kg/day)	†
$Graz$	[14 – 26]	Grazing period (1 being the first week of the year)	†
$K_c$	110	Number of cows above which the heifer selling rate increases	

\* Agricultural statistics.

† Based on expert opinion.

## C Parameters related to infection dynamics

**Table 2:** Parameters for infection and transmission used in the *Mycobacterium avium subsp. paratuberculosis* (*Map*) infection dynamics model within a structublackcol dairy herd (reproduced from [2] - Table 2).

Notation	Value	Definition	Source
$p_X$	$p_L = 0.149$	Probability of <i>in utero</i> transmission for cow in health state $X$ $X =$ latently infected (L)	[4, 26]
	$p_{I_s} = 0.149$	$X =$ subclinically infected (Is)	
	$p_{I_c} = 0.65$	$X =$ clinically affected (Ic)	
$u$	52	Maximal age in the susceptible compartment (weeks)	[6, 23]
$\gamma$	0.1	Susceptibility follows an exponential decrease : $exp(-\gamma(\text{age} - 1))$	[27]
$v_X$		Mean time spent in health state $X$ (weeks)	
	$v_T = 25$	$X =$ transiently infectious (T)	[21]
	$v_L = 52$	$X =$ latently infected (L)	[16, 14]
	$v_{I_s} = 104$	$X =$ subclinically infected (Is)	[13]
$sh_X$	$v_{I_c} = 26$	$X =$ clinically affected (Ic)	†
		Probability of shedding in colostrum or milk for a cow in health state $X$	[20, 19]
	$sh_L = 0$	$X =$ latently infected (L)	
	$sh_{I_s} = 0.4$	$X =$ subclinically infected (Is)	
	$sh_{I_c} = 0.9$	$X =$ clinically affected (Ic)	
$\alpha$	$10^6$	<i>Map</i> infectious dose	[3]
$\beta_l$	$5 \times 10^{-4} \times 7$	Transmission rate if ingestion of an infectious dose (per week)	‡
$\beta_c$	$5 \times 10^{-5} \times 7$	Transmission rate if one infectious dose is present in the local environment (per week)	[21]
$\beta_g$	$9.5 \times 10^{-7} \times 7$	Transmission rate if one infectious dose is present in the global environment (per week)	[21]
$\beta_o$	$5 \times 10^{-6} \times 7$	Transmission rate if one infectious dose is present in the pasture (per week)	‡
$g_X$		Decrease in milk production for cattle in health state $X$ (per week)	[15]
	$g_{I_s} = 1 - 0.11$	$X =$ subclinically infected (Is)	
	$g_{I_c} = 1 - 0.25$	$X =$ clinically affected (Ic)	
$\mu_k$		Removal rate of <i>Map</i> from environment k	[7, 24]
	$\mu_g^{int} = 0.4$	all the environments (per week)	
	$\mu_g^{ext} = 1/14$	all the environments (per week)	
	$\mu_{cp} = 0.17$	collective pens (when empty)	

† Expert opinions.

‡ Parameters' values are assumed.

**Table 3:** Summary of published data and assumed distributions of the quantities of *Mycobacterium avium* subsp. *paratuberculosis* (*Map*) shed, depending on the health state (X) and the route of transmission (r) in the *Map* infection dynamics model within a structblackcol dairy herd (reproduced from [2] - Table 3).

Route of transmission (r)	Health state (X)		Literature		Assumed distributions	
	Minimal value	Maximal value	Minimal value	Maximal value	Mean value	Source
Map direct shedding in milk and colostrum (Map/L)	Subclinically infected	$2.2 \times 10^4$	$8.8 \times 10^4$	$5 \times 10^4$	$5 \times 10^4$	$10^5 \times \text{Beta}(8; 8)$
	Clinically affected	-	-	$5 \times 10^4$	$5 \times 10^4$	$10^5 \times \text{Beta}(8; 8)$
Map indirect shedding in milk and colostrum (faecal contamination) (Map/L)	Subclinically infected	0	$2 \times 10^{10}$	40	40	$1 + 10^3 \times \text{Beta}(1; 25)$
	Clinically affected	700	$2 \times 10^{10}$	$14 \times 10^4$	$14 \times 10^4$	$10^{(3+10 \times \text{Beta}(50; 200))}$
Map shedding in faeces (Map/kg)	Transiently infectious	$6 \times 10^4$	$6.3 \times 10^5$	$3 \times 10^5$	$3 \times 10^5$	$10^6 \times \text{Beta}(8.8; 19)$
	Subclinically infected	$10^4$	$10^{15}$	$2.6 \times 10^6$	$2.6 \times 10^6$	$10^{(4+10 \times \text{Beta}(2.65; 17))}$
	Clinically affected	$10^8$	$10^{15}$	$10^{10}$	$10^{10}$	$10^{(8+10 \times \text{Beta}(2; 17))}$

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