

Structuring the mess: Disentangling disease models for cattle populations



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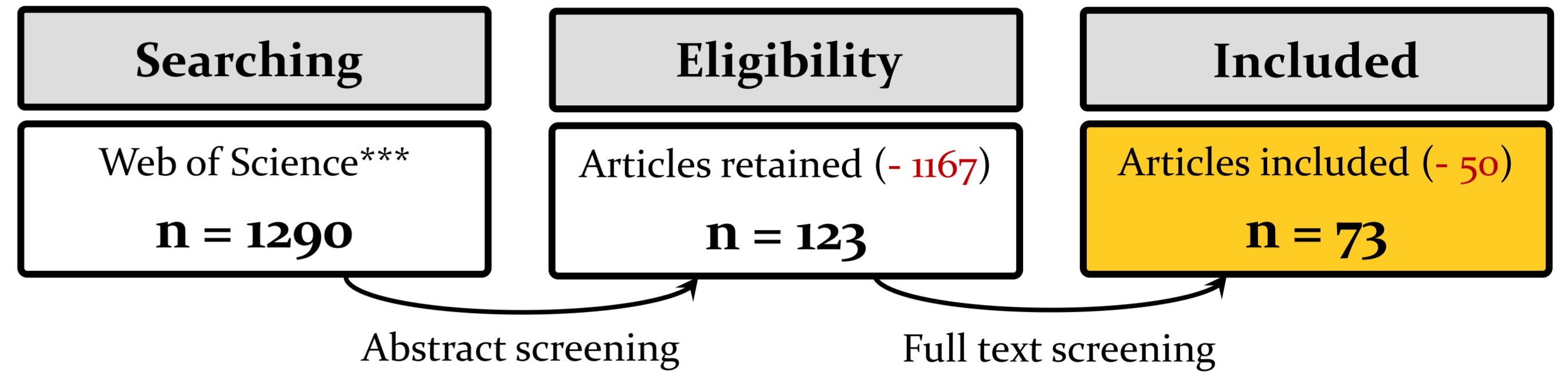
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Objective & Material

Can we structure key characteristics of cattle disease models that have been developed to assist with animal health decision-making?

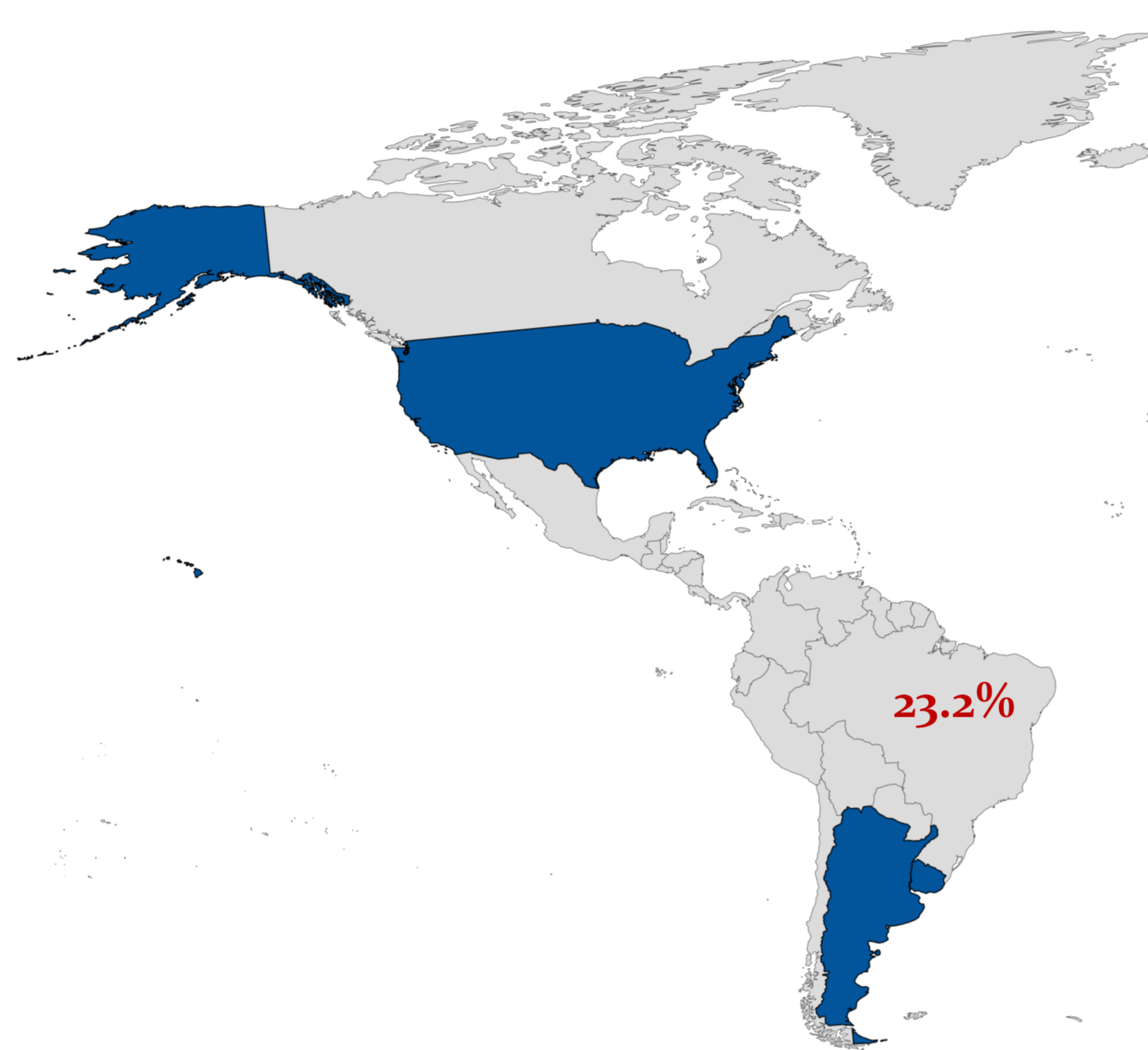
Systematic literature review



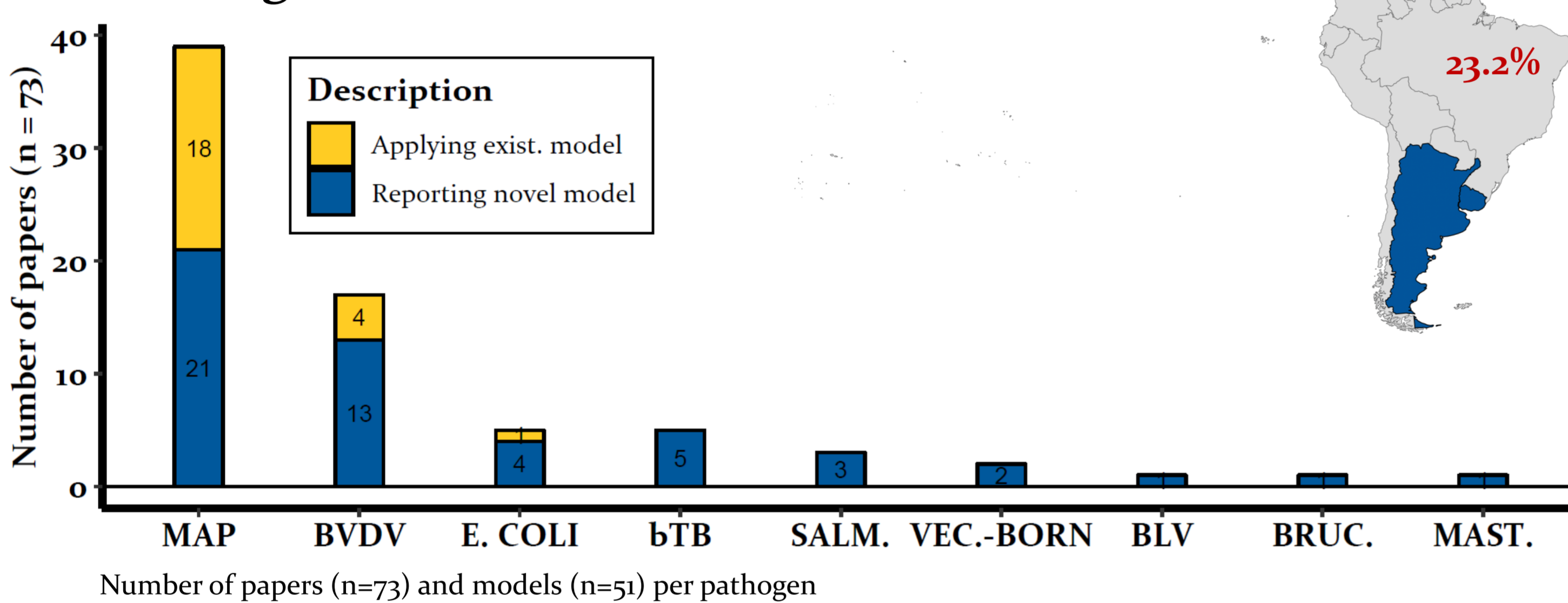
General Model Characteristics

Journals

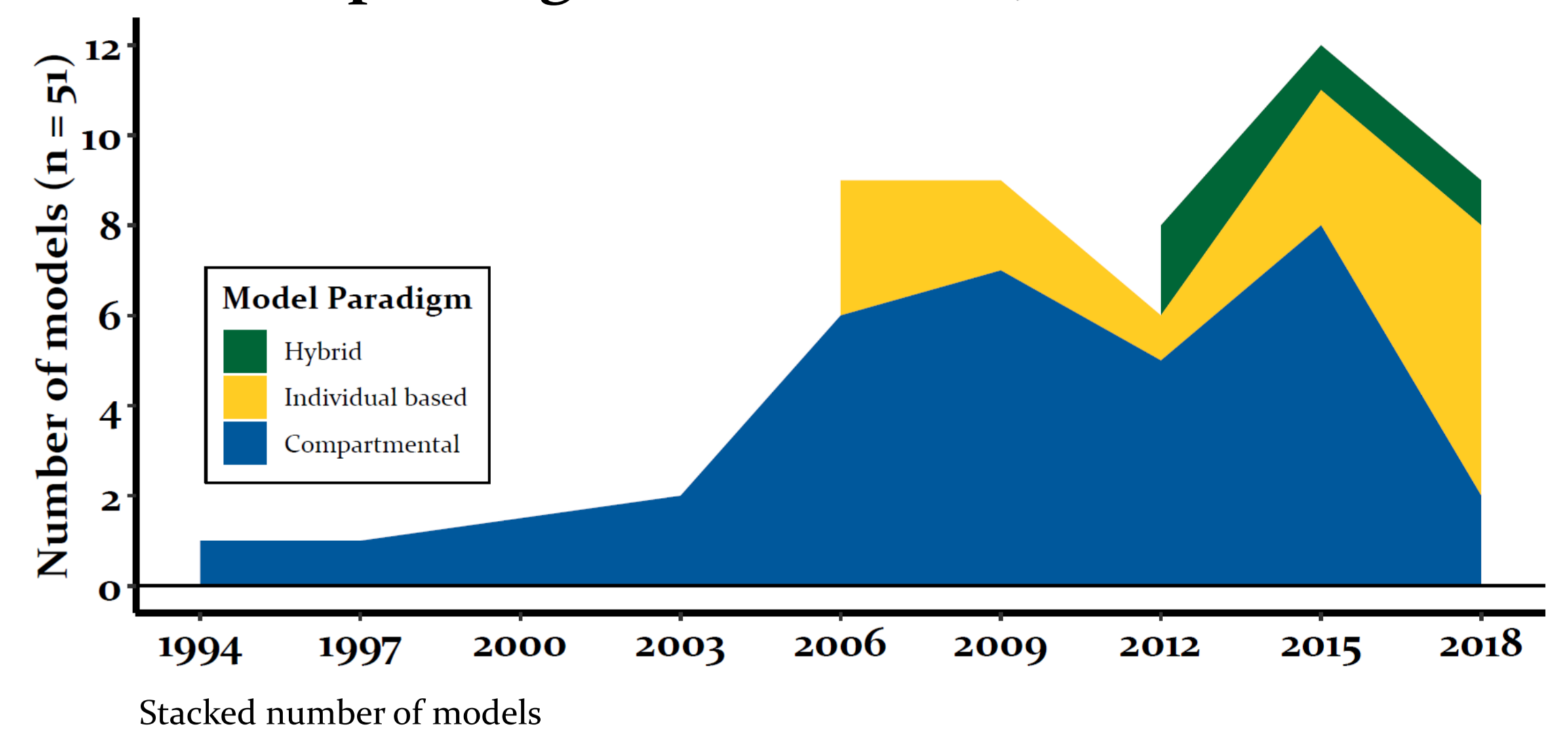
Prev. Vet. Med.:	39.2%
J. Theor. Biol.:	11.8%
Vet. Research:	11.8%
Other:	37.3%



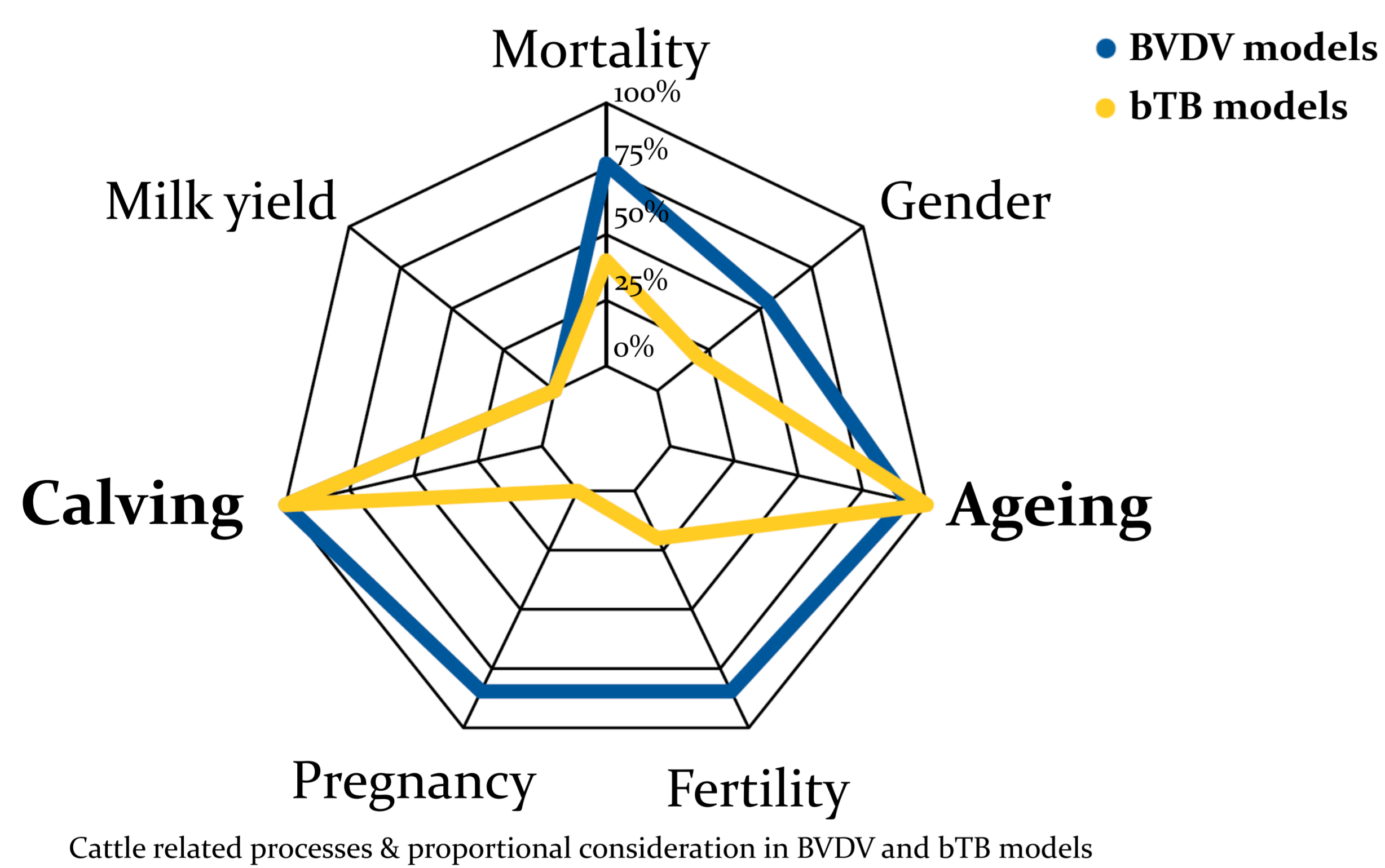
Pathogen distribution



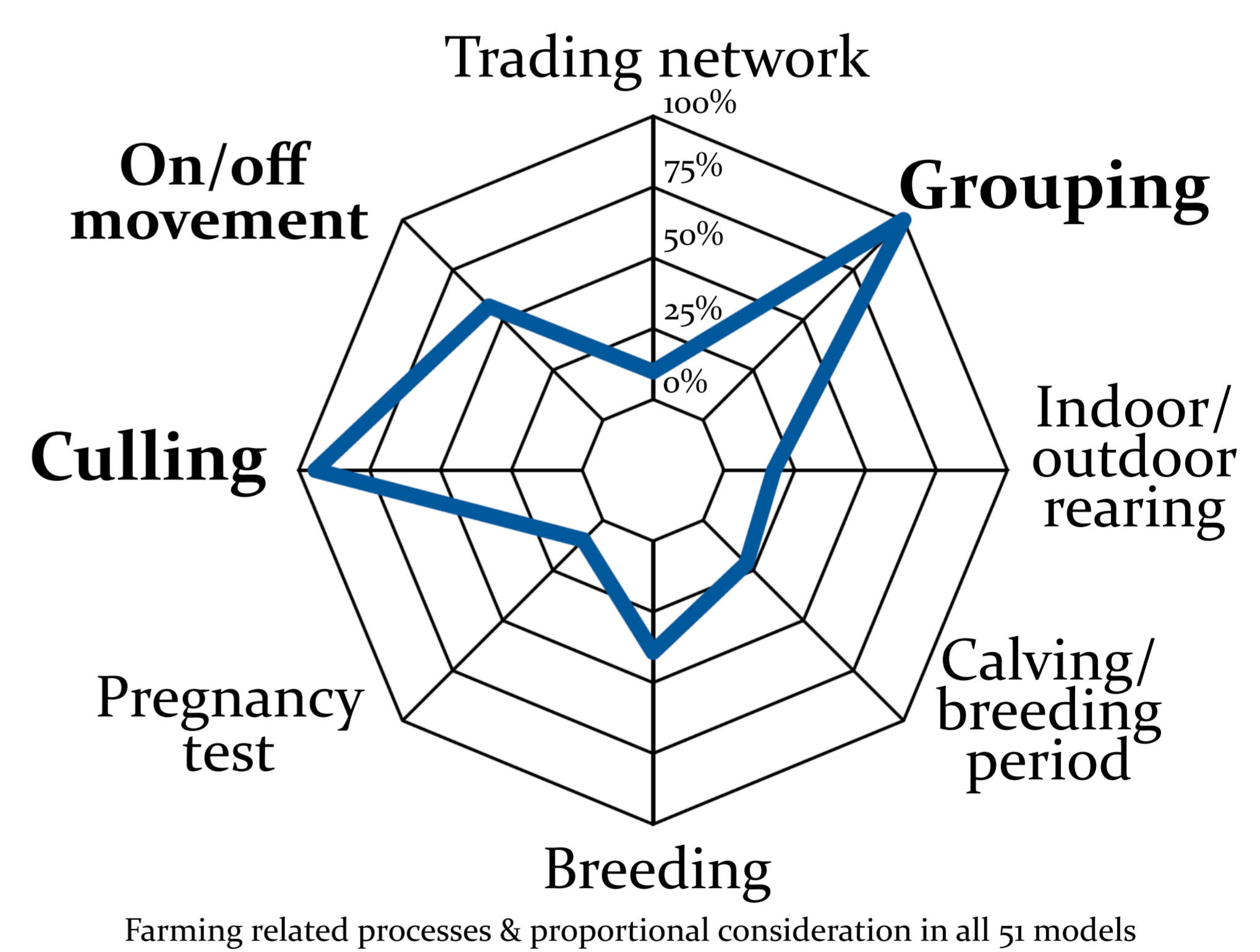
Model paradigms



Biological Processes



Farmer's Processes



Direct Disease Transmission

Individual probability (frequency [N] | density [1] dependent):

$$P_{inf} = 1 - \exp\left(-\beta_{I(x)} \frac{N_{I(x)}}{N|1} + \dots\right)$$

35%

$\beta_{I(x)}$ = Transmission coefficient for infectious state x
 $N_{I(x)}$ = Number of infected animals in state x
 N = Number of all animals

Individual probability (Reed-Frost):

$$P_{inf} = 1 - \left(1 - \frac{k \cdot s}{N}\right)^{N_{I(x)}}$$

28%

k = Number of effective contacts
 s = Susceptibility of each animal
 $N_{I(x)}$ = Number of infected animals in state x
 N = Number of all animals

Cohort rate of transmission:

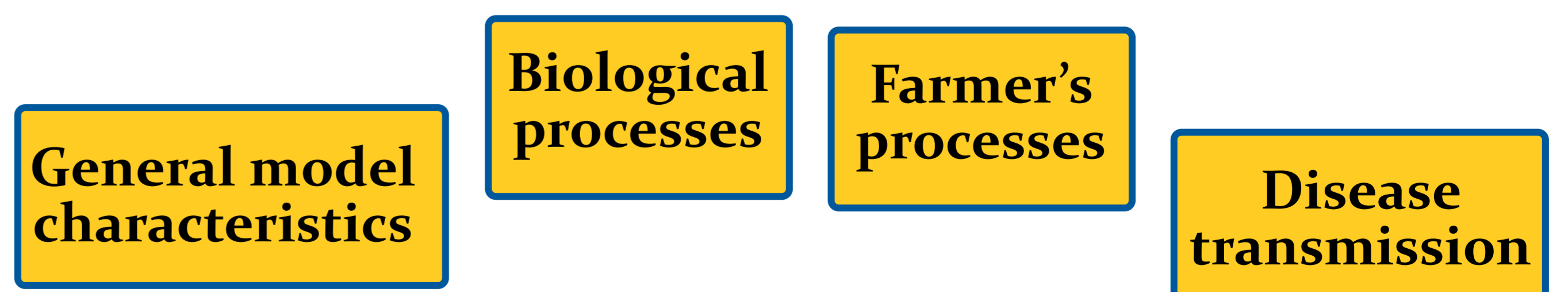
$$\Delta I \sim N_S \cdot \beta_{I(x)} \frac{N_{I(x)}}{N|1}$$

37%

N_S = Number of susceptible animals
 $\beta_{I(x)}$ = Transmission coefficient for infectious state x
 $N_{I(x)}$ = Number of infected animals in state x
 N = Number of all animals

Conclusion

We propose structuring the complexity of cattle disease models by four main features:



It would be of benefit if future model descriptions were to follow this structure.