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Coxiella burnetii within- and between-herd true seroprevalence assessment in domestic ruminants in France accounting for diagnostic uncertainty with latent class

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Context and objectives

Q fever : a worldwide zoonosis still difficult to control

- Human outbreaks regularly occur over the world
- In Europe, most human cases are related to **domestic ruminant exposure**
- **Unbiased estimation of the prevalence** is crucial to detect and assess epidemiological changes

Apparent seroprevalence levels in France and potential bias

First large epidemiological study (Gache *et al.* 2017) :

⇒ **Animal-level seroprevalence** of 22.2% in **cattle**, 41.5% in **goats** and 25.7% in **sheep**



Only apparent seroprevalence were assessed (*i.e.* Se and Sp considered = 100%)
See Poster #231 : **Se = [54% ; 75%]** and **Sp = [97% and 99%]**

⇒ **Without accounting for the diagnostic uncertainty**, results were potentially biased with an under or over estimation of true seroprevalences

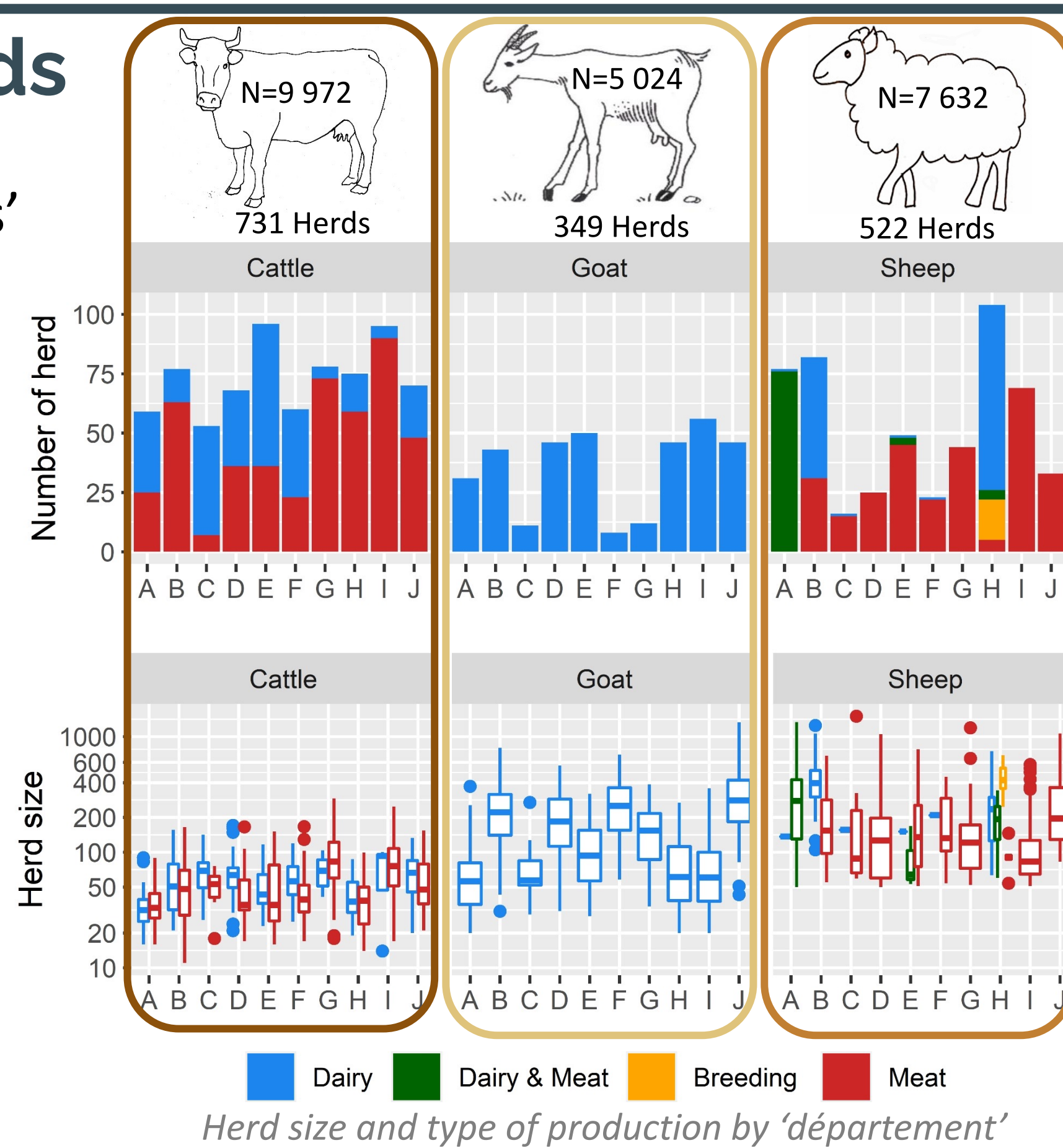
Objectives of the study

- ✓ Reassess the **between-** and **within-herd seroprevalence** in cattle, sheep and goats from the results published by Gache *et al.* 2017
- ✓ Quantify the importance of two potential **risk factors of seropositivity** at the animal and herd level (type of production and herd size)

Material and methods

Data

- Sampling in 10 French 'départements'
- **Random selection** of 19 to 106 herds by 'département' and species
- **Convenience sample** of 10 to 15 animals by herd
- **Serum analysis** in 10 veterinary laboratories with **Priocheck™ ELISA test**
- **Additional informations**
 - ⇒ Herd size = number of :
 - Females for **cattle**
 - Animals for **sheep** and **goats**
 - ⇒ Types of production
 - Dairy/meat** for **cattle** and **goats**
 - Dairy/meat/dairy & meat/breeding** for **sheep**



Hierarchical logistic model

- Two intricate **logistic regression models** for the distribution of :
 - ⇒ The **between-herd prevalence** (BHP_{ij}): the proportion of truly seropositive herds in each 'département'
 - ⇒ The **within-herd prevalence** (WHP_{ij}): the proportion of truly seropositive animals in each seropositive herd
- The **number of tested seropositive animals** in each herd was then supposed to follow a binomial distribution which depends on :
 - ⇒ the number of animals sampled
 - ⇒ the herd status (seropositive or not)
 - ⇒ the WHP_{ij}
 - ⇒ the **sensitivity and specificity of the ELISA test**

Results : Seroprevalence

Median of observed and assessed seroprevalence levels in each species and type of production

	Cattle		Goat		Sheep		
Between-Herd seroprevalences	Meat	16,3*	5,7†	63,5*	47,1†	28,0*	36,7†
		[1,5 ; 19,5]	[5,6 ; 85,5]			21,6†	[17 ; 31,7]
	Dairy	54,4*	47,9†	35,7*	63,6†	78,5*	32,7†
		[20,3 ; 79,2]	[27,2 ; 48,6]			75,0*	[2,7 ; 87]
Within-Herd seroprevalences	Meat	8,3*	39,4†	35,7*	63,6†	12,5*	22,9†
		[27,3 ; 57]	[53,4 ; 73,9]			20,0*	36,9†
	Dairy	21,4*	38,7†	35,7*	63,6†	40,0*	57,8†
		[30,5 ; 49,3]	[44,5 ; 72]			13,3*	18,7†
						52,9*	[1 ; 100]

* Observed median of the apparent seroprevalences
† Assessed median of the true seroprevalence

BHPs assessed† were lower than the observed apparent* **BHPs**

⇒ Account for the **imperfect specificity** of the ELISA test

WHPs assessed† were higher than the observed apparent* **WHPs**

⇒ Account for the **moderate sensitivity** of the ELISA test

Results : Risk factors

Cattle

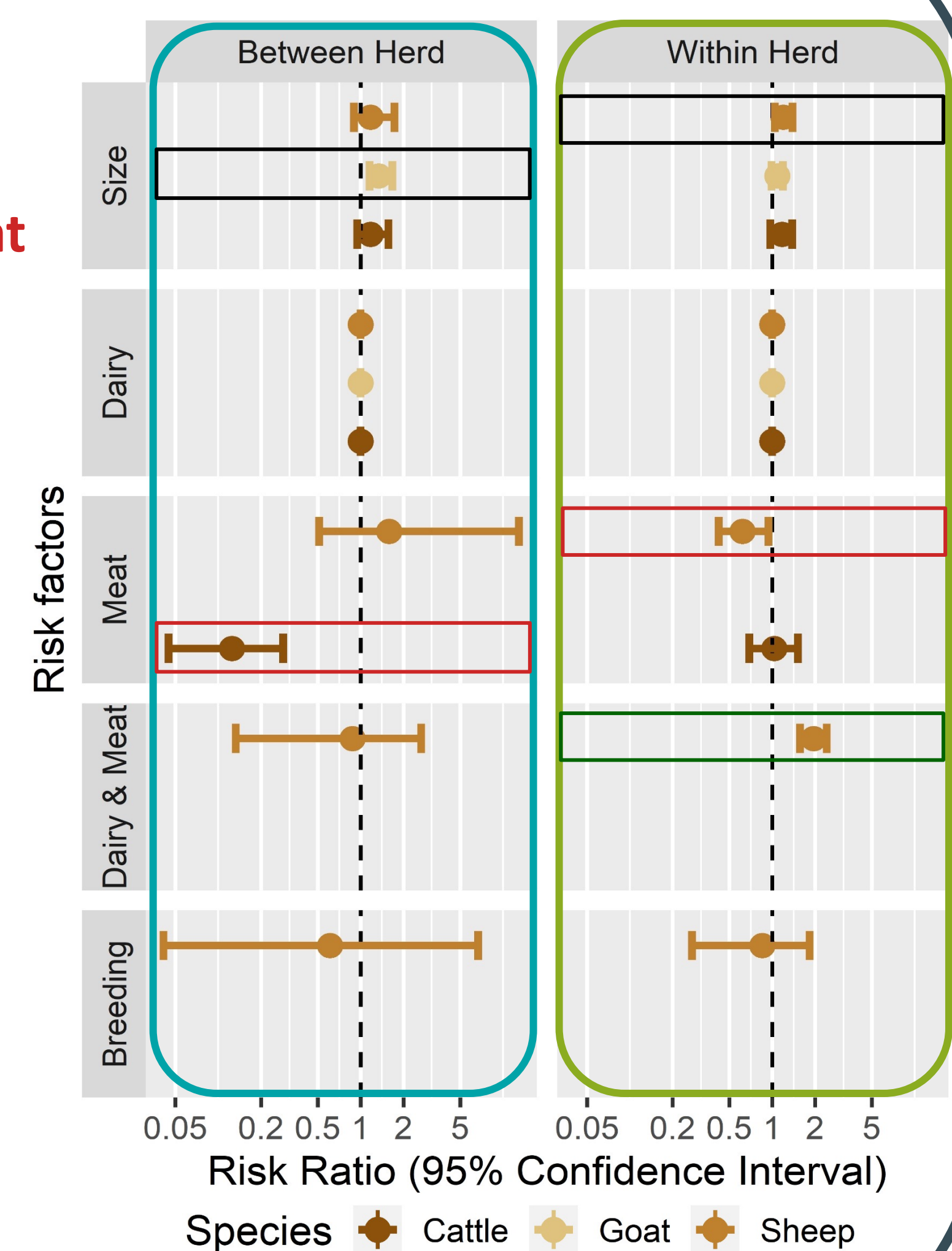
- **BHPs**
 - **7.9x higher in dairy** vs. in **meat** herds

Goats

- **BHPs**
 - **1.4x higher when herd size x2**

Sheep

- **WHPs** were :
 - **1.6x higher in dairy** vs. in **meat**
 - **2.5x higher in dairy & meat** vs. in **meat**
 - **1.3x higher when herd size x2**



Risk ratio of seropositivity of the herd (between-herd) and of the animals in seropositive herds (within-herd). The reference herd is a dairy herd of medium size in each species

Conclusion

- Better knowledge of the true **within-** and **between-herd seroprevalence levels** in France
- Identification and quantification of some relevant risk factors

⇒ **New insights related to the epidemiology of Coxiella burnetii in domestic ruminants in France**

Discussion

- Results must be carefully extrapolated
 - Potential non-representativeness of the 10 'départements' included
 - Random selection of herds without information on abortion occurrence

