



## **Erratum: Apparent prevalence of beef carcasses contaminated with mycobacterium avium subsp. paratuberculosis sampled from danish slaughter cattle (Veterinary Medicine International)**

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*Published in:*  
Veterinary Medicine International

*Link to article, DOI:*  
[10.1155/2014/437037](https://doi.org/10.1155/2014/437037)

*Publication date:*  
2014

*Document Version*  
Publisher's PDF, also known as Version of record

[Link back to DTU Orbit](#)

*Citation (APA):*  
Okura, H., Toft, N., & Pozzato, N. (2014). Erratum: Apparent prevalence of beef carcasses contaminated with mycobacterium avium subsp. paratuberculosis sampled from danish slaughter cattle (Veterinary Medicine International). *Veterinary Medicine International*, [437037]. <https://doi.org/10.1155/2014/437037>

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## *Erratum*

# **Erratum to “Apparent Prevalence of Beef Carcasses Contaminated with *Mycobacterium avium* subsp. *paratuberculosis* Sampled from Danish Slaughter Cattle”**

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Received 15 January 2014; Accepted 4 March 2014; Published 23 March 2014

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A tabulation error appeared in Table 1. The fecal contamination data for Age <2 years and >2 years were incorrectly recorded into the table. Whereas the numbers in the table were transposed incorrectly, the actual statistical analyses remain correct. The revised table is reproduced below.

TABLE 1: Number and group specific percentage (in brackets) of muscle tissue samples from slaughtered cattle from which *Mycobacterium avium* subsp. *paratuberculosis* DNA was detected by IS900 PCR at threshold cycle value of 36, 37, and 38 by age, type of herd, breed type, and fat code. Positive results were defined in relation to possible fecal contamination to the tissue samples.

Variable	Level	Including samples with fecal contamination			Excluding samples with fecal contamination				
		Total number in group	PCR threshold cycle value			Total number in group	PCR threshold cycle value		
			36	37	38		36	37	38
		501	20 (4)	28 (6)	42 (8)	385	16 (4)	22 (6)	34 (9)
Age	<2 years	197	<b>4 (1)</b>	<b>7 (2)</b>	<b>11 (4)</b>	159	3 (1)	4 (2)	8 (4)
	≥2 years	304	<b>16 (8)</b>	<b>21 (11)</b>	<b>31 (16)</b>	226	13 (8)	18 (11)	26 (16)
Herd type	Dairy	313	17 (5)	22 (7)	30 (10)	235	13 (6)	17 (7)	23 (10)
	Beef	115	2 (2)	3 (3)	7 (6)	88	2 (2)	3 (3)	7 (8)
	Veal	73	1 (0)	3 (4)	5 (7)	62	1 (2)	2 (3)	4 (6)
Breed group	Dairy	362	18 (5)	26 (7)	35 (10)	278	14 (5)	20 (7)	27 (10)
	Beef	76	1 (1)	1 (1)	4 (5)	62	1 (2)	1 (2)	4 (6)
	Cross	63	1 (2)	1 (2)	3 (5)	45	1 (2)	1 (2)	3 (7)
Fat code	1	66	1 (2)	3 (5)	8 (12)	56	1 (2)	3 (5)	7 (13)
	2	162	9 (6)	14 (9)	15 (9)	125	7 (6)	10 (8)	11 (9)
	≥3	272	10 (4)	11 (4)	19 (7)	203	8 (4)	9 (4)	16 (8)