

Journal of Veterinary Diagnostic Investigation

<http://vdi.sagepub.com/>

Interlaboratory Comparison of Real-Time Polymerase Chain Reaction Methods to Detect *Coxiella Burnetii*, The Causative Agent of Q Fever

Rebecca M. Jones, Stefan Hertwig, James Pitman, Richard Vipond, Anna Aspán, Göran Bölske, Conall McCaughey, James P. McKenna, Bart J. van Rotterdam, Arnout de Bruin, Robin Ruuls, Rob Buijs, Hendrik-Jan Roest and Jason Sawyer

J VET Diagn Invest 2011 23: 108

DOI: 10.1177/104063871102300118

The online version of this article can be found at:

<http://vdi.sagepub.com/content/23/1/108>

Published by:



<http://www.sagepublications.com>

On behalf of:



Official Publication of the American Association of Veterinary Laboratory Diagnosticians, Inc.

Additional services and information for *Journal of Veterinary Diagnostic Investigation* can be found at:

Email Alerts: <http://vdi.sagepub.com/cgi/alerts>

Subscriptions: <http://vdi.sagepub.com/subscriptions>

Reprints: <http://www.sagepub.com/journalsReprints.nav>

Permissions: <http://www.sagepub.com/journalsPermissions.nav>

>> [Version of Record](#) - Jan 1, 2011

[What is This?](#)

J Vet Diagn Invest 23:108–111 (2011)

Interlaboratory comparison of real-time polymerase chain reaction methods to detect *Coxiella burnetii*, the causative agent of Q fever

Rebecca M. Jones,¹ Stefan Hertwig, James Pitman, Richard Vipond, Anna Aspán, Göran Bölske, Conall McCaughey, James P. McKenna, Bart J. van Rotterdam, Arnout de Bruin, Robin Ruuls, Rob Buijs, Hendrik-Jan Roest, Jason Sawyer

Abstract. The bacterium *Coxiella burnetii*, which has a wide host range, causes Q fever. Infection with *C. burnetii* can cause abortions, stillbirth, and the delivery of weak offspring in ruminants. *Coxiella burnetii* infection is zoonotic, and in human beings it can cause chronic, potentially fatal disease. Real-time polymerase chain reaction (PCR) is increasingly being used to detect the organism and to aid in diagnosis both in human and animal cases. Many different real-time PCR methods, which target different genes, have been described. To assess the comparability of the *C. burnetii* real-time PCR assays in use in different European laboratories, a panel of nucleic acid extracts was dispatched to 7 separate testing centers. The testing centers included laboratories from both human and animal health agencies. Each laboratory tested the samples using their in-house real-time PCR methods. The results of this comparison show that the most common target gene for real-time PCR assays is the *IS1111* repeat element that is present in multiple copies in the *C. burnetii* genome. Many laboratories also use additional real-time PCR tests that target single-copy genes. The results of the current study demonstrate that the assays in use in the different laboratories are comparable, with general agreement of results for the panel of samples.

Key words: *Coxiella burnetii*; Q fever; real-time polymerase chain reaction; ring-trial.

From the Veterinary Laboratories Agency, Weybridge, Addlestone, Surrey, United Kingdom (Jones, Sawyer); Federal Institute for Risk Assessment, Berlin, Germany (Hertwig); Health Protection Agency, Salisbury, Wilts, United Kingdom (Pitman, Vipond); National Veterinary Institute, Uppsala, Sweden (Aspán, Bölske); Royal Hospitals, Belfast HSC Trust, Belfast, United Kingdom (McCaughey, McKenna); National Institute for Public Health and the Environment, Antonie van Leeuwenhoeklaan, Bilthoven, The Netherlands (van Rotterdam, de Bruin); and the Central Veterinary Institute of Wageningen UR, The Netherlands (Ruuls, Buijs, Roest).

¹Corresponding Author: Rebecca Jones, Veterinary Laboratories Agency, Woodham Lane, Addlestone, Surrey KT15 3NB, United Kingdom. r.jones@vla.defra.gsi.gov.uk

With the exception of Antarctica and New Zealand, Q fever is a worldwide endemic disease. The causative agent of Q fever is *Coxiella burnetii*, a small Gram-negative obligate intracellular bacterium.¹ Ruminants are the most common animal reservoir for the bacterium.³ Infection of ruminants is usually subclinical, but clinical signs including abortions, stillbirths, and the delivery of weak offspring can occur.¹

Infected animals can excrete *C. burnetii* in feces, urine, milk, and birth products. The bacterium can survive extracellularly in the environment for extended periods of time (up to 150 days) while still remaining infectious¹ and has been reported¹⁹ to have an infectious dose as low as a

single bacterium. Naive animals become infected either through the inhalation of contaminated aerosols or through the ingestion of infectious matter.³

A zoonotic disease, Q fever may develop into potentially fatal chronic disease in human beings.^{1,20} *Coxiella burnetii* infection in humans can be associated with an outbreak or can be acquired sporadically through occupational exposure to animals or animal products, thus presenting an increased risk of disease.¹³ Outbreaks of Q fever in human beings are often linked to airborne spread of the disease from infected ruminants, particularly during parturition.^{8,9,14,21,23} As a result of its high infectivity, the possibility of spread via aerosols, and the potentially fatal consequences of infection, *C. burnetii* is categorized as a biosafety level 3 agent.

Over recent years, there have been several high-profile Q fever outbreaks in Europe, including a large outbreak that is currently ongoing in The Netherlands.^{9,18,22} The increasing profile of the disease has led to renewed interest in the diagnosis of cases of Q fever both in human beings and animals, and real-time polymerase chain reaction (PCR) is rapidly being adopted throughout Europe to aid in the detection of *C. burnetii*.

A ring-trial was arranged involving 7 European testing centers, comprising both veterinary and human laboratories, to compare a number of the real-time PCR methods currently in use. Seven laboratories participated in the ring-trial, including the following: Veterinary Laboratories Agency (VLA), Weybridge, United Kingdom; Health Protection Agency, Porton Down, United Kingdom; Royal Hospitals, Belfast, United Kingdom; National Institute for Public Health and the Environment, Bilthoven, The Netherlands; Federal Institute for Risk Assessment, Berlin, Germany; National Veterinary Institute, Uppsala, Sweden; and Central Veterinary Institute of Wageningen UR, The Netherlands.

As a result of the designation of *C. burnetii* as a biosafety level 3 agent, there are concomitant difficulties with transporting positive sample material to laboratories located around Europe. Therefore, nucleic acid extracts were prepared from the sample panel at a central laboratory (VLA) and dispatched frozen to the participating testing centers. Samples were of caprine, ovine, and bovine origin and consisted of placental cotyledons or fetal fluid samples from abortion cases. Samples that were designated as “test-positive” had previously been demonstrated to contain *C. burnetii* through the use of immunohistochemistry or through microscopic evaluation of stained smears. Prior to dispatch, the suitability of the sample panel for the present study was assessed using a real-time PCR assay targeting the *IS1111* repeat element.

A small section of placental cotyledon tissue (1–10 mg) was mechanically disrupted using a micropestle and mortar. Both the tissue samples and the fetal fluid samples were subjected to an external overnight lysis. The DNA was then extracted from the digested samples using an extraction robot^a along with the dedicated buffers contained in the associated reagent kit.^a Samples were diluted in nuclease-free water if required.

Table 1. Sample panel composition.

Sample	Species of origin	Sample type	Other information
A	Caprine	Fetal fluid	<i>Coxiella burnetii</i> positive
B	Caprine	Fetal fluid	1 in 10 dilution of sample A
C	Caprine	Fetal fluid	1 in 100 dilution of sample A
D	Caprine	Fetal fluid	<i>C. burnetii</i> positive
E	Caprine	Placental cotyledon	<i>C. burnetii</i> positive
F	Ovine	Placental cotyledon	Negative for <i>C. burnetii</i>
G	Ovine	Fetal fluid	Negative for <i>C. burnetii</i>
H	Bovine	Placental cotyledon	<i>C. burnetii</i> positive
I	Caprine	Placental cotyledon	<i>C. burnetii</i> positive
J	Caprine	Placental cotyledon	<i>C. burnetii</i> positive

Each testing center received a panel of 10 nucleic acid samples, which included extracts from 2 samples that had previously tested negative for *C. burnetii* and 6 samples that had previously tested positive for *C. burnetii*. To assist in distinguishing between the efficacies of different real-time PCR methods, 1 sample (sample A) was subjected to a 10-fold serial dilution to produce samples B and C. The composition of the sample panel is shown in Table 1.

Each participating laboratory tested the panel of nucleic acid extracts using their in-house assay(s) (Table 2). Testing centers were blinded to the *C. burnetii* status of the samples being tested. In some laboratories, the samples were tested using more than one real-time PCR, and at some testing centers, the samples were tested by multiple operators. Each real-time PCR assay was used to test the nucleic acid samples using between 2 and 6 replicate reactions for each template.

The results of the ring-trial (Table 2) demonstrated that samples A, D, E, H, I, and J were positive for *C. burnetii* with all real-time PCR methods used. Samples F and G had been included as samples, which had previously tested negative for *C. burnetii*. Sample G tested negative with all of the real-time PCR assays used, while sample F was negative in the majority of assays, with 1 assay showing amplification in a single reaction.

The inclusion of samples B and C allowed for discrimination between the differing sensitivities of the assays (Table 2). Sample B was positive in all of the assays targeting the *IS1111* repeat element, while only 1 single-copy gene assay was able to detect the *C. burnetii* DNA present in this sample. *Coxiella burnetii* DNA was not detected in sample C by any of the PCR assays targeting single-copy genes, and *C. burnetii* DNA could only be detected in sample C using 2 of the 7 *IS1111* real-time PCR assays.

Strikingly, when the data for each positive sample are examined at the level of quantification cycle, there is close agreement between the results obtained with the real-time PCR methods that target the same region of the *C. burnetii* genome. This is especially surprising given that different PCR

Table 2. Comparison of real-time polymerase chain reaction results generated from 7 participating laboratories.*

Target	Source of assay	A	B	C	D	E	F	G	H	I	J
<i>IS1111</i>	Modified from Marmion et al., 2005 ¹¹	31.3	34.1	36.9	28.6	20.6	N	N	22.7	14.3	14.4
	Marmion et al., 2005 ¹¹	38.3	40.5	IN	35.4	26.0	N	N	28.2	18.6	18.3
	Harris et al., 2000 ⁷	35.4	38.2	N	32.1	24.1	N	N	25.5	17.4	17.3
	Unpublished assay	31.4	33.4	N	29.9	22.8	IN	N	23.7	15.3	15.3
	Klee et al., 2006 ¹⁰	34.5	37.2	N	32.8	24.5	N	N	25.2	15.4	15.5
	Commercial kit ^b	31.1	35.3	N	29.4	21.8	N	N	23.0	14.2	14.3
	Unpublished assay	32.6	36.6	38.3	30.5	22.5	N	N	23.4	14.5	14.3
<i>COM1</i>	Modified from Marmion et al., 2005 ¹¹	37.2	38.9	N	35.0	28.0	N	N	28.8	22.4	21.5
	Marmion et al., 2005 ¹¹	IN	N	N	53.3	43.6	N	N	46.4	36.8	38.6
	Unpublished assay	38.9	IN	N	35.0	27.9	N	N	31.7	21.8	21.7
<i>ICD</i>	Unpublished assay	32.6	N	N	30.5	23.0	N	N	23.8	17.4	17.3
	Unpublished assay	36.9	N	N	33.5	25.8	N	N	26.7	20.2	20.1
<i>16S</i>	Marmion et al., 2005 ¹¹	40.0	N	N	37.5	28.9	N	N	30.2	23.4	24.1
<i>HYPO</i>	Unpublished assay	36.0	IN	N	33.4	25.8	N	N	27.1	19.6	19.5

* N = negative; IN = inconclusive (samples were considered to be inconclusive when amplification was not present in the majority of replicates). For positive samples, the mean quantification cycle value is presented.

reagents and real-time PCR platforms were used in the different testing centers. Additionally, when each assay is used to rank the samples in order of increasing bacterial load, there is again close agreement among the assays.

The present survey of the real-time PCR assays used to detect *C. burnetii* in different European laboratories showed that all of the participating laboratories were using assays targeting the multicopy *IS1111* repeat element. Many laboratories used multiple real-time PCR assays, combining an *IS1111* assay with assays targeting single-copy genes such as *COM1* or *ICD*.

Disease outbreak investigations may cross borders, involving agencies in different countries, and may involve both human and animal health agencies within a country. Given this, it is important that comparability is achieved for *C. burnetii* detection among these different agencies. The results of the current study indicate that the real-time PCR methods used to detect *C. burnetii* are comparable in different testing centers across the European Union and that laboratory test result agreement exists between animal and human agencies within the same country.

The results of the current study indicate that real-time PCR assays that target multiple copy genes, such as the *IS1111* gene, are more sensitive than are those targeting single-copy genes, such as the *COM1* or *ICD* genes. However, many laboratories use additional assays, such as those targeting single-copy genes. This provides an additional level of certainty for any positive result, as it demonstrates that multiple independent sections of the *C. burnetii* genome can be detected in a sample. Additionally, assays targeting the *IS1111* repeat element cannot be used for accurate quantification of the load of *C. burnetii* present within a sample, as this element is present in differing numbers (7–110)¹⁰ depending on the strain of the bacterium present in the sample. The additional use of a PCR targeting a single-copy gene might allow accurate quantification of the bacterial load present within a sample. It is also worth noting that the single-copy methods were able to accurately show the presence of *C. burnetii* DNA in typical

placental cotyledon and fetal fluid samples, such as those used in the current exercise.

Use of real-time PCR provides a sensitive method of detecting *C. burnetii* in animal and human samples. Unlike serological assays, PCR can be used to demonstrate the presence of the organism within clinical material,^{2,4-6,12,15,16} and, by doing so, it provides important information to the veterinarian, which will help to inform the final diagnosis. This may be especially relevant in cases involving ruminant abortions, in which demonstrating the presence of *C. burnetii* within the abortion material may implicate the organism as the cause of the abortion. Critically, subject to a suitable extraction method, real-time PCR might be used to detect the presence of *C. burnetii* within a multitude of different sample types, including bedding material, straw, and birth products (including placental and fetal material), that could be associated with the production of contaminated aerosols, which may be potentially involved in a disease outbreak.^{9,17,23} Polymerase chain reaction assays may also be used to detect *C. burnetii* in milk.¹⁵ The current study has shown that suitable detection methods are available in many laboratories to detect *C. burnetii* in a disease outbreak.

Acknowledgements. The authors would like to thank Jane Errington and Sue Hannon for performing the nucleic acid extractions and the staff from the Veterinary Laboratories Agency Regional Laboratories for providing samples for use in the study. This work was supported by the FZ2100 non-statutory zoonoses surveillance budget, the VLA Test Development Programme, and by the MedVetNet Workpackage 25. British Crown copyright and Crown user rights are reserved.

Sources and manufacturers

- MagNA Pure LC, Roche Diagnostics Ltd., Burgess Hill, United Kingdom.
- Adiavet Cox, Adiagene, Saint Brieuc, France.

References

1. Arricau-Bouvery N, Rodolakis A: 2005, Is Q fever an emerging or re-emerging zoonosis? *Vet Res* 36:327–349.
2. Berri M, Souriau A, Crosby M, Rodolakis A: 2002, Shedding of *Coxiella burnetii* in ewes in two pregnancies following an episode of *Coxiella* abortion in a sheep flock. *Vet Microbiol* 85:55–60.
3. Fournier PE, Marrie TJ, Raoult D: 1998, Diagnosis of Q fever. *J Clin Microbiol* 36:1823–1834.
4. Fournier PE, Raoult D: 2003, Comparison of PCR and serology assays for early diagnosis of acute Q fever. *J Clin Microbiol* 41:5094–5098.
5. Guatteo R, Beaudeau F, Berri M, et al.: 2006, Shedding routes of *Coxiella burnetii* in dairy cows: implications for detection and control. *Vet Res* 37:827–833.
6. Guatteo R, Beaudeau F, Joly A, Seegers H: 2007, *Coxiella burnetii* shedding by dairy cows. *Vet Res* 38:849–860.
7. Harris RJ, Storm PA, Lloyd A, et al.: 2000, Long-term persistence of *Coxiella burnetii* in the host after primary Q fever. *Epidemiol Infect* 124:543–549.
8. Hawker JI, Ayres JG, Blair I, et al.: 1998, A large outbreak of Q fever in the West Midlands: windborne spread into a metropolitan area? *Commun Dis Public Health* 1:180–187.
9. Karagiannis I, Schimmer B, van Lier A, et al.: 2009, Investigation of a Q fever outbreak in a rural area of The Netherlands. *Epidemiol Infect* 137:1283–1294.
10. Klee SR, Tyczka J, Ellerbrok H, et al.: 2006, Highly sensitive real-time PCR for specific detection and quantification of *Coxiella burnetii*. *BMC Microbiol* [serial online] 6:2.
11. Marmion BP, Storm PA, Ayres JG, et al.: 2005, Long-term persistence of *Coxiella burnetii* after acute primary Q fever. *QJM* 98:7–20.
12. Masala G, Porcu R, Daga C, et al.: 2007, Detection of pathogens in ovine and caprine abortion samples from Sardinia, Italy, by PCR. *J Vet Diagn Invest* 19:96–98.
13. Orr HJ, Christensen H, Smyth B, et al.: 2006, Case-control study for risk factors for Q fever in southwest England and Northern Ireland. *Euro Surveill* 11:655.
14. Porten K, Rissland J, Tigges A, et al.: 2006, A super-spreading ewe infects hundreds with Q fever at a farmers' market in Germany. *BMC Infect Dis* [serial online] 6:147.
15. Rodolakis A, Berri M, Hechard C, et al.: 2007, Comparison of *Coxiella burnetii* shedding in milk of dairy bovine, caprine, and ovine herds. *J Dairy Sci* 9:5352–5360.
16. Rousset E, Berri M, Durand B, et al.: 2009, *Coxiella burnetii* shedding routes and antibody response after outbreaks of Q fever-induced abortion in dairy goat herds. *Appl Environ Microbiol* 75:428–433.
17. Rustcheff S, Norlander L, Macellaro A, et al.: 2000, A case of Q fever acquired in Sweden and isolation of the probable ethiological agent, *Coxiella burnetii* from an indigenous source. *Scand J Infect Dis* 32:605–607.
18. Schimmer B, Dijkstra F, Vellema P, et al.: 2009, Sustained intensive transmission of Q fever in the south of The Netherlands, 2009. *Euro Surveill* 14:19210.
19. Tigertt WD, Benenson AS, Gochenour WS: 1961, Airborne Q fever. *Bacteriol Rev* 25:285–293.
20. Tissot-Dupont H, Raoult D: 2008, Q fever. *Infect Dis Clin North Am* 23:505–514.
21. Tissot-Dupont H, Torres S, Nezri M, Raoult D: 1999, Hyperendemic focus of Q fever related to sheep and wind. *Am J Epidemiol* 150:67–74.
22. van der Hoek W, Dijkstra F, Schimmer B, et al.: 2010, Q fever in The Netherlands: an update on the epidemiology and control measures. *Euro Surveill* 15:19520.
23. Welsh HH, Lennette EH, Abinanti FR, Winn JF: 1958, Airborne transmission of Q fever: the role of parturition in the generation of infective aerosols. *Ann N Y Acad Sci* 70:528–540.