



COXIELLA REAL TIME

Reference: ADI143-100

Test for the detection and quantification of *Coxiella burnetii* by real time enzymatic amplification
PCR Test – 100 reactions

For veterinary *in vitro* use only



Sample	Individual analysis	Pool of sample possible*, up to:
Swab (placental, vaginal...)	✓	3
Tissue (placenta, foetal tissues...)	✓	✗
Faeces	✓	✗
Amniotic fluid	✓	✗
Milk	✓	✗

* Depending on the epidemiological case and on the quality of samples.

Kit composition

Content		ADI143-100 Kit
		100 reactions
A5	Amplification solution	2 x 1000 µL tubes with green cap (Ready to use)
COX CTL+	<i>Coxiella burnetii</i> positive control	1 tube with purple cap (To reconstitute)
EPC-Ext	Exogenous extraction control for faeces or acellular samples	2 x 300 µL tubes with yellow cap (Ready to use)
NF-Water	Nuclease-Free Water	1 x 1000 µL tube with white cap (Ready to use)

Revision history

Date	Version	Modifications
01/2020	NE143-011	Last version
01/2023	V01	Change to simplified format

Note: minor typographical, grammar and formatting changes are not included in the revision history.

A. Introduction

Coxiella burnetii is a strictly intracellular Gram-negative bacterium, agent of the Q fever (Burnet and Freeman, 1937; Cox, 1938). The disease is present worldwide. *C. burnetii* is able to infect a large range of hosts including humans. Q fever is a **zoonose** (pathology which can be transmitted to human) mainly transmitted by ovine, bovine and caprine. Q fever usually results from inhalation of contaminated aerosols originating mostly from dropping and body fluids of infected animals.

C. burnetii is present in reproductive apparatus, placenta and fluids produced during parturition or abortion. The bacterium is excreted in milk and faeces of infected animals presenting no clinical signs. Mostly infections are asymptomatic or subclinical by Human, but can be manifested as a flu-like disease, or as hepatitis. A neurological involvement is also possible. Pericarditis and myocarditis are rare. *C. burnetii* can induce abortions, causing economic losses in cattle. Prevention consists mostly in precautions with pregnant females and with fluids from parturition.

B. Test principle

ADIAVET™ COXIELLA REAL TIME test is based on the amplification of specific *Coxiella burnetii* DNA (IS1111 gene). This test is intended to detect simultaneously, in one well:

- *Coxiella burnetii* (FAM labelled probe)
- GAPDH internal control of extraction and amplification specific from an endogenous nucleic acid (HEX labelled probe or its equivalent).
- or
- An exogen control EPC-Ext added during the extraction of acellular samples allows validating extraction and amplification steps (probe labelled with a fluorochrome read in the same spectra as VIC and HEX).

The positive control COX CTL+, included in the kit, enables the quantification of positives samples. The measure unit is the equivalent genome (EG) number (or bacteria) per mL.

C. Storage conditions

On receipt, the kit should be stored at <-15 °C until the expiration date.

It is recommended to make fractions of A5 solution if it should be defrosted more than 3 times.

Do not thaw more than 3 times.

Store away from sunlight.

Do not mix reagents of two different batches.

D. Material required but not provided

- Real-time Thermal cycler and device.
- Instrument for homogenous mixing of tubes.
- Pipettes of 1 - 10 µL, 20 - 200 µL and 200 - 1000 µL.
- Nuclease-free filtered pipette tips.
- Nuclease-free microtubes of 1,5 mL and 2 mL.
- Powdered-free latex or nitrile gloves.
- Nuclease-free water.
- Kit for nucleic acids extraction.

Additional kits for method adoption and PCR

- **LD_{PCR} Positive Control – COX (Ref.: ADC14LD)**
Confirmation of performances – LOD_{PCR} of kit.

E. Warnings and precautions

- For veterinary *in vitro* use only.
- For animal use only.
- For professional use only.

- All instructions must be read before performing the test and strictly respected.
- Do not use reagents after the expiration date.
- Do not use reagents if the packaging is damaged.
- Do not open PCR wells or tubes after amplification.
- Do not mix reagents from different batches.
- Used material must be disposed of in compliance with the legislation in force regarding environmental protection and biological waste management.
- This kit contains products of animal origin. Certified knowledge of the origin and/or sanitary state of the animals does not totally guarantee the absence of transmissible pathogenic agents. It is therefore recommended that these products be treated as potentially infectious and handled observing the usual safety precautions (do not ingest or inhale).

F. Nucleic acids extraction

1. Extraction kits

Nucleic acids must be extracted from the samples before using the kit. The RNA/DNA purification kits listed below are recommended by Bio-X Diagnostics:

Product name	Extraction system	Number of tests and reference
ADIAMAG	Magnetic beads	200 tests: ref. NADI003 800 tests : ref. NADI003-XL

For the extraction, consult the user manual version, available on the website, indicated on the certificate of analysis included in the PCR kit of interest.

The validated extraction protocols are described in the kit validation file. Other purification kits can be used if they have been validated by the user.

After extraction, nucleic acid extracts can be kept on ice or at +2/8 °C for a few hours or until use. For long term storage, they must be kept at a temperature below -15 °C or -65 °C.

2. Controls

Using controls allow to verify the reliability of the results. Controls can be included by series of analysis according to the recommendations defined by the standards in force (Cf. AFNOR U47-600...).

Controls	Validation of	How to proceed
No Template Control (NTC)	Absence of amplification contamination	5 µL NF-Water in a well per run
COX CTL+ (Dilution pure to 1/10 000)	<i>C. burnetii</i> target amplification and standard range	5 µL CTL+ for each range point in a well per run
Negative extraction control	Absence of contamination for the extraction and amplification	1 extraction (water or lysis buffer) per run
Positive extraction control	Extraction and amplification	1 extraction (Positive sample between 1 et 100X LOD _{Method}) per run

G. Procedure

1. Use of EPC-Ext

EPC-Ext must be added to each faeces or acellular sample.

Aliquot and store the solution at a temperature below -15 °C according to the size of extraction series. Do not thaw more than 3 times.

Add **5 µL** of EPC-Ext in the first nucleic acids extraction lysis buffer.

2. Use of COX CTL+

Carefully use this CTL+. It contains a large amount of DNA that could be a source of contamination.

The tube COX CTL+ contains a titrated solution of *C. burnetii* DNA as EG/mL or as *C. burnetii*/mL.

Add **200 µL** of « **NF-Water** » per tube.

Homogenize the tubes using a mixer, such as vortex, > 20 seconds and until dissolution of the blue pellet.

After reconstitution, aliquot and store the solution at a temperature below -15 °C until the kit expiration date. Do not thaw more than 3 times.

If quantification is needed, prepare the following ranges in Nuclease-free water:

Dilution	COX CTL+ concentration (<i>C. burnetii</i> /mL)
Pure	4x10 ⁶
1/10	4x10 ⁵
1/100	4x10 ⁴
1/1000	4x10 ³
1/10000	4x10 ²

Use **5 µL** of each dilution in the dedicated wells (see § « Amplification », Step 2).

In the case of a qualitative test, use **5 µL** of the 1/1000 dilution in one of the dedicated wells (see § « Amplification », Step 2).

3. Amplification

Warning:

- Before starting, rehydrate or thaw reagents at room temperature in the dark.
- Homogenize all reagents and samples before use.
- Store reagents at a temperature below -15 °C after use.

Step 1: Dispense **20 µL** of amplification solution (A5) per well.

Step 2: Dispense **5 µL** of nucleic acids extracts and **5 µL** of controls in each dedicated well.

Use NF-Water for the No Template Control (NTC).

Step 3: Cover the wells with an appropriate optical film or caps.

Step 4: Start the PCR analysis.

The following program is defined for ABI Prism thermocyclers (like 7500, QuantStudio5, Step-one...) from Applied Biosystems, for Mx3000, Mx3005P and AriaMx from Agilent, for LightCycler from Roche Diagnostics, for Rotor-Gene Q from Qiagen, for CFX96 and Chromo 4 from Biorad and for MIC from BioMolecular System.

DNA standard program	
2 min. 50 °C	
10 min. 95 °C	
15 sec. 95 °C*	45 cycles
60 sec. 60 °C**	

*30 sec. 95 °C for MX3000 and MX3005P

** Reading and parameters for fluorescence acquisition:

Fluorochrome	Absorbance (nm)	Emission (nm)
FAM	494	520
HEX or equivalent	530	549
ROX	575	602

Note: The Quencher is non-fluorescent. The A5 solution contains a passive reference read in the same spectra as ROX for ABI machines.

For other thermal cycler instruments, please contact your sales representative or the customer relations department.

H. Interpretation of results

1. Validation and interpretation of qualitative results

Display all curves and position the threshold line for each fluorochrome.

a. Test validation

Amplification is valid if the following results are obtained.

Expected Ct (Threshold Cycle) values for the CTL+ are indicated on the certificate of analysis of the kit.

Controls	Amplification		Validation of
	FAM	HEX or equivalent	
No Template Control (NTC)	No	No	Absence of amplification contamination
COX CTL+ (1/1000)	Yes	No	Target amplification
Extraction negative control	No	No	Absence of extraction contamination
Extraction positive control	Yes	Yes/No	Extraction and amplification steps

b. Results interpretation

Nucleic acids extraction and amplification are **valid** for each sample if at least one typical amplification curve is observed in FAM and/or HEX or equivalent.

Amplification		Interpretation
FAM	HEX or equivalent	<i>C. burnetii</i>
No	Yes	Undetected
Yes	Yes	Detected
Yes	No	Detected
No	No	Undetermined

« **Undetermined** »: no characteristic amplification curve.

Possible causes:

Defective PCR due to inhibitors, set up error, absence of samples, degraded samples and/or issue with nucleic acids extraction (loss or destruction of nucleic acids).

Recommendations:

Set up a new PCR assay using pure nucleic acids extracts and 10x dilutions in Nuclease-free water.

If the assay is inconclusive, perform a new nucleic acids extraction.

2. Validation and interpretation of quantitative results

a. CTL+ range

COX CTL+ dilution	Concentration (<i>C. burnetii</i> /mL)	FAM amplification	Validation of
Pure	4x10 ⁶	Yes	<i>C. burnetii</i> target amplification and calibration curve set up
1/10	4x10 ⁵	Yes	
1/100	4x10 ⁴	Yes	
1/1000	4x10 ³	Yes	
1/10000	4x10 ²	Yes	

To interpret quantitative results, set up a calibration curve (number of cycles = f (Log concentration)), determine the curve equation ($y = ax + b$) and check PCR efficiency ($Eff\% = \left(10^{\frac{-1}{a}} - 1\right) \times 100$).

The calibration curve is valid if:

- The 5 points of the range are amplified. However, one point of the range can be omitted if that point is not one of the extreme points.
- The coefficient of correlation R^2 is higher than 0,9.
- Efficiency between 75 et 125 %.
- Points of the range are spread homogenously.

b. Quantification interpretation

Quantification of a positive sample is only possible in the quantification domain of the method use (see validation data).

FAM amplification	Sample status for <i>C. burnetii</i>
No signal	Undetected Nucleic acid undetected
Signal < LQ _{METHOD}	Detected Nucleic acid detected with a quantity under the LQ _{METHOD}
LQ _{METHOD} < signal < LQ _{max}	Detected Quantifiable nucleic acid
Signal > LQ _{max}	Detected Nucleic acid detected with a quantity over the LQ _{MAX}

In the case of a "quantifiable" sample, *C. burnetii* concentration is determined using the calibration curve equation:

$$x = 10^{\left(\frac{y-b}{a}\right)} \times F$$

Where x: concentration (in *C. burnetii*/mL if F is omitted).
y: Ct value in FAM for the positive sample to quantify.
b: intercept.
a: slope.
F: multiplying coefficient (optional).

The multiplying coefficient is determined according to the sample matrix and extraction method.

Multiplying coefficient examples with ADIAMAG extraction kit according to the NEKF user manual:

Matrix	Multiplying coefficient (F)	Unit
Vaginal swab, placenta swab, milk, foetal liquide	0.6	<i>C. burnetii</i> / mL
Faeces	3	<i>C. burnetii</i> / g
Tissue	3	<i>C. burnetii</i> / g

Symbols

Symbols	Signification
	Catalog number
	Manufacturer
	Temperature limitation
	Use by
	Batch code
	Consult Instructions for Use
	Contain sufficient for "n" tests
	For veterinary <i>in vitro</i> use only – For animal use only
	Keep away from sunlight

Bibliography

- Burnet, F. M., and M. Freeman (1937). Experimental studies on the virus of Q fever. Med. J. Aust. 2:299-302
- Cox, H. R. (1938). A filter-passing infectious agent isolated from ticks. III. Description of organism and cultivation experiments. Public Health Rep. 53:2270-2276
- Norme AFNOR NF U47-600-1 (Février 2015), Méthodes d'analyse en santé animale – PCR (réaction de polymérisation en chaîne - Partie 1: exigences et recommandations pour la mise en œuvre de la PCR en santé animale.
- Norme AFNOR NF U47-600-2 (Février 2015), Méthodes d'analyse en santé animale – PCR (réaction de polymérisation en chaîne - Partie 2: exigences et recommandations pour le développement et la validation de la PCR en santé animale

1 | Extract nucleic acids with

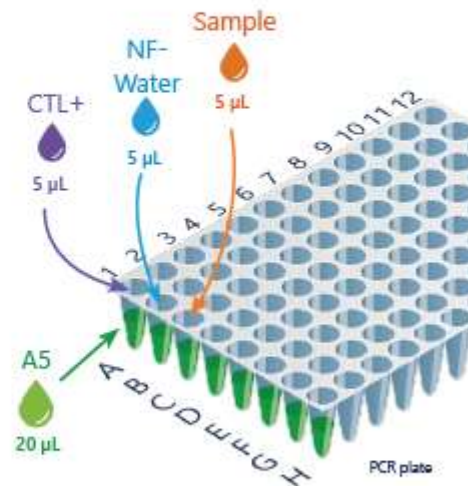
**Adia^X
Mag**



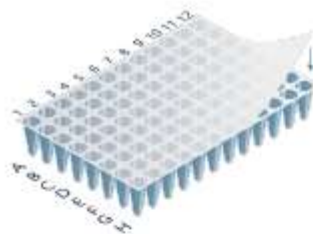
Scan me to
discover Adiamag™

2 | Distribute 20 μ L of A5 amplification
solution

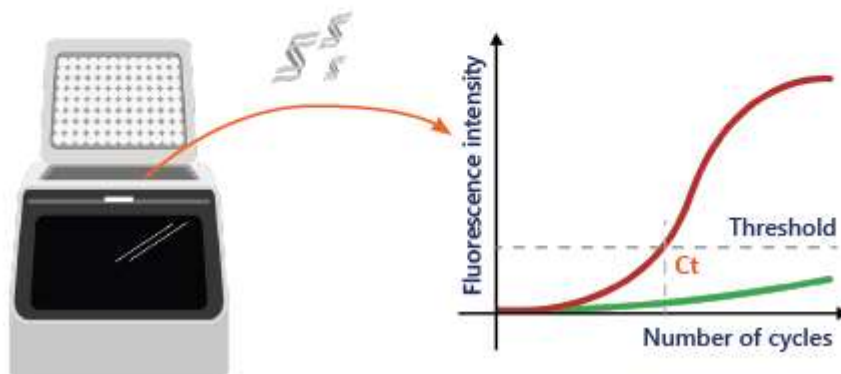
3 | Distribute 5 μ L of nucleic acids, CTL+
and NF-Water



4 | Seal the wells



5 | Start PCR analysis



*The notes do not replace the instructions for use of
which they are a summary.