

## Xpert qDetect *Campylobacter jejuni*

#GDK05.0100 (100 rxns)  
 (FOR RESEARCH ONLY)



**Product:** Although rarely fatal to humans, *Campylobacter*-associated food poisoning is nowadays a leading cause of bacterial gastroenteritis worldwide. Moreover, infections can also cause extraintestinal diseases and severe long-term complications. *Campylobacter jejuni* accounts for approximately 80% of all *Campylobacter* infections in humans.

These Gram-negative bacteria are associated with poultry and animal feces and infections are linked to the consumption of contaminated water and a wide variety of food products of animal origin, including raw milk, poultry, and undercooked meat.

Traditional microbiological detection and confirmation methods are long established and well developed but typically require 4-5 days in order to obtain results. Moreover, *Campylobacter* spp. require special culture conditions, including a microaerobic environment and long exposure to normal atmosphere during sample processing may inhibit growth and subsequent detection. This qPCR Detection Kit provides a fast, sensitive and reliable method for the detection of *Campylobacter jejuni* based on qPCR using specific FAM-labeled probes, requiring less than 2 days. This immense time reduction allows taking appropriate action, if needed, much sooner.

This kit is compatible with instruments equipped with FAM and ROX channels. The detection limit is approximately 500fg of *Campylobacter jejuni* DNA allowing the detection of, after enrichment, as little as 1-10 cells per 25g of food sample, with a specificity of 100%.

**Applications:** Detection of *Campylobacter jejuni* in food and animal feeding stuffs (following bacterial enrichment and subsequent DNA extraction (see: "prior to use")).

**Contents:** The qPCR Detection Kit (#GDK05.0100) *Campylobacter jejuni* contains sufficient reagents for 100 qPCR reactions.

Component	GDK05.0100
Cj Mix A	2x 840 µl
Cj Mix B	210 µl
Positive Control (Cj+)	70 µl
Negative Control (Cj-)	70 µl

**Note:** This product does not include reagents and other materials required for DNA extraction.

**Properties:** Fast, Easy and Reliable  
 Low limit of Detection  
 Validated according to ISO 22118:2011  
 100% Specificity  
 Compatible with instruments equipped with **FAM** and **ROX** channels

**Storage:** -20°C and protected from light for at least 1 year. Minimize repeated freeze/thawing, consider preparation of aliquots.

## Prior to use:

### Safety Considerations

*Campylobacter jejuni* is pathogenic to human, hence isolation and identification must be carried out by trained laboratory personnel, in a properly equipped Containment Level 2 laboratory. Care must be taken in the sterilization and disposal of all test materials. All procedures must be performed in the designated area of the laboratory. Disposable gloves must be worn during all procedures. All relevant National and Local Regulations must be met.

### Laboratory Set-Up

In order to prevent erroneous results due to contamination and/or degradation of DNA, it is strongly recommended to have dedicated materials, equipment and area for the DNA extraction, separated from the preparation of PCR procedures. Workflow in the laboratory should be unidirectional, starting with DNA isolation and moving to preparation of PCR amplification and finally to (q)PCR and Detection. Moreover, all molecular biology procedures, including DNA extraction, require qualified personnel to prevent risk of erroneous results and/or inadequate interpretation of results.

### Bacterial Enrichment

This kit is for the detection of DNA from *Campylobacter jejuni* present in total DNA previously purified from a bacterial pre-enrichment culture of a defined sample of food or animal feed. Representative food samples should be prepared conform standard procedures and under GLP conditions<sup>[1]</sup>. It is highly recommended to carry out the pre-enrichment according to ISO 10272-1:2017<sup>[2]</sup> or BAM (Chapter 9).<sup>[3]</sup>

### DNA extraction

Following incubation, cells are harvested from 1ml of the enriched culture by centrifugation at 10,000g-12,000g for 5 minutes. After washing the pellet with 0.9% Sodium Chloride (NaCl), add 200µl of Lysis Buffer (e.g. 10mM Tris-HCl pH 8.0 containing 1mM EDTA and 0.1% Tween<sup>®</sup>20) to the pellet. Resuspend by vortexing or pipetting up and down in the tube. Incubate at 95°C-100°C for 10-15 minutes using a water bath or heating block. Centrifuge 10,000g-12,000g for 2 minutes and transfer the cleared supernatant to a new microtube. Use 2µl of the supernatant as template DNA sample in the qDetect protocol.

Alternatively, DNA is extracted from the sodium chloride-washed pellet using an appropriate kit for the extraction of DNA from Gram-negative bacteria (not included) using the manufacturer's instructions. The obtained highly-purified DNA, free of possible PCR inhibitors, can be used immediately or stored at -20°C for up to several years for future analysis. Use 2µl of the eluate as template DNA sample in the qDetect protocol.

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## References

1. Public Health England (2016) Preparation of samples and dilutions, plating and sub-culture. Microbiology Services. Food, Water & Environmental Microbiology Standard Method FNES26 (F2); Version 2.
2. Microbiology of the food chain - Horizontal method for the detection and enumeration of *Champylobacter* spp. - Part 1: Detection method (ISO 10272-1:2017)
3. FDA's Bacteriological Analytical Manual, Chapter 7 (revised Dec 2000). An online edition of BAM Chapter 7 is available at <https://www.fda.gov/food/laboratory-methods-food/bam.chapter-7-champylobacter>

## qDetect - Basic Protocol

Unambiguous detection of *Campylobacter jejuni* (and of the internal control) requires a suitable calibration of both FAM and ROX channel. Please refer to the manufacturer's instruction of the real time PCR cyclers.

### 1. Mix for each qPCR reaction:

Component	Volume
Cj Mix A	16 µl
Cj Mix B	2 µl

In order to minimize risk of contamination, reagent loss and improve pipetting accuracy, we recommend to prepare a mastermix for multiple samples (N), always including a negative control, and a positive control, by mixing all components (N+2), except template DNA (nor control DNA), dividing the mixture equally into each PCR tube (18 µl each), briefly spin tubes (or tap down) and then add 2 µl template DNA or control DNA directly in the mixture.

### 2. Set-up qPCR cycling:

N° cycles	Temp	Time	Acquisition
1x	50°C	2 min	No
1x	95°C	5 min	No
40x	95°C	30 sec	No
	60°C	30 sec	Yes
	72°C	30 sec	No

After an initial cycle of 2 min at 50°C and 5 min at 95°C (Enzyme activation and denaturation of template), cycle 40 times for 30 seconds at 95°C, 30 seconds at 60°C and 30 seconds at 72°C. Acquire data for the detection of *Campylobacter jejuni* on the **FAM** channel. Probe to detect specific amplification of the internal control, which is included in Cj Mix A and which is amplified simultaneously with the target DNA, should be detected in the **ROX** channel.

## Results

### Controls

In order to validate the assay, controls must have the following results. If the signal of one of the controls does not match, the whole experiment, including all samples, must be repeated.

Control	FAM channel	ROX channel
Negative Control	Ct=N/A	positive
Positive Control	positive	unimportant*

N/A = Not applicable (signal below threshold).

\*unimportant: is expected to be positive, however, if negative but other controls match expected results, this makes no difference

### Samples

For each sample, there are 4 possible outcomes, as summarized in the table below.

FAM channel	ROX channel	Result
positive	positive	positive
positive	Ct=N/A	positive
Ct=N/A	positive	negative
Ct=N/A	Ct=N/A	inhibition*

\*) in case both *Campylobacter jejuni* (FAM channel) and Internal Control (ROX channel) have signals below threshold, but all the controls resulted in signals as expected, the sample must be retested, as the qPCR reaction was inhibited. Inhibition often is the result of a too high DNA concentration and therefore it is recommended that retesting should be carried out with a 10-fold dilution of the original DNA sample.