



Instituto de Medicina Genómica SL

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All products sold by the imegen are subjected to rigorous quality control. The **imegenTM Pork ID kit** has passed all internal validation tests, ensuring the reliability and reproducibility of each assay.

For any questions about the applications of this product or its protocols, please contact our Technical Department:

Phone number: 963 212 340 **e-Mail**: info@imegen.es

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1. Product information

Kit description

Identification of meat species presence in food samples is an essential step in order to verify the origin and traceability of the used raw materials, as well as a necessary quality control for handling and cleaning processes of production lines.

imegen-Pork ID Kit allows determining the presence of DNA of Pork (Sus scrofa) in any food.

Pork DNA detection is done by real time PCR using two TaqMan®-MGB probes. One of them, labelled with FAMTM dye, specifically detects one mitochondrial DNA sequence of Pork. The second probe is labelled with VIC® and detects an Internal Positive Control, which is used to rule out inhibitors in the sample and check the correct functioning of the assay.

The detection limit has been calculated upon standard samples consisting of mixtures of raw Pork meat and other species. **imegen-Pork ID Kit** can detect blends containing a percentage below 0.1% [w/w] of Pork meat. The limit of detection in processed samples varies depending on the composition and food processing.

To ensure the representativeness of the results, we recommend the use of a DNA extraction method that allows you to process a large amount of sample [10-20 g]. If you do not have a procedure with these features, we recommend the use of **Food Extraction Kit** [Part No: IMG-262].

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Content and storage of the kit

The kit contains needed reagents to perform the real time PCR reactions:

- Amplification primers for the PCR systems.
- The system contains two probes. One of them labelled with FAM[™] for the specific Pork DNA detection and other labelled with VIC[®] for the IPC detection.
- Pork Positive control (0.1% of Sus scrofa).

The kit contents the necessary reagents to perform 48 reactions:

Reagents	Color	Amount	Storage
Pork Master Mix	Green pad	360 µL	-20°C
General Master Mix	White pad	600 μL	4°C
Positive Control	Green cap	60 μL	-20°C

Table 1. Kit components and storage temperature of imegen Pork ID Kit

Equipment and material required but not supplied

In the following table the equipment requirements for using **imegen-Pork ID Kit** are shown:

Equipment			
1	Real-time PCR Thermal Cycler with channels for detection of FAM TM (520 nm) and VIC [®] (550 nm)		
2	Micropipettes (10 μl, 20 μl and 200 μl)		
3	Table top centrifuge with adaptors for 96 well PCR plates and/or 0.2 ml tubes		
4	Vortex		

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Materials				
1	Optical 96-well reaction plates or 0.2 ml optical			
ı	tubes			
2	Optical adhesive film for 96 well plates or optical			
	adhesive covers for 0.2 ml tubes			
3	Disposable micropipette filter tips			
4	1.5 ml sterile tubes			
5	Powder-free latex gloves			

Detection and quantification limits

The detection limit is the minimal amount of DNA that can be detected. This limit has been estimated using standard manufactured in our laboratory.

imegen-Pork ID Kit can detect blends containing a percentage above 0.01% of Pork DNA. The limit of detection in processed samples varies depending on the composition and food processing.

Specificity

The specificity of the kit was tested through comparison with the NCBI sequence database and was also experimentally tested with success on a collection of reference DNAs. See the results in the table below:







Meat Species	Results
Sheep	ND
Goat	ND
Horse	ND
Buffalo	ND
Donkey	ND
Deer	ND
Pork	D
Wild Boar	D
Chicken	ND
Turkey	ND
Fish	ND
Duck	ND
Ostrich	ND
Goose	ND
Human	ND
Soy	ND
Wheat	ND

Table 2. Specificity of imegen Pork ID Kit

D: Detected

ND: Not Detected

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2. Real time PCR

PCR reactions preparation

imegen-Pork ID Kit is designed to determine, in a single PCR reaction, the presence or absence of Pork DNA and the internal positive control.

We recommend using, the positive control included in this kit for each run.

To estimate the amount of necessary reagents, we recommend make calculations taking into account the number of samples and controls to be simultaneously analysed, and then considering one more reaction, or increase a 10% the volume of each reagent.

The recommended protocol for preparation of amplification reactions is showed below:

- 1. Thaw the Pork Master Mix and the Positive Control vial.
- 2. Vortex each reagent and keep cold.
- 3. Add into a 1.5 mL tube, the following reagents:

Reagents	Amount per reaction
Pork Master Mix	7.5 μL
General Master Mix	12.5 μL

Table 3. Reagents amount per reaction

- 4. Vortex and spin the 1.5 mL tube and dispense 20 μ l per well or tube of 0.2 ml.
- Add 5 μl of each DNA sample at 10 ng/μl, into the appropriate wells. We recommend making each sample analysis in duplicate.
- 6. Add 5 µl of Positive Control and Negative Controls* into the appropriate wells.

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- 7. Cover the well plate with optical film or the tubes with optical cover and spin in the centrifuge.
- * We strongly recommend using an extraction negative control for each run of extractions carried out. This control consists in one tube to which no sample is added and which is summitted to the same extraction process as the other samples. Likewise, we recommended using a PCR negative control for each PCR run; this tube contains no DNA but all PCR reagents.

PCR amplification program

This kit is compatible with the Real-time PCR platforms 7500 FAST, StepOne Real-Time PCR System (Thermo Scientific), and any other Real-time PCR platforms equipped with FAM and VIC channels.

Probes	Receptor	Quencher
Pork-P	FAM TM	MGB
IPC-P	VIC®	MGB

Table 4. Probes information

The following instructions should be taken into account in order to setup the amplification program:

Reaction volume: 25 μL

Targets: FAM and VIC

In case the quencher has to be defined, select MGB for all probes. If the real time PCR system does not take into account the quenchers, select only the receptors [FAM and VIC].

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If the real time PCR system is a 7500 Fast or a StepOne Real-Time PCR system [Thermo Scientific] select Quantitation- Standard curve as a type of experiment and include ROXTM as a reference.

Ramp rate: standard

Optimal program:

Fields	Step 1 Enzyme activation	Step 2 PCR		
Cycle	1 initial cycle	36 cycles		
Number		Denaturation	Primers binding/Extention	
Temperature	95°C	95°C	60°C	
Time 10 minutes		15 seconds	1 minute*	

Table 5. Optimal PCR program

Note: This program has been validated on a StepOne Real-Time PCR System from Applied Biosystems. If you use another brand or model of thermal cycler, you may need the amplification program to be adjusted. Please contact our service department for advice.

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^{*}Fluorescence detection



3. Results analysis

Before analysing the samples results, it should be checked if obtained results in controls are as expected:

- **Positive Control**: The result must always be positive in all amplification reactions, both in the FAMTM channel as VIC[®].
- Negative controls: Amplification must be only detected in the VIC[®] channel. In this channel an internal positive control (IPC) is detected, which determines the absence of inhibition in the sample.

IPC

It must be checked that the IPC (VIC°) is positive in all samples, with a Ct similar to the Positive Control. A negative result in the IPC indicates the presence of inhibitors in the sample. It should be noted that IPC result may be negative in samples where a lot of Pork DNA (FAM^{TM}) is detected, because the PCR reagents are exhausted before amplification of the IPC begins.

Pork

Amplification in the FAM^{TM} channel indicates presence of Pork DNA in the sample.

It is necessary to check if sample Ct is less than the Ct_{cut-off} in order to determine if one reaction of amplification is positive. Any reaction of amplification with Ct upper than Ct_{cut-off} may be considered as negative. The Ct_{cut-off} is equal than the positive control Ct (0.1%) plus 3.32.

$$Ct_{cut-off} = 3.32 + Ct_{Positive\ Control}$$

Note: Any sample with a Ct equal than Ct_{cut-off} contains approximately 0.01% of Sus scrofa DNA.

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In samples where no amplification in the FAMTM channel is seen, we can conclude that no Pork DNA is detected or that its amount in the sample is below than the detection limit.

The following table shows graphically the results that may be obtained from one sample analysis, as well as the interpretation that should be done from the obtained result:

Pork Ma	aster Mix	
Pork	IPC	Interpretation
-	+	No Pork DNA is detected
+	+	Pork DNA is detected
-	-	PCR inhibitors presence in the sample*
+	-	Sample with big amount of Pork DNA

Table 6. Results interpretation

The following table shows graphically the results that may be obtained from the analysis of different assay controls, as well as the interpretation that should be done from the obtained result:

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^{*} If presence of inhibitors in the sample is detected, we recommend checking whether there has been an excess of DNA in the reaction (the recommended maximum is 250 ng). If the amount of DNA is right, we recommend repeating DNA extraction. If the problem persists, please contact our technical department.



	Pork Master Mix		
Controls	Pork	IPC	Interpretation
	+	+	Expected result
Positive control	-	-	PCR Amplification Failure ¹
	-	+	Expected result
Extraction Negative Control	+	+	Contamination in the Pork DNA extraction procedure ²
PCR Negative	-	+	Expected result
Control	+	+	PCR contamination with Pork DNA ³

Table 7. Possible results and their interpretation

Recommendations:

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¹ PCR Amplification Failure: Check amplification program and configuration of fluorescence capture. Amplification failure may be due to a setup technical problem.

² Contamination in the Pork DNA extraction procedure: Contamination may be due to some error made in the process of sample handling, reagents contamination, or environmental contamination. Check DNA extraction protocol, wipe the laboratory where DNA extraction process was performed and take care to avoid any contamination during sample homogenization. If necessary, use new aliquots of the reagents used in DNA extraction.

³ PCR contaminations with Pork DNA: Contamination of PCR reactions may be due to an error made in the process of sample handling, contamination of the reagents or environmental contamination. Thoroughly clean the laboratory where the PCR process was performed, as well as equipment. If necessary, use new aliquots of the reagents used in the PCR. Prepare the PCR reaction containing the Positive Control last to avoid cross contamination.