

Kylt® Erysipelothrix rhusiopathiae

Real-Time PCR Detection









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Revision No.	Amendments
003	changed kit composition (2x qPCR-Mix and Primer-Probe-Mix instead of Reaction-Mix, from batch: 23ER:02 on), new layout for test evaluation
002	Deletion of alternative storage at +2°C to +8°C

A. General

- Kylt® Erysipelothrix rhusiopathiae kits are intended for the specific detection of bacterial DNA of *Erysipelothrix rhusiopathiae* (ER). The kits are suitable for the analysis of samples from birds, swine and ruminants, such as tissues and organs (e.g. liver, heart sac, endocardium, cardiac valve, lung, spleen, cecal tonsils, lymph nodes, kidney, central nervous system), blood and synovia and pure or mixed colony material / isolates derived from cultural processes of the aforementioned samples.
- The qualitative testing with Kylt® Erysipelothrix rhusiopathiae kits is based on a duplex Real-Time PCR: In one reaction setting, the target genes for ER as well as for the exogenous control (Internal Amplification Control (IAC)) are amplified in parallel by respective primer pairs in the Polymerase Chain Reaction (PCR). Amplified target gene fragments are detected via fluorescently labeled probes during the PCR reaction in real-time (Real-Time PCR). The probes specific for detection of amplified ER and the exogenous control target genes are labeled with fluorescent dyes FAM and HEX, respectively, and their emitted fluorescence is separately optically measured by the Real-Time PCR thermal cycler. By means of both individual analyses in one reaction vessel per sample and the Negative Control and Positive Control per run the ER-specific status of a sample can be evaluated in the end. This way, results can be achieved within a few hours after sample receipt.
- These kits were developed for use by trained laboratory personnel following standardized procedures. This Direction For Use must be followed strictly.

B. Reagents and Materials

■ The following Kylt® Erysipelothrix rhusiopathiae kits are available and comprise the following reagents:

Reagent	Colour of Lid	100 Reactions Article No 31770	25 Reactions Article No 31771	Store at
2x qPCR-Mix	Otransparent	4 x 280 μl	1 x 280 μl	≤-18 °C
Primer-Probe-Mix	orange	4 x lyophilizate (final 150 μl each)	1 x lyophilizate (final 150 μl each)	≤ -18 °C
Positive Control	red	4 x lyophilizate (final 50 μl each)	2 x lyophilizate (final 50 µl each)	≤ -18 °C
Negative Control	blue	1 x 1 ml	1 x 1 ml	≤-18 °C

- After receipt, the components are immediately stored at ≤ -18 °C. Avoid repeated freezing and thawing of all the reagents and keep them thawed as short as possible. If occasional processing of few samples only is expected you may prepare appropriate aliquots of reagents before storage at ≤ -18 °C. Prepare aliquots in such a way that freeze-thaw-cycles are reduced to a maximum of three. The Negative Control can alternatively be stored at +2°C to +8°C.
- The components are to be used within the indicated shelf life (see box label). The components of different batches may not be mixed.
- Before its first use, rehydrate the <u>Positive Control</u>: add 50 μ l of Negative Control per vial, briefly incubate at room temperature and mix thoroughly by repeated vortexing. It is recommended to generate aliquots of suitable volumes and store them at \leq -18 °C.
- The <u>Primer-Probe-Mix</u> needs to be stored protected from abundant light. Do not expose to direct (sun)light. Before first use, rehydrate the lyophilized Primer-Probe-Mix: add 150 µl of the Negative Control per vial of Primer-Probe-Mix, briefly incubate at room temperature and mix by pulse-vortexing. Generate aliquots of suitable volumes and store them at ≤ -18 °C.

C. Equipment and Reagents not included

- This detection method can be used on all commercially available Real-Time PCR thermal cyclers that detect the emitted fluorescence of the fluorescent dyes FAM and HEX (emission 520 and 550 nm, respectively). Note that default normalization option against ROX (e.g. using ABI cyclers) must be deactivated.
- Apart from the disposables, the following further devices are needed and are not included in the Kylt® Erysipelothrix rhusiopathiae kits:
 - DNA preparation kit / protocol (e.g. Kylt® DNA Extraction-Mix II or Kylt® RNA / DNA Purification products)
 - Table top microcentrifuge
 - Vortex
 - Micropipettes covering volumes of 1 µl to 1000 µl
 - Centrifuge for PCR tubes or plates
- Accessory Kylt® products: see chapter F "Related and Accessory Products".
- We recommend the exclusive use of certified Nuclease-free disposables as well as powder-free protective gloves. Please wear gloves during the entire experimental procedure. Gloves need to be changed frequently, especially after spillage or suspected contaminations.

D. Control Reactions

- The <u>Positive Control</u> allows for control of the specificity and efficiency of the reagents and the reaction itself, including the performance of the Real-Time PCR and of the Real-Time PCR thermal cycler.
- The <u>Negative Control</u> allows for exclusion of contaminations. The sample testing is only valid if both, Positive and Negative Controls, are used and verified for validity in every Real-Time PCR run.
- The <u>Internal Amplification Control</u> is included in the Reaction-Mix in a defined copy number; it is co-amplified (channel HEX) with every single reaction to detect possible inhibitory effects of the DNA preparation on the Real-Time PCR itself and thus to verify true-negative results.
- If appropriate sampling is unsure we recommend to analyze the samples in parallel with Kylt® Host Cells Real-Time RT-PCR Detection for presence of amplifiable nucleic acids derived from host cell material, see chapter F "Related and Accessory Products".

E. Protocol (see also "Protocol At A Glance" at the end of this Direction For Use)

- The overall protocol of the analysis consists of the following main workflow:
 - 1. Sample Preparation
 - 2. DNA Preparation
 - 3. Reaction Setup and Amplification (Real-Time PCR)
 - 4. Data Analysis Validity and Qualitative Result
- We recommend proceeding through the protocol without interruption to avoid potential degradation of the processed samples and reagents. If necessary, you may store the final DNA preparation at ≤ -18 °C until further processing. Avoid repeated freezing and thawing of the DNA preparations.

1. Sample Preparation

- We recommend <u>pooling</u> of at most five samples or samples from five individuals, respectively, per DNA preparation.
- Tissue and organ samples are homogenized thoroughly in sterile buffer (see above) and a suitable volume is used for the DNA preparation.
- Material derived from cultural processes, i.e. <u>colony material</u>, is directly transferred into respective tubes for Kylt® DNA Extraction (please refer to 2 "DNA Preparation"), such as conical screw cap tube; therefore a little amount of a single colony is picked with a sterile loop wire or sterile pipette tip and transferred to the tube.

2. DNA Preparation

a) Kylt® DNA Extraction (requires Kylt® DNA Extraction-Mix II)

- This method is recommended to be applied for cultural harvests, i.e. pure colony material, only. For detailed information, please refer to the Direction For Use of Kylt® DNA Extraction-Mix II.
- For detailed information on the DNA preparation process, please refer to the Direction For Use or Standard Operating Procedure of the specific kit or in-house method, respectively.

b) Kylt® RNA/DNA Purification products

- All kinds of sample matrices, including pure isolates, tissues and organs may be processed with Kylt® RNA/DNA Purification products (please refer to chapter F "Related Products").
- For detailed information on the DNA preparation process, please refer to the respective Direction For Use.

c) Alternative methods

- All kinds of sample matrices, including pure isolates, tissues and organs may be processed with appropriate DNA preparation kits or appropriate in-house methods.
- For detailed information on the DNA preparation process, please refer to the Direction For Use or Standard Operating Procedure of the specific kit or in-house method, respectively.

3. Reaction Setup and Amplification (Real-Time PCR)

- Before its first use, rehydrate the <u>Positive Control</u> (add 50 μl of Negative Control) and the <u>Primer-Probe-Mix</u> (add 150 μl of Negative Control).
- Before each use, briefly vortex and spin down the 2x qPCR-Mix, rehydrated Primer-Probe-Mix and Negative Control.
- To determine the total number of reactions needed, count the number of samples and add two more for the Negative Control and the Positive Control.
- Prepare the Master-Mix using the components listed below. A larger volume of a ready to use Master-Mix can be prepared and stored at ≤ -18 °C for convenient use over a longer period of time up to the expiry date given on the label. In case of frozen storage the Master-Mix should be aliquoted in such a way that freeze-thaw-cycles are reduced to a maximum of three.
- Vortex, spin down and add 16 μl of the finalized Master-Mix to each of the PCR tubes or plate wells ("cavities").

	Volume (μl)	
Reagent	per Reaction	e.g. n=7
2x qPCR-Mix	10 μΙ	70 μΙ
Primer-Probe-Mix	6 μΙ	42 μΙ
Total Master-Mix	16 µl	112 μl dispense 16 μl per reaction
DNA (Negative Control / sample DNA / Positive Control)	4.0 µl	
Total Reaction	20.0 µl	

- Keep exposure of the 2x qPCR-Mix, Primer-Probe-Mix and prepared Master-Mix to (sun)light as short as possible and return it back to appropriate storage temperature right after application. Avoid the formation of bubbles when pipetting the Master-Mix, samples and controls.
- Add 4 μl of the Negative Control to the corresponding cavity and seal it individually, if possible.

- To minimize risk of potential cross-contaminations, 4 μl of the <u>Positive Control</u> are added to the corresponding cavity after all previous samples and control reactions are set up. Before each use, briefly vortex and spin down the rehydrated Positive Control (see also chapter B "Reagents and Materials").
- If not already done, finally seal the cavities. It is recommended to briefly spin them down before the start of the Real-Time PCR run.
- Place the cavities in the Real-Time PCR thermal cycler and run the test with Kylt® Profile II as given below.

Kylt® Profile II					
Step No	Description	Temperature	Duration		
1	Activation of Polymerase	95 °C	10 min		
2	Denaturation	95 °C	15 sec		
3	Annealing & Extension	60 °C	1 min	42 cycles	
4	Fluorescence Detection	channels FA	J		

- Kylt® Profile II allows for combined run of this and most other Kylt® qPCR detection methods.
- Alternatively, the <u>Kylt® Profile I</u> given below can be applied. Kylt® Profile I allows for combined run of this and most other Kylt® qPCR detection methods as well as Kylt® RT-qPCR detection products that need Reverse Transcription, such as those for detection of viral RNA.

Kylt® Profile I					
Step No	Description	Temperature	Duration		
1	Reverse Transcription	50 °C	10 min		
2	Activation of Polymerase	95 °C	1 min		
3	Denaturation	95 °C	10 sec		
4	Annealing & Extension	0° C	1 min	42 cycles	
5	Fluorescence Detection	channels FA	J		

- In the event of a combined Real-Time (RT-)PCR run, make sure all necessary channels are detected.
- Please follow the specified instructions of your Real-Time PCR thermal cycler as recommended by the manufacturer.

4. Data Analysis - Validity and Qualitative Result

General

■ The amplification data can be processed automatically using the specific software tool of your Real-Time PCR thermal cycler. Alternatively, the threshold can be set manually considering the following directions: The threshold should cross the FAM-curve and the HEX-curve in the linear increase of their slope (log scaling of the y-axis). By setting the threshold, the crossing points with the HEX- and FAM-curves determine the respective cycle threshold (Ct), which is negatively correlated with the initial concentration of copies of the target genes in the Real-Time PCR reaction.

- Only curves with the typical exponential amplification, meaning the curve of the raw data shows a flat baseline at the beginning, followed by a clear (exponential) slope in fluorescence and possibly reaching a plateau-phase (y-axis in log scaling), should be regarded as positive.
- The actual test analysis starts with the validity check of the entire Real-Time PCR run. Afterwards, by means of the Internal Control the validity of each sample reaction and its true test result can be verified according to the Ct-value of the Internal Control channel (HEX). Finally, the *Erysipelothrix rhusiopathiae*-specific status of each sample is analyzed (FAM).

Test Evaluation - Control Reactions

■ The **Real-Time PCR test run** is only **valid** if the curves of the control reactions can be evaluated as follows:

Control Boostions	Channel		
Control Reactions	HEX	FAM	
Negative Control	positive	negative	
Positive Control	positive	positive	

- The FAM-Ct-value of the Positive Control has to be > 15 and ≤ 35 .
- The HEX-Ct-values of the Positive and Negtive Control have to be ≤ 40 .

Test Evaluation - Samples

Target Channel		Signal		
Internal Control	HEX	positive	positive / negative	negative
Erysipelothrix rhusiopathiae	FAM	negative	positive	negative
The sample is <i>Erysipelothrix rhusiopathiae</i>		negative	positive	inhibited

- A sample is negative for *Erysipelothrix rhusiopathiae*, if its HEX-curve is positive ($Ct \le 40$), but its FAM-curve is negative.
- A sample is positive for Erysipelothrix rhusiopathiae, if its FAM-curve is positive (Ct ≤ 42), independent of the HEX-curve.
- **A sample** is **inhibited**, if neither the FAM-curve nor the HEX-curve are positive.
- **Recommendation:** In the case of an inhibited sample the test may be repeated with a dilution of the DNA preparation at e.g. 1:10 (9 volumes Negative Control + 1 volume DNA Extract or eluted DNA). The Negative Control is used as the diluting agent. Preferably, the entire DNA preparation process is repeated using Kylt® RNA/DNA Purification products or appropriate alternative.
- Convenient and reliable sample data entry, Real-Time PCR start, final qualitative analysis and documentation can be conducted with the Kylt® Software, please inquire.

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F. Related and Accessory Products

Product	Article No	Reactions	Description
Kylt® DNA Extraction-Mix II	31398	100	Simplified and economic DNA extraction
Kylt® RNA / DNA Purification	31315	50	Combined RNA and DNA purification from veterinary samples (spin-column based).
Kylt® RNA / DNA Purification HTP	31826	4 x 96	Magnetic bead based combined RNA and DNA purification kit for veterinary diagnostic samples. Suitable for Kylt® Purifier and Kylt® Purifier 48.
Kylt [®] Purifier	31436	1 unit	Purification system for magnetic bead based kits. Up to 96 samples are processed in under 30 minutes. Intended for high-throughput laboratories.
Kylt® Purifier 48	31436	1 unit	Purification system for magnetic bead based kits. Up to 48 samples are processed in under 30 minutes. Intended for low to medium throughput laboratories.
Kylt® Purifier Spin Tips	31434	5 Sets	Plate with 96 separate spin tips, used by the Kylt® Purifer to mix the well contents by stirring. Sufficient for 480 samples.
Kylt® Purifier Plates	31435	20 Plates	Plates to be used for the several reactions and reagents during automated nucleic acid purification. Sufficient for 320 to 480 samples (depending on device and protocol).
Kylt® Host Cells	31106 / 31107	100 / 25	Kit to detect animal host cells; to verify sample taking process.

G. Ordering information

For a fast and efficient service please send your order to *orders.kylt-de@san-group.com* and please provide the following information:

- Delivery and Invoice address
- Purchaser contact telephone number
- End user name and telephone number (if different)
- Purchase order number
- Product name and cataloge number
- Quantity and size of products
- Indicate if your account is VAT exempt

Production:

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Development, manufacturing and distribution of Kylt® In-Vitro Diagnostica is certified according to ISO 9001:2015.



ISO 9001:2015



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PROTOCOL AT A GLANCE

Real-Time PCR Setup

Pulse-vortex and spin down

Vortex

Dispense Master-Mix and add 4 µl NC, DNA preparation, PC

+ 4 µl NC, DNA preparation, PC

+ 16 µl Mix

Seal cavities, spin down (recommended), and start cycler

Analysis

positive sample

compared to threshold

threshold

threshold

baseline

negative sample

phase