thermoscientific

PathoProof Complete-16 kit

INSTRUCTIONS OF USE

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Revision history:

Revision	Date	Description
3	8.1.2018	New manual layout design. Added Applied Biosystem QuantStudio 5 support. Included example picture to dilute samples in manual Large kit extraction. Updated kit component information. Updated ordering information for PCR consumables. Minor text changes.

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1. Introduction

⚠ WARNING! Read the Safety Data Sheets (SDSs) and follow the handling instructions. Wear appropriate protective eyewear, clothing, and gloves. Safety Data Sheets (SDSs) are available from thermofisher.com/support.

Thermo ScientificTM PathoProofTM Complete-16 kit is designed for accurate same-day identification of mastitis-causing microorganisms from bovine milk using real-time polymerase chain reaction (PCR). PCR technology is used for amplifying DNA in a test tube enabling further analysis of DNA. In the PathoProof kits, real-time PCR detects and amplifies the DNA of mastitis-causing microorganisms in a milk sample. The microorganism in question is identified based on its unique DNA. The kit includes all the necessary reagents for bacterial DNA extraction and PCR. The test has been optimized for use with even the most challenging fresh, frozen and preserved milk samples.

Real-time PCR has become the "gold standard" method for food pathogen testing and quality assurance. Based on this advanced technology, PathoProof Complete kits offer several advantages over the conventional culture-based methods:

- Results are obtained substantially faster.
- Risk of carry-over contamination in the laboratory is minimized because the tests are performed in closed reaction vessels that are not opened after the run.
- Fewer "no growth" results because the test identifies and quantifies DNA, so it accurately detects viable, dead and growth-inhibited microorganisms.
- Applicable for use with preserved milk samples, thus eliminating the need for cooling during sample transportation.
- Can be integrated into milk recording programs using preserved milk.

The PathoProof Mastitis Complete-16 kit identifies 15 mastitis-causing microorganisms and the β -lactamase penicillin resistance gene in staphylococci (including Staphylococcus aureus and all major coagulase-negative staphylococci). The microorganisms and the β -lactamase gene are detected in four separate PCR reactions. The PathoProof Mastitis Complete-16 kit identifies the following targets:

- Corynebacterium bovis
- Enterococcus spp. (including E. faecalis and E. faecium)
- Escherichia coli
- Klebsiella oxytoca and/or pneumoniae
- Mycoplasma bovis
- Mycoplasma spp.
- Prototheca spp.
- Serratia marcescens
- Staphylococcus aureus
- Staphylococcal β-lactamase gene
- Staphylococcus spp.
- Streptococcus agalactiae
- Streptococcus dysgalactiae
- Streptococcus uberis
- Trueperella pyogenes and/or Peptoniphilus indolicus
- Yeast

2. Kit components and storage conditions

PF1650 - PathoProof Complete-16 kit is for use with:

- Applied Biosystems[™] QuantStudio[™] 5 Real-Time PCR System, 96-well, 0.1 mL
- Applied Biosystems[™] 7500 Real-Time PCR Systems
- Applied Biosystems[™] 7500 Fast Real-Time PCR Systems

PF1600 - PathoProof Complete-16 kit is for use with the Agilent™ Mx3005P™ Real Time PCR System.

The PathoProof Complete-16 kit contains all the necessary reagents (except ethanol) for DNA extraction and real-time PCR. The kits are stable for six months from the packaging date when stored and handled properly. The tables in Section 2.1 - 2.3 list the components included in the kits.

2.1 KingFisher kit

The Thermo ScientificTM KingFisherTM kit uses the Thermo ScientificTM KingFisherTM Flex instrument for sample preparation. The kit is designed for high throughput laboratories, and/or for laboratories requiring a less hands-on time protocol. The KingFisher kit is sufficient for 4 x 96 reactions when at least 40 samples are analyzed per run.

The PF1650LKF and PF1600LKF kits contain the following components:

Component	Size	Storage conditions
Buffer AW1 ¹	242 ml	
Buffer AW2 ²	130 ml	
Tween® 20 Solution	250 ml	
Buffer AE	110 ml	
Buffer RLT ¹	3x35 ml	Room Temperature
Suspension G2	21 ml	
F-871L Lysis Solution 1 ²	200 ml	
Collection Microtubes (racked)	4x96	
Caps for Collection Microtubes	1x55	
F-872L Lysis Solution 2	4x10 ml	
F-873L Proteinase K	2x12 ml	-20 °C
F-882 PathoProof Master mix	16x1.1 ml	-20 0
F-929 Universal Amplification Standard	350 µl	
F-1651L ³ Primer Mix 1	2x1.1 ml	
F-1652L ³ Primer Mix 2	2x1.1 ml	
F-1653L ³ Primer Mix 3	2x1.1 ml	
F-1654L ³ Primer Mix 4	2x1.1 ml	-20 °C
F-961L ⁴ Primer Mix 1	2x1.1 ml	Protect from light
F-962L ⁴ Primer Mix 2	2x1.1 ml	
F-963L ⁴ Primer Mix 3	2x1.1 ml	
F-964L ⁴ Primer Mix 4	2x1.1 ml	

- 1. Contains chaotropic salt. Not compatible with disinfecting agents containing bleach.
- 2. Contains sodium azide as a preservative.
- 3. Included in PF1650LKF kit for use with Applied Biosystems QuantStudio 5, 7500 and 7500 Fast instruments only. Not compatible with Agilent Mx3005P instrument
- Included in PF1600LKF for use with Agilent Mx3005P instruments only with the following filter sets: FAM[™], Cy5[™], ROX[™], HEX[™]/JOE[™] and ATTO[™]. Not compatible with Applied Biosystems QuantStudio 5, 7500 or 7500 Fast instruments.

2.2 Manual Large kit

The Manual Large kits are designed for laboratories performing DNA extraction on large numbers of samples in each session. The extraction is performed in 8 well strips and in 96 Column Well Plates. The kit is sufficient for 4×96 reactions when at least 20 samples are analyzed per run.

The PF1650L and PF1600L kits contain the following components:

Component	Size	Storage conditions	
96 Column Well Plates	4		
S-Blocks ¹	2		
Collection Microtubes (racked)	4x96		
Elution Microtubes CL (racked)	4x96		
Caps for Collection Microtubes	4x55		
Caps for Elution Microtubes	1x50	Room Temperature	
Microporous Tape Sheet	25 sheets	noom remperature	
Buffer AL ²	2x54 ml		
Buffer AW1 ²	95 ml		
Buffer AW2 ³	66 ml		
Buffer AE	110 ml		
F-871L Lysis Solution 1 ³	200 ml		
F-872L Lysis Solution 2	4x10 ml		
F-873L Proteinase K	2x12 ml	-20°C	
F-882 Master Mix	16x1.1 ml	-20 G	
F-929 Universal Amplification Standard	350 μl		
F-1651L ⁴ Primer Mix 1	2x1.1 ml		
F-1652L ⁴ Primer Mix 2	2x1.1 ml		
F-1653L ⁴ Primer Mix 3	2x1.1 ml		
F-1654L ⁴ Primer Mix 4	2x1.1 ml	-20°C	
F-961L ⁵ Primer Mix 1	2x1.1 ml	Protect from light	
F-962L ⁵ Primer Mix 2	2x1.1 ml		
F-963L ⁵ Primer Mix 3	2x1.1 ml		
F-964L ⁵ Primer Mix 4	2x1.1 ml		

- 1. Reusable, wash or autoclave after use.
- 2. Contains chaotropic salt. Not compatible with disinfecting agents containing bleach.
- 3. Contains sodium azide as a preservative.
- 4. Included in PF1650L kit for use with Applied Biosystems QuantStudio 5, 7500 and 7500 Fast instruments only. Not compatible with Agilent Mx3005P instrument
- Included in PF1600L for use with Agilent Mx3005P instruments only with the following filter sets: FAM[™], Cy5[™], ROX[™], HEX[™]/JOE[™] and ATTO[™]. Not compatible with Applied Biosystems QuantStudio 5, 7500 or 7500 Fast instruments.

2.3 Manual Small kit

The Manual Small kit is suitable for laboratories handling small numbers of samples in each DNA extraction session. The DNA extraction is performed in individual tubes. The kit is sufficient for 50 reactions when at least 12 samples are analyzed per run.

The PF1650S and PF1600S kits contain the following components:

Component	Size	Storage conditions
Spin Columns	50	+2-8°C
Collection Tubes (2 ml)	150	
Buffer AL ¹	12 ml	
Buffer AW1 ¹	19 ml	Room Temperature
Buffer AW2 ²	13 ml	1100III Temperature
Buffer AE	12 ml	
F-871S Lysis Solution 1 ²	30 ml	
F-872S Lysis Solution 2	4x1.3 ml	
F-873S Proteinase K	1.8 ml	-20°C
F-882 Master Mix	2x1.1 ml	-20 C
F-929 Universal Amplification Standard	350 μl	
F-1651S ³ Primer Mix 1	300 μl	
F-1652S ³ Primer Mix 2	300 μl	
F-1653S ³ Primer Mix 3	300 μl	
F-1654S ³ Primer Mix 4	300 μl	-20°C
F-961S ⁴ Primer Mix 1	300 μΙ	Protect from light
F-962S ⁴ Primer Mix 2	300 μΙ	
F-963S ⁴ Primer Mix 3	300 μΙ]
F-964S ⁴ Primer Mix 4	300 μΙ	

- 1. Contains chaotropic salt. Not compatible with disinfecting agents containing bleach.
- Contains sodium azide as a preservative.

 Included in PF1650S kits for use with Applied Biosystems QuantStudio 5, 7500 or 7500 Fast instruments only. Not compatible with Agilent Mx3005P instrument.

 Included in PF1600S kit for use with Agilent Mx3005P instruments only with the following filter sets: FAMTM, Cy5TM,
- ROX[™], HEX[™]/JOE[™] and ATTO[™]. Not compatible with Applied Biosystems QuantStudio 5, 7500 or 7500 Fast instruments.

2.4 Real-time PCR reagent description

2.4.1 PathoProof Complete-16 kit for Applied Biosystems QuantStudio 5, Applied Biosystems 7500 and Applied Biosystems 7500 Fast instrument

- **F-882**: PathoProof Master Mix. PCR master mix in an optimized buffer, containing MgCl₂, deoxynucleoside triphosphates and hot start DNA polymerase.
- **F-929**: PathoProof Universal Amplification Standard. Control DNA for all targets for use as a positive control and for calibration of PathoProof Norden Lab Studio software with the real-time PCR instrument and the reagents.
- **F-1651**: PathoProof Complete-16 kit PCR primer mix for reaction 1, including oligonucleotides for identification of *S. aureus*, *Enterococcus* spp., *C. bovis*, *M. bovis* and an Internal Amplification Control.
- **F-1652**: PathoProof Complete-16 kit PCR primer mix for reaction 2, including oligonucleotides for identification of staphylococcal β-lactamase (penicillin resistance) gene, *E. coli*, *S. dysgalactiae*, *Mycoplasma* spp. and an Internal Amplification Control.
- **F-1653**: PathoProof Complete-16 kit PCR primer mix for reaction 3, including oligonucleotides for identification of *Staphylococcus* spp. (including all relevant coagulase-negative staphylococci), *S. agalactiae*, *S. uberis*, *Prototheca* spp. and an Internal Amplification Control.
- **F-1654**: PathoProof Complete-16 kit PCR primer mix for reaction 4, including oligonucleotides for identification of *Klebsiella* spp., *Serratia marcescens*, *T. pyogenes*, *P. indolicus*, yeast and an Internal Amplification Control.

2.4.2 PathoProof Complete-16 kit for Agilent Mx3005P instrument

- **F-882**: PathoProof Master Mix. PCR master mix in an optimized buffer, containing MgCl₂, deoxynucleoside triphosphates and hot start DNA polymerase.
- **F-929**: PathoProof Universal Amplification Standard. Control DNA for all targets for use as a positive control and for calibration of PathoProof Norden Lab Studio software with the real-time PCR instrument and the reagents.
- **F-961**: PathoProof Complete-16 PCR primer mix for reaction 1, including oligonucleotides for identification of *S. aureus*, *Enterococcus* spp., *C. bovis*, *M. bovis* and an Internal Amplification Control.
- **F-962**: PathoProof Complete-16 PCR primer mix for reaction 2, including oligonucleotides for identification of staphylococcal β-lactamase (penicillin resistance) gene, *E. coli*, *S. dysgalactiae*, *Mycoplasma* spp. and an Internal Amplification Control.
- **F-963**: PathoProof Complete-16 PCR primer mix for reaction 3, including oligonucleotides for identification of *Staphylococcus* spp. (including all relevant coagulase-negative staphylococci), *S. agalactiae*, *S. uberis*, *Prototheca* spp. and an Internal Amplification Control.
- **F-964**: PathoProof Complete-16 PCR primer mix for 4, including oligonucleotides for identification of *Klebsiella* spp., *Serratia marcescens*, *T. pyogenes*, *P. indolicus*, yeast and an Internal Amplification Control.

3. Materials required but not supplied

The materials required but not supplied with the PathoProof Complete-16 kit are listed below. Please contact your local Technical Support team for ordering details.

- · Disposable powder-free gloves
- Ethanol (96–100%); do not use denatured ethanol
- Pipettes
- · Sterile pipette tips with filter
- PCR plates compatible with the real-time PCR instrument
 - o For QuantStudio 5 and Applied Biosystems 7500 Fast system: order catalogue number 4346907 MicroAmp™ Fast Optical 96-Well Reaction Plate
 - For Applied Biosystems 7500 system: order catalogue number AB-1400 96-well PCR Plate
 - For Agilent Mx3005P systems: order catalogue number AB-0600 96 -well PCR Plate
- Optically clear PCR vessel caps: order catalogue number AB-0866 Optically clear flat 8 Cap Strips
- · Vortex mixer
- Microcentrifuge compatible with the spin tubes
- Either 1.5 ml or 2 ml tubes for sample preparation
- · Strip tubes for sample preservation after DNA extraction (optional)
- Real-time PCR instrument compatible with PathoProof kits:
 - o Applied Biosystems QuantStudio 5 Real-Time PCR System, 96-well, 0.1 mL
 - Applied Biosystems 7500 and 7500 Fast Real-Time PCR System with the following dyes calibrated: FAM, CY5, Texas Red, VIC and TAMRA
 - Agilent Mx3005P qPCR System with filter sets for: FAM, CY5, ROX, HEX/JOE and ATTO.

Depending on the extraction platform used, materials or instruments not supplied with the PathoProof Complete-16 kit are listed below. Please contact your local Technical Support team for ordering details.

3.1 KingFisher kits

- KingFisher Flex instrument
- Plate centrifuge capable of 1500 xg
- 96 DeepWell Plate
- KingFisherFlex 96 Tip Comb
- Adhesive Plate Seal for storing eluted samples: order catalogue number AB-0558 Adhesive Sealing Sheets

3.2 Manual Large kits

- Decapping tool
- Thermo Scientific[™] Heraeus[™] Multifuge X3 plate centrifuge
- Two dry bath incubators capable of +37°C and +55°C
- Two dry bath 96 blocks for Collection Microtubes

3.3 Manual Small kits

- Microcentrifuge for 1.5 or 2 ml tubes that is capable of 20 000 xg
- Two dry bath incubators for +37°C and +55°C
- Two or more dry bath blocks for 1.5 ml or 2 ml tubes

4. General guidelines

The following general guidelines should be followed throughout the PathoProof Complete-16 kit protocol:

- Use protective gloves
- · Thaw all frozen reagents thoroughly prior to use
- Mix all solutions well before use
- Spin down reagents after mixing

4.1 Avoiding carryover contamination

Due to their high sensitivity, real-time PCR kits are susceptible to carryover contamination of DNA. The contaminating DNA is typically an amplification product from a real-time PCR run, but can also originate from samples containing high quantities of target bacterial DNA.

The PathoProof Complete-16 kit does not require opening of the real-time PCR vessels once the real-time PCR program has been started. While this kit design significantly reduces the risk of cross-contamination, the following general guidelines should be followed, in addition to other precautions mentioned in this instruction manual, in order to minimize such risks:

- · Designate physically separated working areas for:
 - o DNA extraction (handling of milk samples or other samples containing bacteria)
 - o PCR preparation
- Use different laboratory equipment (disposable gloves, pipettes, pipette tip boxes, vortexes, centrifuges, laboratory coats etc.) in each working area.
- Change gloves frequently and always before leaving an area.
- Use aerosol-resistant pipette tips.
- Use new and/or sterilized plastic ware.
- After starting a real-time PCR run, do not open the real-time PCR vessels under any circumstances.
- Always dispose the real-time PCR vessels into a dedicated, closed trash container and make sure that the vessels do not open accidentally.

5. PathoProof Assay Workflow

This section contains instructions for performing DNA extraction, real-time PCR setup and PCR run. The instructions for performing results analysis with Thermo Scientific™ PathoProof™ Norden Lab Studio™ software are given in the PathoProof Norden Lab Studio software manual. An overview of the laboratory workflow is presented in **Figure 1** below.

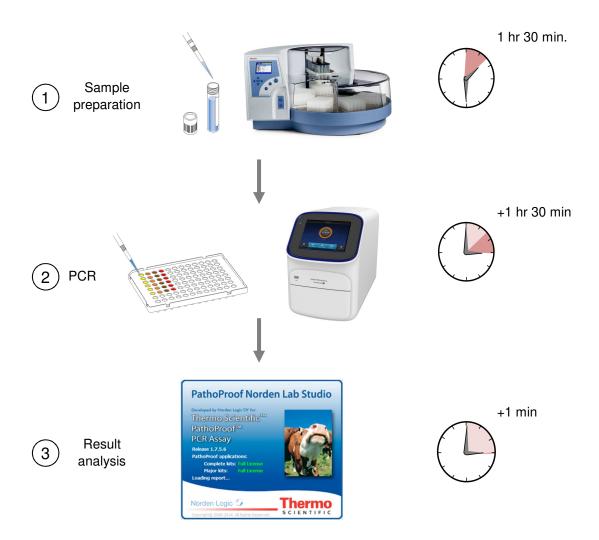


Figure 1. Schematic illustration of the PathoProof assay workflow in three easy steps.

- 1. DNA is extracted from fresh, frozen or preserved bovine milk samples.
- 2. Extracted DNA is used to set up the PCR reactions in a 96-well PCR plate. Four separate PCR reactions are run for each sample.
- 3. The obtained data is interpreted, reported and data-based with Norden Lab Mastitis Studio software.

PathoProof Norden Lab Studio software is an application designed for interpreting, reporting and storing the results obtained using PathoProof kits. This software facilitates data analysis and is highly recommended as an integral part of the procedure for the PathoProof Complete-16 kit.

When using PathoProof kit for the first time, it is necessary to calibrate PathoProof Norden Lab Studio software. Calibration may also be required when changing real-time PCR plastics or when the real-time PCR instrument has undergone maintenance. The calibration runs and experiment runs must be performed using the same real-time PCR instrument, the same type of vessels/plastics, and same optically clear caps for sealing the PCR plate.

Dye calibration is also required for Applied Biosystems[™] 7500 instruments that have not previously been calibrated for dyes FAM, CY5, TAMRA, VIC and TEXAS RED.

5.1 DNA extraction

Before starting:

- Buffers AW1 and AW2 are supplied as concentrates. Before using them for the first time, add the amount of ethanol (96–100%) indicated on the bottles.
- Equilibrate Buffer AE to room temperature.

KingFisher kits only:

- Buffer RLT is provided in 35 ml bottles. To prepare the mixture for the extraction, add 35 ml ethanol (96-100%) and 7 ml Suspension G to the Buffer RLT bottle. Vortex the Suspension G for 3 min before using it for the first time and 1 min before subsequent use. The mixture is stable for one year.
- The 96 Tip Combs are supplied as packages of 2. If opening a new package, store the other 96 Tip Comb within an unused 96 DeepWell Plate.
- Wash and Elution plates can be filled, sealed with Cap Mats or Adhesive Plate Seals and preserved at
 room temperature for one month. If using Adhesive Plate Seals ensure that the rims of the plates are
 dry and clean before sealing. Ensure that no liquid has evaporated from the wells (especially from the
 corner wells) before using the prefilled plates for extraction. Small amount of evaporation (10-25%)
 does not affect the efficiency of the extraction. Prefilled plates should not be shaken or heated.

Manual Large and Small kits only:

• Set two incubators with the proper block inserts at +37°C and +55°C.

5.1.1 DNA extraction for KingFisher kits

1. Prepare eight 96 DeepWell Plates according to the table below.

Plate name	Reagent	Volume per well
Tip Comb Plate	Place Tip Comb plate into 96 DW Plate	-
Sample	Proteinase K F-873L	40 μΙ
Wash 1	Buffer AW1	800 μΙ
Wash 2	Buffer AW1	500 μΙ
Wash 3	Buffer AW2	500 μΙ
Wash 4	Buffer AW2	500 μΙ
Wash 5	Tween® 20 Solution	600 μΙ
Elution	Buffer AE	150 μΙ

- Add reagents only to wells allocated for each sample and the extraction negative control. Example sample layout for a total of 60 samples and negative control is presented in Figure 2.
- Mark the seven 96 DeepWell Plates accordingly from Sample, Wash 1 to Wash 5 and Elution for later identification.

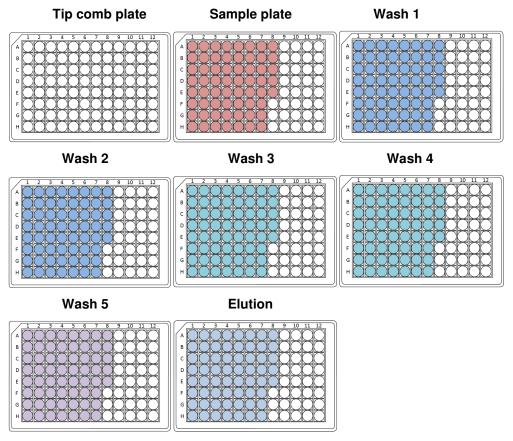


Figure 2. Filling the Thermo Scientific™ KingFisher™ 96 DeepWell plates for KingFisher kit. This setup is an example for 60 samples and 1 extraction negative control.

- After filling the plates, put these plates aside for the sample preparation process (steps 2 5). The
 prefilled plates can be stacked (96 Tip Comb plate at the top) to avoid contamination of the
 extraction reagents.
- 2. Depending on the sample type, prepare samples for centrifugation using the instructions below:
 - For individual quarter milk samples: Add 400 µl of Lysis Solution 1 and 400 µl milk sample into racked Collection Microtubes. Avoid pipetting milk clots into the reaction vessels.
 - <u>For DHI samples:</u> Prepare a fresh mix containing the following volumes per sample: 5 μl Proteinase K and 400 μl Lysis Solution 1. If using a multichannel pipette, add in an extra 1-2 reactions to ensure sufficient volume. Add 400 μl of the Lysis Solution 1/Proteinase K mix and 600 μl milk sample into racked Collection Microtubes.

In addition to the milk samples, include at least one negative control (reagents only) in each DNA extraction run. Seal the tubes using the caps provided for Collection Microtubes.

- 3. Mix well by vortexing or shaking and centrifuge for 3 min at 3000 rpm.
- 4. The presence of fat on top of the supernatant following centrifugation is normal. Remove the supernatant and fat layer on top of supernatant using a pipette without disturbing or removing the pellet. After liquid removal, approximately 50-100 μl of residual fat/liquid on top of the pellet is normal and does not need to be removed.

- 5. Resuspend the pellet carefully in 100 µl Lysis Solution 2 by pipetting up and down and transfer the sample into a well of the sample plate using a pipette.
- 6. Switch on the KingFisher Flex instrument at the power switch. If using the KingFisher Flex instrument through a computer, open the Thermo Scientific™ BindIt™ software from the computer connected to the KingFisher Flex instrument.
- 7. Select the "PathoProof" protocol from the DNA/RNA menu under factory protocols. Press "Start" to start the run.
- 8. Open the cover of the instrument and load the plates into the instrument as indicated on the KingFisher Flex display. After each plate press "Start" and after the worktable has rotated load the next plate. After loading the last plate close the protective cover and press "Start" to initiate the run.

After completion of the sample incubation steps, the extraction protocol pauses for 2 minutes and offers the option to remove the sample plate. Remove the sample plate immediately when the instrument pauses or wait for the full 2 minutes before removing the plate. At the end of the next step, take caution to only add the plate back when the instrument asks for insertion of the sample plate. This will avoid spillages and potential injury from the moving instrument turntable.

9. Take the sample block from the KingFisher instrument, shake the mixture containing Buffer RLT, ethanol and Suspension G thoroughly for 10 s and add 440 µl to each sample in the sample plate. Load the plate back into the instrument when the instrument asks to do so. Press "Start" to continue the run.

If dispensing the mixture into wells with a multichannel dispenser, make sure tips do not touch the well walls or the liquid in the wells.

- 10. After the samples have been processed remove the plates as instructed by the KingFisher Flex display.
 - If the eluted samples are going to be stored, seal the wells of the elution plate with a Cap Mat or an Adhesive Plate Seal.
 - The presence of a small amount of beads in the eluted DNA is normal and does not affect the PCR step.

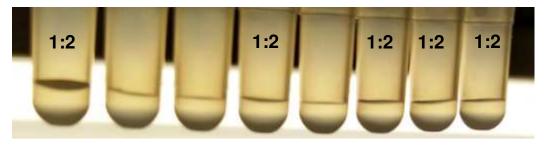
Proceed to Section 5.2 for real-time PCR setup.

5.1.2 DNA extraction for Manual Large kit

Centrifugation of 96 Column Well Plates is performed on the Heraeus Multifuge X3 plate centrifuge.

- 1. Prepare a fresh mix containing the following volumes per sample: 7 μl Proteinase K and 350 μl Lysis Solution 1. If using a multichannel pipette, add in an extra 1-2 reactions to ensure sufficient volume.
- 2. Vortex or shake the milk samples thoroughly. Add 350 µl of Lysis Solution 1/Proteinase K mix and 350 µl milk sample into racked Collection Microtubes. Avoid pipetting milk clots to the reaction vessels. In addition to the milk samples, include at least one negative control (reagents only) in each DNA extraction. Seal the tubes using the caps provided for Collection Microtubes.
- 3. Vortex briefly and incubate at 55°C for 5 min.

- 4. Centrifuge for 5 min at 5000 rpm.
- 5. The presence of fat on top of the supernatant following centrifugation is normal. Remove the supernatant and fat using a pipette without disturbing or removing the pellet. Residual fat/liquid on top of the pellet is normal and does not need to be removed.
- 6. Resuspend the pellet in 100 μl Lysis Solution 2 by pipetting up and down. Seal the tubes using new caps for Collection Microtubes.
- 7. Incubate at 37°C for 10 min.
- 8. Prepare a fresh mix containing the following volumes for each sample: 20 μl Proteinase K and 200 μl Buffer AL. If using a multichannel pipette, add in an extra 1-2 reactions to ensure sufficient volume. Add 220 μl of the mix to each sample, taking care not to wet the rims of the Collection Microtubes. Seal the tubes using new caps for Collection Microtubes.
- 9. Mix thoroughly by shaking vigorously for 15 sec. For efficient lysis, it is essential that the samples and Buffer AL are mixed immediately and thoroughly to yield a homogeneous solution. Hold the racked Collection Microtubes with both hands and shake up and down vigorously.
- 10. Incubate at 55°C for 10 min. Centrifuge briefly at 3000 rpm to remove drops from inside the caps. Stop the centrifuge as soon as it reaches 3000 rpm.
- 11. Add 200 μl ethanol (96–100%) to each tube. Seal the tubes using new caps for Collection Microtubes. Shake vigorously for 5 sec. Centrifuge briefly at 3000 rpm to remove drops from inside the caps. Stop the centrifuge as soon as it reaches 3000 rpm.
- 12. Place the 96 Column Well Plate on top of an S-Block. Seal unused wells with Microporous Tape Sheet and mark the column numbers for later identification in the next extraction.
- 13. Remove the possible viscous clots from the mixture prepared in step 11 using a pipette. Carefully apply the supernatant to the 96 Column Well Plate. Take care not to wet the rims of the wells to avoid aerosol formation during centrifugation.
- 14. Seal the 96 Column Well Plate with Microporous Tape Sheet. Load the S-Block and 96 Column Well Plate onto the carrier. Place the carrier in the rotor bucket and centrifuge at 6000 rpm for 4 min.
- 15. Place the 96 Column Well Plate on top of an empty S-Block. Remove the tape. Carefully add 500 μ l Buffer AW1 to each well. Seal the 96 Column Well Plate with a new Microporous Tape Sheet. Centrifuge at 6000 rpm for 4 min.
- 16. Place the 96 Column Well Plate on top of an empty S-Block. Remove the tape. Carefully add 500 μ l Buffer AW2 to each well. To ensure evaporation of residual ethanol in the following centrifugation step, do not seal the plate. Centrifuge at 6000 rpm for 15 min.
- 17. Place the 96 Column Well Plate on top of a rack of Elution Microtubes. To elute the DNA, add 100 μl Buffer AE to each well. Seal the 96 Column Well Plate with a new Microporous Tape Sheet, and incubate for 1 min at room temperature. Centrifuge at 6000 rpm for 4 min.
- 18. Inspect the tubes and samples. If samples can be seen to have a concave meniscus or show an uneven elution volume, dilute the eluted samples using AE buffer or distilled water.



19. Seal the wells using the caps for Elution Microtubes.

Proceed to Section 5.2 for real-time PCR setup.

5.1.3 DNA extraction for Manual Small kits

Centrifugation of Spin Columns is performed using a Microcentrifuge capable of 20 000 xg.

- 1. Prepare a fresh mix containing the following volumes per sample: $7 \mu I$ Proteinase K and 350 μI Lysis Solution 1. Add in an extra 1 2 reactions to ensure sufficient volume.
- 2. Vortex or shake the milk samples thoroughly. Add 350 μl of Lysis Solution 1/Proteinase K mix and 350 μl milk to each reaction vessel. Avoid pipetting milk clots into the reaction vessels. In addition to the milk samples, include at least one negative control (reagents only) in each DNA extraction.
- 3. Vortex briefly and incubate at 55°C for 5 min.
- 4. Centrifuge for 5 min at 5000 xg.
- 5. The presence of fat on top of the supernatant following centrifugation is normal. Remove the supernatant and fat using a pipette without disturbing or removing the pellet. Residual fat/liquid on top of the pellet is normal and does not need to be removed.
- 6. Resuspend the pellet in 100 μl Lysis Solution 2 by pipetting up and down.
- 7. Incubate at 37°C for 10 min.
- 8. Prepare a fresh mix containing the following volumes per sample; 20 μl Proteinase K and 200 μl Buffer AL. Add in an extra 1 2 reactions to ensure sufficient volume. Add 220 μl of the mix to each reaction vessel. Mix by vortexing for 5 10 sec.
- 9. Incubate at 55°C for 10 min. Briefly centrifuge the reaction vessels to remove drops from inside the caps.
- 10. Add 200 μl ethanol (96–100%) to each sample and mix by pulse-vortexing for 15 sec. It is essential that the sample, the Buffer AL and the ethanol are mixed thoroughly to yield a homogeneous solution. Do not use alcohols other than ethanol, as this may result in reduced DNA yields. After mixing, briefly centrifuge the reaction vessels to remove drops from inside the caps.
- 11. Remove the possible viscous clots from the mixture using a pipette. Then, carefully apply the supernatant to the Spin Column (placed inside a 2 ml collection tube) without wetting the rim. Close each Spin Column to avoid aerosol formation during centrifugation. Centrifuge at 20,000 xg (~14,000 rpm) for 1 min. Place the Spin Column in a clean 2 ml Collection Tube (provided), and discard the tube containing the filtrate.
- 12. Carefully open the Spin Column and add 500 μl Buffer AW1 without wetting the rim. Close the cap and centrifuge at 20,000 xg (~14,000 rpm) for 1 min. Place the Spin Column in a clean 2 ml Collection Tube (provided), and discard the Collection Tube containing the filtrate.
- 13. Carefully open the Spin Column and add 500 μl Buffer AW2 without wetting the rim. Close the cap and centrifuge at 20,000 x g (~14,000 rpm) for 3 min.
- 14. Place the Spin Column in a clean 1.5 ml microcentrifuge tube (not provided), and discard the Collection Tube containing the filtrate. Carefully open the Spin Column and add 100 μl Buffer AE. Incubate at room temperature for 1 min, and then centrifuge at 20,000 x g (~14,000 rpm) for 3 min to elute the DNA. Discard the Spin Column.

Proceed to Section 5.2 for real-time PCR setup.

5.2 Real-time PCR setup

Throughout the procedure, follow the general guidelines detailed in Section 4.

Important notes before starting:

- Use at least one negative (no template) PCR control for each of the four PCR solutions in each real-time PCR run.
- If all the samples are expected to be negative for the targets, it is advisable to include the PathoProof Universal Amplification Standard and/or a positive DNA extraction control (such as a milk sample previously tested positive with PathoProof Complete-16 kit) in each real-time PCR run.
- The experiment runs must be performed using the same real-time PCR instrument, the same type of vessels and the same sealing method (optically clear caps) that was used when calibrating the PathoProof software (see PathoProof Norden Lab Studio software manual).
- The purified DNA can be stored at +5°C for a few days and for longer periods at -20°C.
- 1. Vortex briefly the thawed PathoProof Master Mix and PathoProof Primer Mixes 1- 4 and spin down.
- 2. Prepare four separate PCR solutions by combining PathoProof Master Mix and PathoProof Primer Mixes 1, 2, 3 and 4 in four separate microcentrifuge tubes. Use the formula on the next page to calculate the volumes required. The formula provides excess volume to compensate for volume loss due to reagent pipetting steps.

PCR solution	Reagent	Volume
PCR solution 1	PathoProof Master Mix	Ν x 11 μl
FCA Solution 1	PathoProof Primer Mix 1	Ν x 5.5 μΙ
PCR solution 2	PathoProof Master Mix	Ν x 11 μl
FOR SOLUTION 2	PathoProof Primer Mix 2	N x 5.5 μl
PCR solution 3	PathoProof Master Mix	Ν x 11 μl
FOR SOLUTION 3	PathoProof Primer Mix 3	Ν x 5.5 μl
PCR solution 4	PathoProof Master Mix	Ν x 11 μl
L Solution 4	PathoProof Primer Mix 4	Ν x 5.5 μΙ

N = Number of samples including:

- Negative PCR control (necessary in every run)
- PathoProof Universal Amplification Standard (positive control, optional)
- Negative DNA extraction control (recommended)
- Positive DNA extraction control (optional)
- DNA from extracted milk samples
- 3. Vortex the PCR solutions briefly and spin down.
- 4. Add 15 μl of the PCR solutions 1, 2, 3 and 4 into the four wells allocated for each sample. **Figure 3** shows an example for allocating PCR solutions 1 4 for five samples and one negative control.

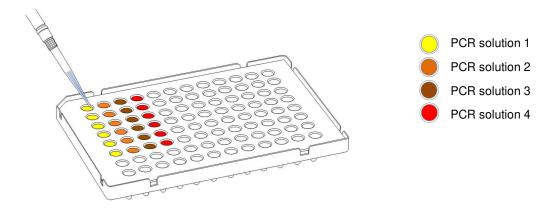


Figure 3. Example setup for pipetting PCR solutions in Complete-16 kit.

Always add PCR solution 1 to plate columns 1, 5 and 9, PCR Solution 2 to columns 2, 6 and 10, PCR solution 3 to columns 3, 7 and 11 and PCR solution 4 to columns 4, 8 and 12. This order must be maintained for correct data analysis when using the PathoProof Norden Lab Studio software.

 Add 5 μl of the eluate from the DNA extraction protocol (Section 5.1) into the four wells allocated for each sample. Figure 4 shows an example for pipeting 5 samples and negative control into previously prepared PCR solution plate.

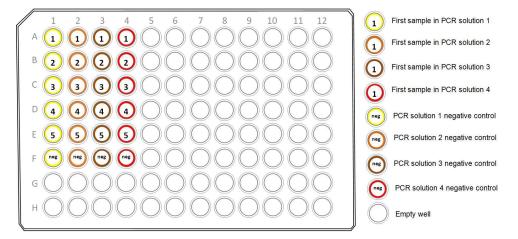


Figure 4. Example setup for 5 samples and one Negative PCR control. Negative PCR control reactions are always allocated to the last available wells.

- a. If the PathoProof Universal Amplification Standard is included in the test setup, add 5µl of the standard into four consecutive wells.
- b. In the wells allocated for negative (no template) controls, add 5µl of sterile water or Elution Buffer (AE).
- 6. Close the 96-well PCR plate with **optically clear caps** and centrifuge using a suitable plate centrifuge (3000 rpm, 5 sec).

Note: It is critically important to use the optically clear caps. Do not use adhesive seal or heat seal to close the PCR plates as performance of the assay will be adversely affected.

Proceed to instrument-specific instructions in Section 5.3.

5.3 Real-Time PCR instrument settings, PCR run and data export

Instrument specific template files needed for PathoProof calibration and sample runs will be installed into the PathoProof Norden Lab Studio directory and a link to these files will be created onto the desktop during the software installation process. PathoProof software (and the instrument specific template files) is available to download at www.nordenlogic.com.

5.3.1 Applied Biosystems QuantStudio 5 Real-Time PCR System

1. Open template file

Open PathoProof template file for QuantStudio 5 instrument. Wait for instrument initialization.

2. Load the plate

Open tray by selecting tray icon from instrument touch screen in the instrument and load the plate. Close the tray by selecting the tray icon again.

3. Name samples

Name the samples as needed (samples may be named after the run) by selecting Plate-section and then selecting Advanced Setup. PathoProof template files have sample place holder names (similar to Figure 4) that can be renamed.

Select wells that are not going to be used and then select "Clear" from right click options.

Optional: Sample names can be imported into run template file by selecting $File \rightarrow Import\ Plate\ Setup$ and then searching Sample layout file generated with Sample layout tool. Assay specific Sample layout tools have been saved to each assay template file folder.

4. Start the run

To start the sample run, select **START RUN** -button in the Run tab, select the instrument from drop down menu and input run file name to save the file. Run is initialized once save-button has been selected.

5. Export raw data for the PathoProof software

After the real-time PCR run, select **Export**.

- 1. Update the export file name from **File Name** -field if needed.
- 2. Ensure that the file type is QuantStudio and file extension is .xls
- 3. Next to **Location**, search preferred file saving location.
- 4. Make sure that the Sample Setup and Amplification Data -boxes have been selected under Content.
- 5. Make sure to select Unify the above content into one file under Options.
- 6. Generate export file from the run by selecting **Export**.

Follow the instructions in your PathoProof Norden Lab Studio software manual in order to import the data into PathoProof Norden Lab Studio software.

5.3.2 Applied Biosystems 7500 Real-Time PCR System

PathoProof assays are only compatible with instrument software versions 2.0.6 and above.

Before using the PathoProof assay, the Applied Biosystems 7500 or 7500 Fast Real-Time PCR System needs to be calibrated for the following dyes:

- FAM
- VIC
- TAMRA
- CY5
- TEXAS RED

If the Applied Biosystems 7500 or 7500 Fast Real-Time PCR System does not have these dyes calibrated, please contact your local Technical Support team to order Spectral Calibration Kits I and II for ABI7500 systems.

1. Open template file

Open PathoProof template file for ABI7500 instrument. Wait for instrument initialization.

2. Load the plate

Push the tray door to open it and load the plate. Close the tray door.

3. Name samples

Name the samples as needed (samples may be named after the run) by selecting **Setup** -section on left and then **Plate Setup**. PathoProof template files have sample place holder names (similar to Figure 4) in section **Define Samples** that can be renamed. Select wells that are not going to be used and then select "Clear" from right click options.

Optional: Sample names can be imported into run template file by selecting **File** \rightarrow **Import Plate Setup** and then searching Sample layout file generated with Sample layout tool. Assay specific Sample layout tools have been saved to each assay template file folder.

4. Save the run

Save the file in a separate location and with different file name for later identification and to ensure, that the default PathoProof template file is not over written.

5. Start the run

Start the run by clicking the **Start run** -button in the Setup Screen.

6. Export raw data for the PathoProof software

After the real-time PCR run, choose Analysis option from the left side of the screen.

Select Export.

- Make sure that the Sample Setup and Amplification Data -boxes have been selected next to 1. Select data to export.
- Make sure that One file is selected next to 2. Select one file or separate files
- 3. Update the Export File Name if needed and ensure that the file extension is .xls from section 3. Enter export file properties
- Select Open file(s) when export is complete and Save current settings as default -check boxes
- 5. Generate export file from the run by selecting **Start Export** and Follow the instructions in your PathoProof Norden Lab Studio Software manual in order to import the data into PathoProof Norden Lab Studio software.

5.3.3 Agilent Mx3005P qPCR System

PathoProof Complete-16 kit is only compatible with instrument that has the following filter sets:

- FAM
- CY5
- ROX
- HEX/JOE
- ATTO

If your Agilent instrument does not have these filter sets, please contact Thermo Fisher Scientific technical support: microbiology.techsupport.uk@thermofisher.com.

Before the first run and after instrument maintenance, verify that the Filter Gain Settings are as follows (from the Instrument selection -> Filter Set Gain Settings):1x for CY5, 4x for ATTO, 1x for ROX, 2x for HEX/JOE and 4x for FAM

1. Open template file and switch lamp on

Open PathoProof template file for Mx3005P instrument. Wait for instrument initialization. Switch the instrument's lamp on by clicking the lamp icon.

2. Name samples

Name the samples as needed (samples may be named after the run if preferred).

3. Clear empty wells

Select the empty wells and click "Clear Selected Wells" in the panel at the right.

4. Save the run

Save the file in a separate location and with different file name for easy identification and to ensure, that the PathoProof template file is not over written.

5. Load the plate and Start the run

Open instrument door and pull from the handle of heat lid tray. Insert the PCR plate under the heat lid and push the heat lid back to it's original location. Close the instrument door.

Start the run by clicking the **Start run** -button in the Thermal Profile Setup.

6. Export raw data for the PathoProof software

Generate export file from the run by:

- Click the Analysis button, then the Results tab.
- · Select File menu from the top left.
- Select Export Chart Data → Export Chart Data to Text file → Format 1 -- Vertically Grouped by Plot...

Now follow the instructions in your PathoProof Norden Lab Studio software manual in order to import the data into PathoProof Norden Lab Studio software.

6. Troubleshooting

Possible problems in real-time PCR are presented first, followed by problems in DNA extraction.

6.1 DNA extraction

This table presents explanations for the following problems: little or no DNA was obtained from the milk samples, or the extracted DNA is contaminated with inhibitory substances.

Possible explanation(s)	Recommended action
Inefficient lysis due to reduced Proteinase K activity.	Be sure to store the Proteinase K solution at -20 °C or, when in use, on ice. Always be sure to prepare fresh mixes of Proteinase K with Lysis Solution 1 and with Buffer AL. Repeat the DNA purification procedure with a new sample.
Inefficient cell lysis due to insufficient mixing of the sample with Buffer AL.	Repeat the DNA purification procedure with a new sample. Be sure to mix the sample and Buffer AL immediately and thoroughly by pulse-vortexing.
No ethanol or incorrect kind of ethanol added to the lysate before loading onto the column or the 96 Column Well Plates.	Repeat the purification procedure with a new sample. Use 96–100% ethanol. Do not use denatured ethanol, isopropanol or lower percentage ethanol.
Buffer AW1 or AW2 prepared incorrectly.	Repeat the purification procedure with a new sample. Ensure that the Buffer AW1 and AW2 concentrates were diluted with the correct volumes of 96–100% ethanol as indicated on the bottles. Do not use denatured alcohol, isopropanol or lower percentage ethanol.
Buffers AW1 and AW2 used in the wrong order.	Ensure that Buffers AW1 and AW2 are used in the correct order in the protocol. Repeat the purification procedure with a new sample.
	Ensure that the Spin Column or the 96 Column Well Plate does not come into contact with the filtrate prior to elution.
Residual Buffer AW2 in the eluate.	Before DNA elution, ensure that the 96 Column Well Plate is centrifuged without the Microporous Tape Sheet in order to allow ethanol to evaporate from the sample.
Spin Column or 96 Column Well Plate not incubated at room temperature (15–25 °C) for 1 min before elution.	After the addition of Buffer AE, the Spin Column or the 96 Column Well Plate should be incubated at room temperature for at least 1 min.
DNA not eluted efficiently.	To increase elution efficiency, pipet Buffer AE onto the center of the spin column or the 96 Column Well Plate and incubate for 5 min at room temperature before centrifugation.

6.2 Real-time PCR

This table presents explanations for the following problems: IAC Ct values are not within acceptable range, positive target amplification signal in negative control, acceptable IAC Ct in all samples but no positive results, false positive signal read in most of the samples

Problem	Possible explanation(s)	Recommended action	
The Ct values of the Internal	Missing reagents in the PCR setup.	Repeat the real-time PCR	
Amplification Controls are not within the acceptable range in samples* and in negative controls**.	Incorrect volume of master mix and/or primer mix in wells.	Make sure that the correct amounts of primer mix and master mix are added into the correct wells.	
Unacceptable IAC amplification signals for all samples* but acceptable IAC signals for the Negative controls.	PCR inhibitors originating from the DNA extraction are present in the test samples.	Refer to Chapter 5.1 "DNA extraction".	
Unacceptable IAC amplification signals for all parallel reactions of	Missing reagents in the wells of the sample or incorrect volume of reagent.	Repeat the real-time PCR for the sample.	
one sample* but acceptable IAC signals in the other samples and the Negative controls.	The PCR inhibitor concentration of the sample is too high.	Dilute the extracted DNA sample (use 1:5 and 1:10 dilutions for example) and repeat the real-time PCR. See Chapter 5.1 "DNA extraction".	
	Missing reagents in the well concerned or incorrect volume of reagent.	Repeat all parallel real-time PCR reactions of the sample.	
Unacceptable IAC amplification signals for one reaction of one sample* but acceptable IAC	Bacterial DNA is present in a high quantity (+++) in the sample (confirm the presence of bacterial DNA by inspecting the amplification curves).	No action required.	
amplification signals for the other parallel reactions of the sample, the other samples and the Negative controls.	PCR inhibitors originating from the DNA extraction are present in the test samples at such high concentration that they inhibit the amplification of less robust IAC (the higher the inhibitor concentration, the more IACs will fail starting from primer mix 1)	Dilute the extracted DNA sample (use 1:5 and 1:10 dilutions for example) and repeat the real-time PCR for all parallel reactions of that sample. See Chapter 5.1 "DNA extraction".	
Unacceptable IAC amplification signals** for one reaction of Negative	Missing reagents in the wells of the negative control.	No action required because the IACs in the	
control or all parallel reactions of Negative control but acceptable IAC amplification signals in sample wells.	Incorrect volume of master mix and/or primer mix in wells.	samples are acceptable	
Positive bacterial target amplification signals in wells for Negative control.	A carryover contamination may have occurred in the laboratory.	Take precautions detailed in Section 4.1 to minimize contaminations.	
Acceptable IAC amplification signals in all wells, but all targets negative.	Failed DNA extraction.	Refer to Chapter 5.1 "DNA extraction". Include PathoProof Amplification Standard and/or a positive DNA extraction control, such as a milk sample previously tested positive with PathoProokit.	
	There were no bacteria present in the samples.		
Amplification signal is not smooth and/or is linear	Adhesive seal or heat seal was used in closing PCR plate. Heating protocol used in PathoProof will stretch the seal on well, which may result in abnormal signal reads.	Refer to section 4.3 in PathoProof Norden Lab Studio IFU to interpret and exclude false positive signal reads from sample. To avoid false positive signal reads, use optically clear caps	

^{*} If the Ct values of the Internal Amplification Controls are not within the acceptable range, PathoProof Norden Lab Studio displays a warning icon () beside the sample name in the run viewer and in the upper left corner of the sample viewer. Additionally, the word "Failed" appears after the IAC Ct values in the sample viewer.

^{**} If the Ct values of the Internal Amplification Controls are not within the acceptable range in the negative controls, PathoProof Norden Lab Studio displays a warning icon () beside the run name in the database view and in the upper left corner of the negative control viewer. Additionally, the word "Failed" appears after the IAC Ct values in the negative control viewer

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