

PathoSEEK[®] Listeria Detection Assay v2 with MaGiC Lysis Kit

User Guide v2

Real Time PCR (qPCR) assay for the detection of *Listeria monocytogenes* in dried cannabis flower and infused products matrices

Table of Contents

Introduction	3
Process Overview	3
Limit of Detection	4
Kit Components	4
Shelf Life and Storage	5
Required Equipment and Supplies	5
Safety Precautions and Recommendations for Best Results	6
Intended User	7
Sample Preparation	7
Sample Lysis	9
MaGiC Lysis for Flower, Gummy Infused Products, Chocolate and Oil Extracts	9
Real-Time Quantitative PCR (qPCR) Setup Protocol with PathoSEEK Amplification Mix	10
Running the BioMolecular Systems MIC	13
Data Analysis with BioMolecular System MIC	13
Running the Agilent AriaMX	16
Data Analysis: Agilent AriaMX	19
PathoSEEK® Listeria Assay v2 Data Analysis Quick Reference Table	19
Running the BioRad CFX96	25
Data Analysis BioRad CFX96	30
PathoSEEK® Listeria Data Analysis Quick Reference Table	30
Troubleshooting Guide	32
Glossary and Definitions	33
References	33
Revision History	34
DISCLAIMER	34

Introduction

Current regulations require cannabis flower and cannabis products to be free of *Listeria monocytogenes*. The PathoSEEK® Listeria Assay v2 with MaGiC Lysis Kit will detect this species using a qPCR (Quantitative Polymerase Chain Reaction) assay in cannabis flower, cannabis concentrates, infused edibles and infused non-edibles.

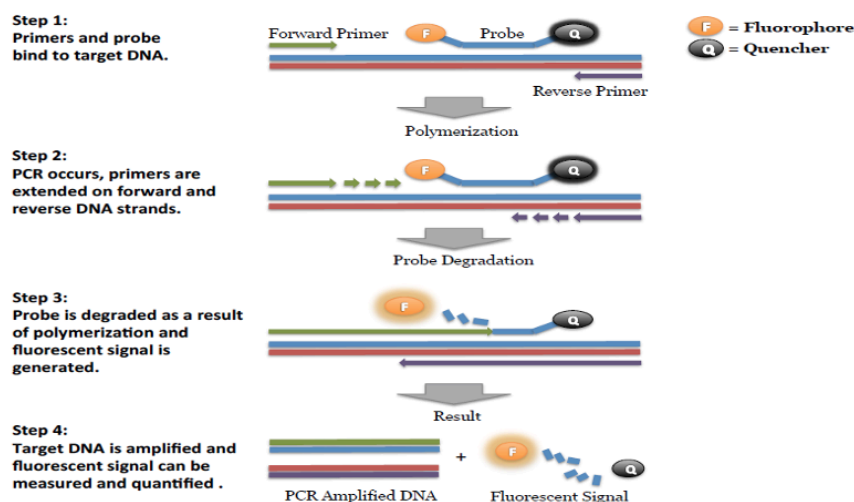
Process Overview

The PathoSEEK® Listeria Detection Assay v2 employs a multiplexing strategy with an internal cannabis control (IC) that is introduced at the lysis step to ensure accurate detection of *Listeria monocytogenes* and internal control DNA in every reaction. Below is a simplified depiction of the MaGiC lysis in Figure 1 and qPCR assays in Figure 2.

Figure 1: Overview of MaGiC Lysis



Figure 2: Overview of qPCR



Limit of Detection

The method is able to detect down to 1 CFU per test portion.

Kit Components

PathoSEEK® Listeria Detection Assay v2 with MaGiC Lysis Kit, **P/N 420527**

(Kit contains sufficient reagents for 200 reactions)

Component Name	Qty Provided	Storage Conditions
MaGiC Lysis Reagent	1 Bottle (12 mL)	RT
MaGiC Stabilization Buffer	1 Bottle (24 mL)	RT
PathoSEEK Amplification Mix <i>Includes 2 tubes nuclease free water for resuspension</i>	4 Vials (50 rxns/each)	RT / -20 °C*
PathoSEEK Listeria Assay v2	1 Tube (200 µL)	-20 °C

Note: Actual fill volumes include overage

*The PathoSEEK Amplification Mix can be stored lyophilized at Room Temperature for up to 2 years. Once re-hydrated it must be stored at -20 °C for up to 3 months.

Additional Required Reagents Not in Kit:

Item P/N	Item Name	Qty Provided	Storage Conditions
420226	Listeria Enrichment Broth (via Hardy Diagnostics)	225 mL (Pack of 12)*	2-8°C
420337	Internal Cannabis Control	1 Tube (50 µL)	-20 °C
420385	PathoSEEK® Listeria Positive Control	1 Tube (50 µL)	-20 °C
420184	PCR Grade Water	500 mL Bottle	2-25°C

*Listeria Enrichment Broth is a Made-To-Order product that requires a minimum order quantity as listed above. The lead time between date of order and ship date is approximately 3-4 weeks.

Shelf Life and Storage

Once received, each kit component must be stored at its designated storage condition. Reagents stored properly can be used until the expiration date indicated on each component label.

Required Equipment and Supplies

Equipment:

- Bio Molecular Systems Mic 4-Channel PCR Instrument - **Medicinal Genomics P/N 420241**
 - BMS supplied or lab supplied Windows PC
 - Mic Tubes and Racked Caps - **Medicinal Genomics P/N 420244**
 - Mic Tubes and Caps (Bulk) - **Medicinal Genomics P/N 420243**
- Agilent AriaMx Real-Time PCR System G8830A—Containing the following Optical Channels: FAM and HEX. **Medicinal Genomics P/N 420387**
 - Agilent HP Notebook PC option 650 or lab supplied Windows PC
 - Optical Strip Caps.—Agilent #401425.

Note: If using adhesive seals instead of strip caps, use Applied Biosystems MicroAmp Optical Film Compression Pad, Fisher Scientific, #43-126-39 to prevent evaporation and cross contamination between wells.

- Bio-Rad CFX96 Touch™ Real-Time System.
 - Bio-Rad supplied or own Windows PC
- Adjustable, variable volume pipettes (single or multichannel).—P10, P20, P200, and P1000
- Adjustable, variable volume filter pipettes tips.—For P10, P20, P200, and P1000
- 1.5 mL Tube Benchtop Cryogenic Rack— VWR #89004-558 or equivalent
- Freezer—Capable of maintaining -20 ± 2 °C
- Tabletop Mini Plate Centrifuge—Fisher Scientific #14-100-143 or equivalent
- Tabletop Mini Centrifuge—VWR #10067-588 or equivalent
- Vortex-Genie Pulse—Scientific Industries, SKU: SI-0236 or equivalent
- Incubator—Capable of maintaining 37 ± 2 °C, VWR #97025-630 or equivalent.
- 96-Well PCR Plate Cryogenic Rack (optional) —VWR #89004-570

- 1.5 µL Tube Benchtop Cryogenic Rack (optional if crushed ice is not an option) — VWR #89004-558 or equivalent
- Eppendorf Tube Rack
- Scientific Scale—Capable of measuring to milligrams
- Refrigerator—Capable of maintaining 2–8 °C

Supplies:

- 96-well Optical qPCR plate — **Medicinal Genomics P/N 100164**
- Adhesive optical seal for qPCR plates — **Medicinal Genomics P/N 100177**
- Filter Bags—**Medicinal Genomics P/N 100008** (Whirl-Pak #B01385WA)
- Crushed ice
- Beaker or Solo Cup. (optional)
- 1.5 mL Eppendorf Tubes
- 15 mL or 50 mL conical tubes
- Pipette - Aid Portable Pipetting device for serological pipettes, VWR 89166-464 or equivalent
- 25mL Serological Pipette—VWR 89130-890 or 89130-900 or equivalent
- 10% bleach

Safety Precautions and Recommendations for Best Results

Environment

The quality of results depends on the strict compliance with Good Laboratory Practices (for example, the EN ISO 7218 standard), especially concerning PCR:

- Never circulate lab equipment from one workstation to another
- Always use a positive and negative control for each series of amplification reactions
- Periodically verify the accuracy and precision of pipette, as well as correct functioning of the instruments
- Change gloves often, especially if you suspect contamination
- Clean workspaces periodically with 10% bleach and other decontaminating agents
- Use powder-free gloves and avoid fingerprints and writing on tube caps. Both can interfere with data acquisition.

Safety Precautions

- *Listeria monocytogenes* is a Gram positive bacterium that is ubiquitous in the environment and can be found in soil, water, and air (1) (2). *Listeria monocytogenes* is known to cause food-borne disease in humans called Listeriosis. Ingestion may not always result in illness and the severity of disease varies from person to person (1). Immunocompromised individuals are at greater risk for developing listeriosis and experiencing negative outcomes like severe illness or death (1). Listeriosis may lead to other diseases like septicemia and meningitis and in the United States *Listeria monocytogenes* is the 5th leading cause of bacterial meningitis (2). Listeriosis may be treated with the use of antibiotics but approximately 30% of cases are fatal in individuals defined as at-risk (2).
- Assay users should observe standard microbiological practices and safety precautions when performing this assay. Wear protective gloves, lab coats, eye/face protection as indicated by your quality system.
- It is the responsibility of each laboratory to handle waste and effluents processed according to their nature and degree of hazardousness and to treat and dispose of them in accordance with applicable local, state, and federal regulations.

Intended User

The PathoSEEK® Listeria Assay v2 with MaGiC Lysis Kit and all components required to perform the method are intended for use by trained personnel familiar with laboratory techniques associated with pathogenic organism detection.

Sample Preparation

1. Aliquot Listeria Broth (LB).

Note: LB is a very good growth medium for microbes. Therefore, it is best to pour the approximate amount of LB into another sterile tube or container so as to not contaminate the stock bottle.

2. Wipe down the workspace with a 10% bleach solution, including the bench top and all equipment being used.

3. If not already at room temperature, remove LB from the 2-8 °C refrigerator (it should come to room temperature, 20–28 °C, before use).
4. Prepare consumables. Label all the filter bags with “[sample name] [date]”.
5. Label lysis plate with date.
6. Before weighing out the sample to be tested, make sure that the entire sample is broken up and thoroughly homogenized. A well-homogenized sample will ensure more accurate testing.
7. *Cannabis flower, n grams* —Weigh flower sample material into one side of the mesh liner inside the Whirl-Pak bag. Add $9 \times n$ mL of LB to each test portion. Close the Filter bag by folding the top over three times. Mix for 1 minute by hand. Incubate for 16 - 24 h at 37 ± 2 °C.
8. *Infused products and concentrates, n grams.* — Weigh the Infused Product matrix into a 15 mL conical tube, 50 mL conical tube, or Whirl-Pak bag depending on Infused Product volume. Add $9 \times n$ mL of LB to each test portion. Vortex or homogenize sample and LB. Incubate for 16 - 24 h at 37 ± 2 °C.
9. If processing multiple samples, be sure to change gloves between each sample to ensure that there is no cross-contamination between samples during the weighing process. See flower example in Figure 3.



Figure 3: Homogenized cannabis flower and LB.

Sample Lysis

MaGiC Lysis for Flower, Gummy Infused Products, Chocolate and Oil Extracts

1. Remove the Whirl Pak Bag, 15, or 50 mL tube from the incubator.
 - a. Flower sample in Whirl Pak Bag: Mix thoroughly by hand manipulating and/or squeezing the contents for the flower bag for 1 minute.
 - b. Non flower Infused Product in 15 or 50 mL tube: vortex sample thoroughly for 30 seconds.

2. Dilute Internal Cannabis Control (IC) to 1:50,000 for all matrices.

Note: Prepared 1:50k dilution of IC has been tested up to 24 freeze thaws with minimal effect on resulting Cq value when stored at -20°C when not in use.

- a. Ensure stock IC is fully thawed, vortexed and spun down before use.
 - b. Dilute stock IC 1:100.
 - i. Add 2µL of IC to 198µL of Nuclease free Water. Close tube, vortex and quick spin.
 - c. Dilute the 1:100 dilution of IC to 1:10,000.
 - i. Add 2µL of 1:100 IC dilution to 198µL of water. Close tube, vortex and quick spin.
 - d. Dilute 1:10,000 dilution of IC to 1:50,000.
 - i. Add 20µL of 1:10,000 IC dilution to 80µL of water. Close tube, vortex and quick spin.
3. Remove 10 µL of enriched sample from the bag or tube and put into the well of a 96 well PCR plate.
4. Add 50 µL of MaGiC Lysis Buffer and pipette tip mix 15 times. Repeat for all wells being tested using a fresh tip for each transfer.
5. Transfer 5 µL of 1:50,000 diluted IC control to the 60 µL of sample. Repeat for all sample wells. Be sure to use a fresh tip every time.
6. Seal the top of the plate and spin down in a centrifuge.
7. Place plate on a thermal cycler and run MaGiC Lysis program
 - a. 95 °C for 10 minutes.
 - b. 4 °C for 10 minutes or “forever”. If the thermal cycler does not go down to 4°C set to 25°C.

Note: When using the Agilent AriaMX, a compression pad should be placed over the adhesive seal to prevent evaporation during thermal cycling.

8. Once the plate has been at 4 °C for 10 minutes or 25 °C for 10 minutes, remove it from the thermal cycler and spin the plate down to remove evaporation from the plate seal. Carefully remove the plate seal.
9. Add 100 µl of MaGiC Stabilization buffer to each sample well. Pipette mix 5 times.
10. Seal the plate with a plate seal and store in -20 °C freezer if not proceeding directly to qPCR setup.
 - a. Samples with stabilization buffer added are stable at -20 °C for 1 month.
 - b. Extracted samples which have been frozen must be tip mixed 15 times prior to setting up qPCR or performing dilutions.
11. Proceed to qPCR

Real-Time Quantitative PCR (qPCR) Setup Protocol with PathoSEEK Amplification Mix

1. Remove Amplification Mix, Listeria Detection Assay, and assay positive control from the -20 °C freezer.
 - a. If Lyophilized Amplification Mix has not been rehydrated, rehydrate a vial in 550 µl of Nuclease Free water. Swirl or Pipette mix. After resuspension, store at -15 to -20 °C when not in use.
 - b. Allow all frozen reagents to defrost at room temperature. Once defrosted, place tubes on ice.
2. Before preparing the master mix, invert or vortex and spin down the reagents.
 - a. Listeria Detection Assay v2 tube – Vortex quickly followed by a pulse spin down in a microcentrifuge.
 - b. Assay Positive Control tube – Vortex quickly followed by a pulse spin down in a microcentrifuge.
 - c. Amplification Mix – Invert the bottle 5-10 times to mix or briefly vortex.
 - d. Return all reagents to the ice.
3. Prepare Master Mix in a 1.5 mL tube (the probe mix also contains the internal cannabis control probe mix). Label tube “Master Mix”. Always prepare enough master mix for an additional 1 or 2 reactions to account for pipetting and dead volumes. Be sure to include 2 extra reactions for the

qPCR positive and negative controls. For example, if testing 10 samples, you would need to make enough master mix for 13 or 14 reactions, which would account for 1 or 2 excess. See Table 1 as a reference.

Table 1: PathoSEEK Amplification Master Mix Reagent Volumes

Reagent	Volume for 1 Reaction
Amplification Mix	10 µL
Listeria Detection Assay v2	1 µL
Nuclease Free Water	4 µL
Total	15 µL

- a. Once combined, cap tube and vortex to mix.
 - i. Pulse spin down tube in microcentrifuge.
 - ii. Place the Master Mix tube on ice until used.
4. Prepare a 1:10 dilution of the assay positive control
 - a. Add 1 µL of Positive Control to 9 µL nuclease free water (found in the kit), vortex to mix well and spin down the tube.

Note: It is best to add the largest volume reagent first, in this case the 9 µL water then the 1 µL of positive control, pipette mix or vortex control dilution to ensure control DNA is in solution.
5. For the negative control, use water that was used to rehydrate your Amplification Mix.
6. Use a 96-well optical qPCR plate or optically clear qPCR tubes.
7. Transfer samples into qPCR plate or tubes
 - a. Carefully remove the seal from the Extraction Plate.
 - i. If lysed samples were frozen, let the DNA thaw completely and spin the plate to avoid cross contamination between samples. Tip mix thawed samples well before transferring to the qPCR plate or tubes.
 - b. Transfer 5 µL of each sample into the corresponding qPCR tube or well on the qPCR plate.
 - c. Add 5 µL of the diluted Positive Control to the corresponding positive control well or tube.
 - d. Add 5 µL of water to the corresponding negative control well or tube.

Note: ALWAYS use a fresh tip for every liquid transfer into the qPCR plate

8. Add 15 µL of Master Mix to each corresponding sample well, positive control well, and negative control well in the qPCR plate or tubes. Gently tip mix a few times after each addition of qPCR master mix. Be careful not to introduce bubbles during this mix.
9. Seal the plate with strip caps or an adhesive seal or seal qPCR tubes with strip caps.
10. For the Agilent AriaMX or Biorad CFX, spin down qPCR plate or tubes for at least 1 minute in plate or tube microcentrifuge to bring well contents to the bottom of wells and help to get rid of reaction bubbles.

Note: Check for bubbles in the wells (minimal bubbles on the surface of the liquid is acceptable). If bubbles remain in the wells, *spin down* for another minute.

11. For the Agilent AriaMX: If using an adhesive seal; place the reusable compression pad (gray side down) on the plate directly lining up the holes in the pad with the wells in the plate.
12. Place the sealed plate or tubes onto the PCR instrument.
13. Follow the software specific instructions to initiate the run.

Running the BioMolecular Systems MIC

1. Open the BMS Workbench software and create a new file.
2. Select qPCR Run.
3. Select the appropriate template by clicking the “+” sign next to assays or ensure that the appropriate thermal cycling conditions are entered: Hot start at 95 °C for 5 minutes, followed by 40 cycles of 95 °C for 15 seconds and 65 °C for 90 seconds.
4. Click on the Mic icon in the upper right-hand corner, and select “Start run” from the menu that appears.



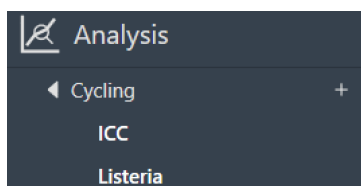
5. A pop-up will appear asking for the reaction volume, which will be auto-filled with the correct volume based on the template chosen. Close the lid and the instrument will start.

Note: If the cover isn't closed, the program will not start.

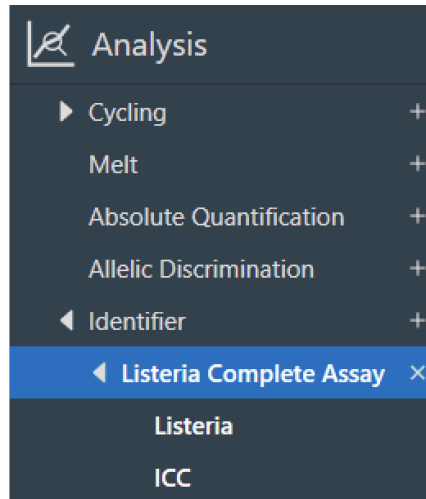
6. Let the Mic run to completion before analyzing the data.

Data Analysis with BioMolecular System MIC

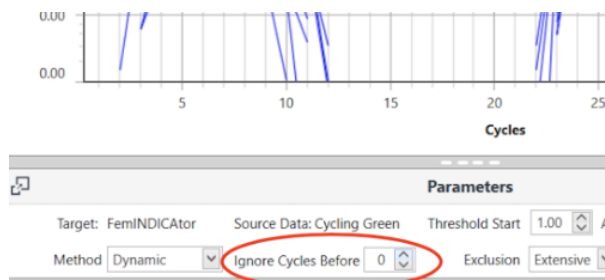
1. When the run has completed, click on the “+” sign next to the “Cycling” tab and select IC (Internal Cannabis Control). The selection will appear under the “Cycling” tab. Next, select *Listeria*.



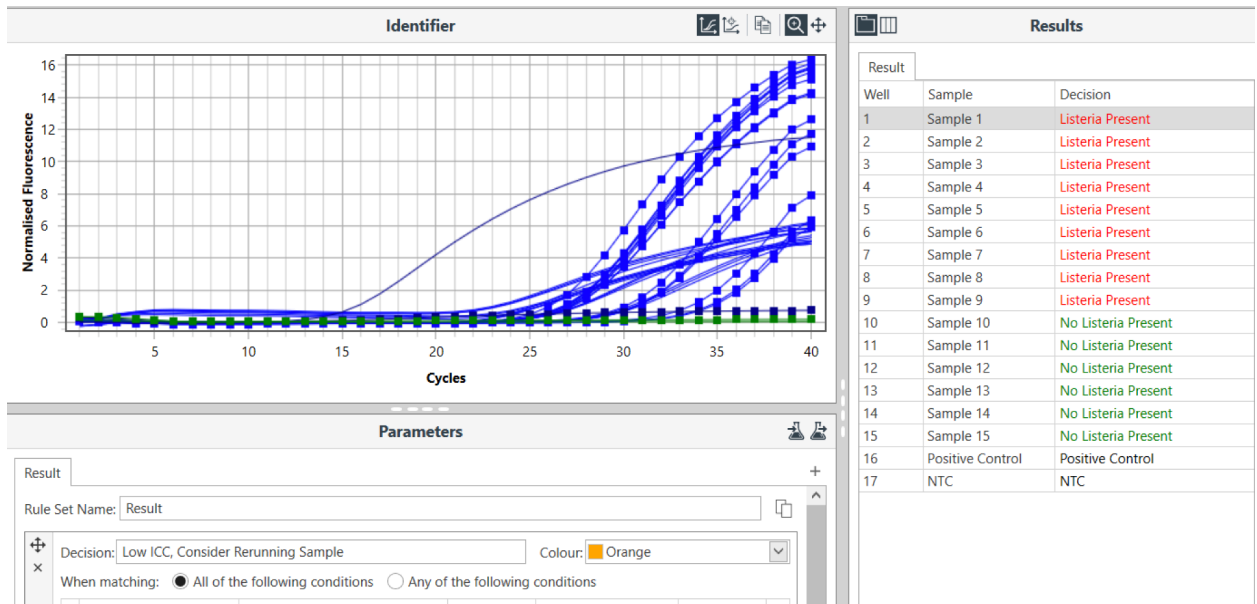
2. When these have been expanded, select the “+” sign next to the “Identifier” tab and select the available report.



3. If the automatic calls are blank, then one of the cycles is out of threshold. To fix this, find “Ignore Cycles Before” in the “Parameters” section, and increase its value one whole number at a time until a gray bar appears on the graph. This might have to be done with one, or all of the filters.



4. To review an automatic report, click the “+” sign next to the “Identifier” tab then select the relevant “Complete Assay”. This feature will call the samples tested as detect, non-detect, or inconclusive based on the qPCR data. These results will be displayed on the right side of the screen.



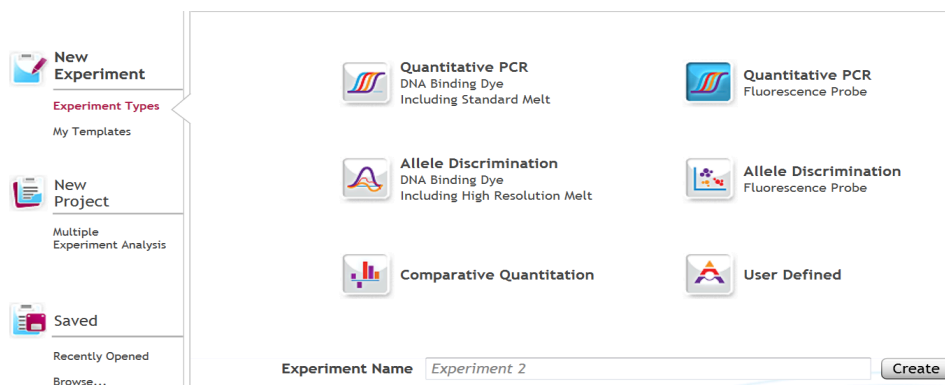
5. Data may be exported by selecting the “Report” tab, then clicking the Export icon.

Running the Agilent AriaMX

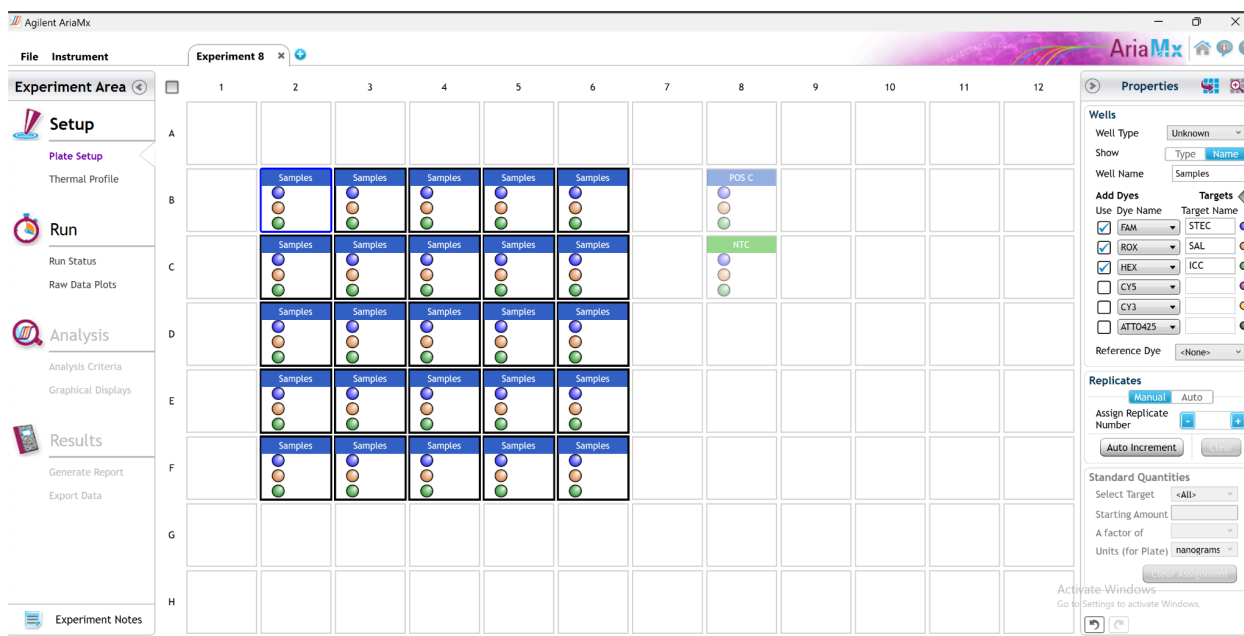
The following species will be detected on the following Fluorophores

Target	Optical Channel
<i>Listeria</i>	FAM
Internal Cannabis Control	HEX

1. Create a New Experiment on the Agilent qPCR instrument.
 - a. Select “Quantitative PCR: Fluorescence Probe” from Experiment Types.

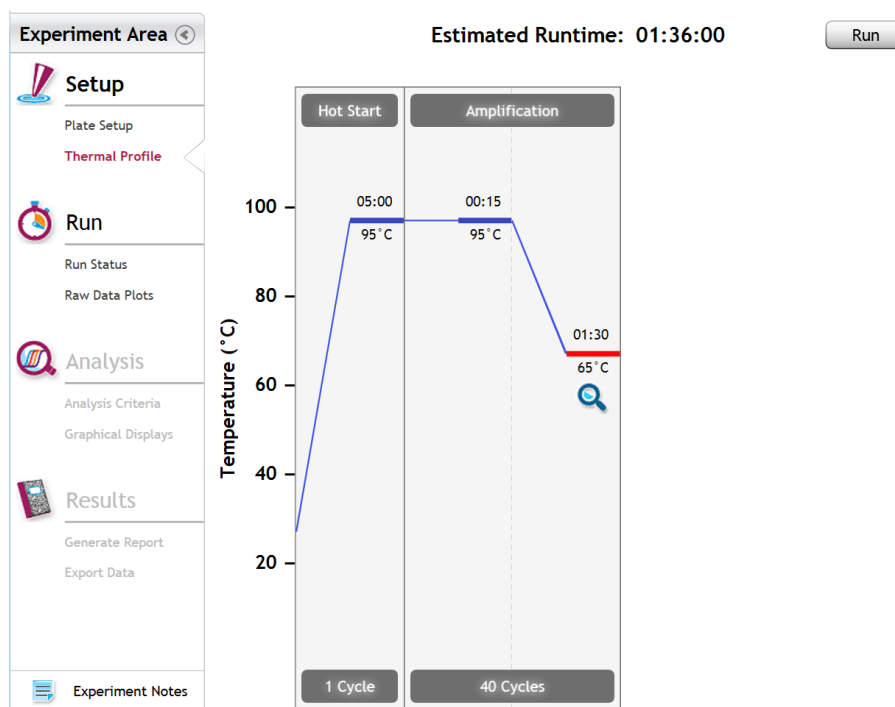


- b. Under Setup > Plate Setup, highlight **only** wells that contain reactions. Select FAM and HEX under **Add Dyes**. Having fluorescence detection turned on in empty wells can affect the calculations which the software makes to obtain the corrected data, ΔR log.



2. Change the well types to reflect your plate set up. All wells should be set to Unknown except the negative control may be set to NTC well type. Add Target names to the dyes under **Targets**:
 - a. FAM: *Listeria*.
 - b. HEX: Internal Cannabis Control
3. Under Setup > Thermal Profile, create the following PCR thermal profile
 - a. Hot Start at 95 °C for 5 minutes, followed by 40 cycles of 95 °C for 15 seconds and 65 °C for 90 seconds.

Note: A compression pad should be placed on top of the plate with an adhesive seal to prevent cross contamination.



4. Close the lid and click “Start Run”.
5. Save the experiment with the [User] and [date].
6. When the qPCR run is complete, immediately dispose of the plate or tubes. Do not remove the plate seal or tube caps to avoid contamination in the lab.

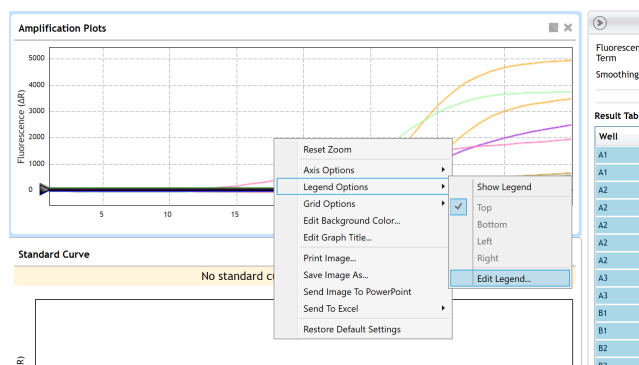
Data Analysis: Agilent AriaMX

PathoSEEK® Listeria Assay v2 Data Analysis Quick Reference Table

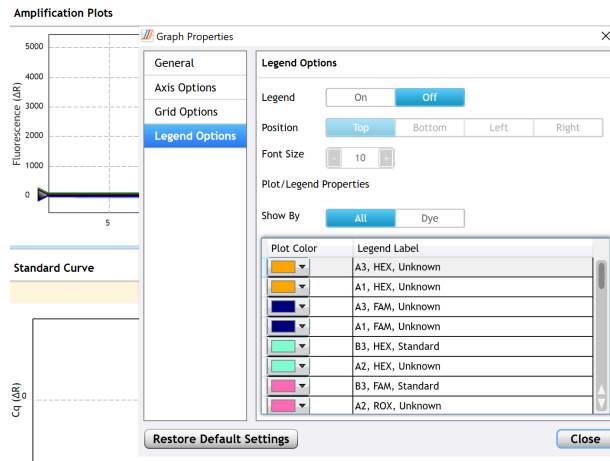
Table 2: All Matrices

PathoSEEK Assay	Cq Value	Fluor	Negative Control (Cq)	CFU threshold (CFU/g)
<i>Listeria</i>	≤ 40	FAM	No Cq	Presence/Absence
Internal Cannabis Control (IC)*	≤35	HEX	*Internal cannabis control verifies the presence or absence of cannabis DNA	
Assay Positive Control	≤35	FAM/ROX		

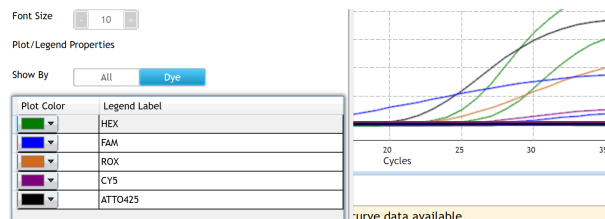
1. Open the Data Analysis window when the run is complete.
2. Highlight the wells of interest in the Analysis Criteria under Analysis, then select Graphical Displays
 - a. Amplification plots will be available for viewing
 - b. The Cq values will appear to the right in the table
 - c. Right click inside the graph, select Edit Legend under Legend Options



- d. Change “All” to “Dye”

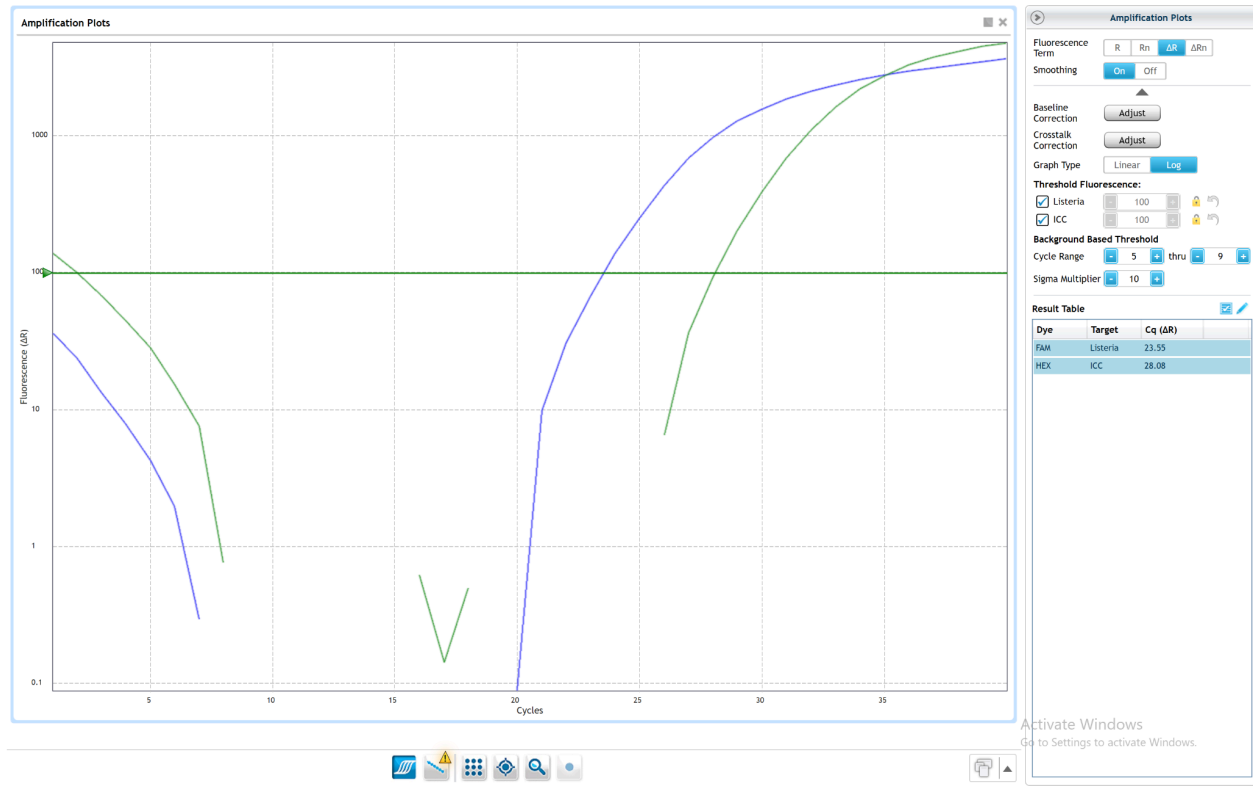


- e. All user settings for Plot/Legend Properties will be removed. Do you want to proceed?
Select “Yes”.
- f. This will assign a single color to each fluorophore.



3. To analyze the results

- a. Start by turning the graph to Log Scale with a right click on the chart, select Axis options, enable y-axis log scale. Expand the amplification plots settings by clicking on the triangle.



- b. Manually adjust thresholds to 100 RFU for the HEX and FAM fluorophores. (Target names will have previously been identified.)

Graph Type Linear Log

Threshold Fluorescence:

☒ Listeria - + 🔒 ↺

☒ ICC - + 🔒 ↺

Background Based Threshold

Cycle Range - + thru - +

Sigma Multiplier - +

4. Controls

- a. Positive Control, on the FAM fluorophores, have Cq values ≤ 35 .
 - i. Visually confirm with the curve on the graph.

- b. Negative Control, on the FAM Fluorophores, have no Cq Value.
 - i. Visually confirm with the curve on the graph.
- 5. Unknown *Listeria* Targets
 - a. A “presence” or failing result for the unknown *Listeria* target(s).
 - i. A Cq of ≤ 40 on the FAM channel indicates the presence of *Listeria*.
 - 1. Visually confirm with the curve on the graph. It is very important to check the raw data view (R) to confirm with the amplification curve when a presence result occurs.
 - a. Review the raw data for each sample: Select the wells of interest, use the plate view of the Raw Data Plots to scroll across the plate to confirm amplification in sample wells and positive control wells. This review also ensures that thermal cycling is uniform across the plate.

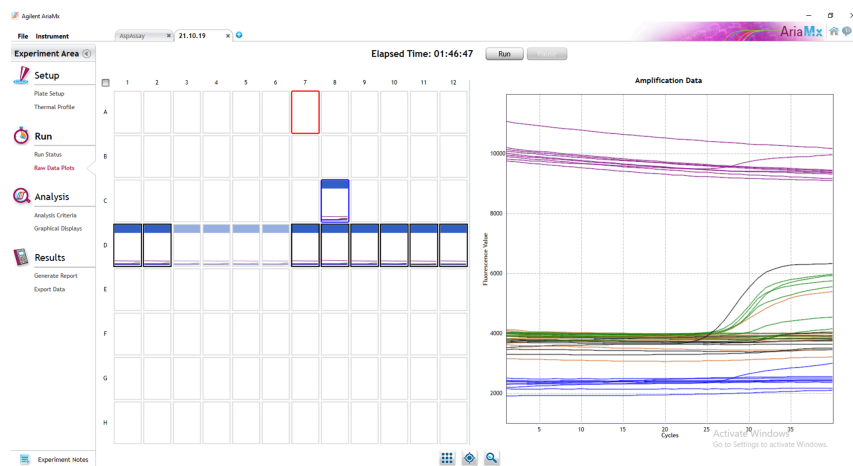


Figure 4: Raw data (R) plots for which the cycle quantification (Cq) will be generated.

- 2. Look at the raw data in the "Graphical Displays" view. For every dye that gives a Cq make sure the baseline is flat and the fluorescence signal grows rapidly over 5-7 cycles as seen below. (To view the raw data, select "R" next to "Fluorescence Term" at the top right-hand side of the screen.)

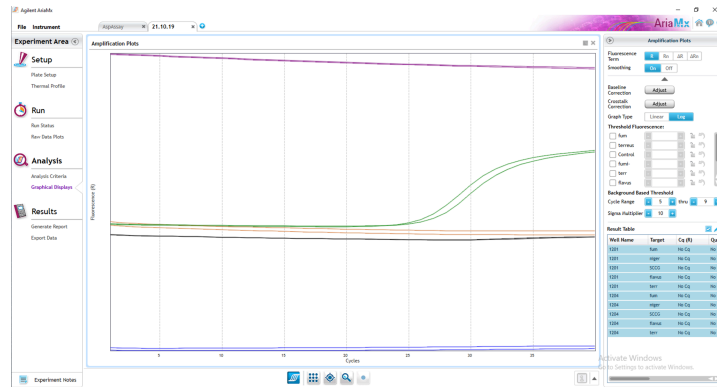
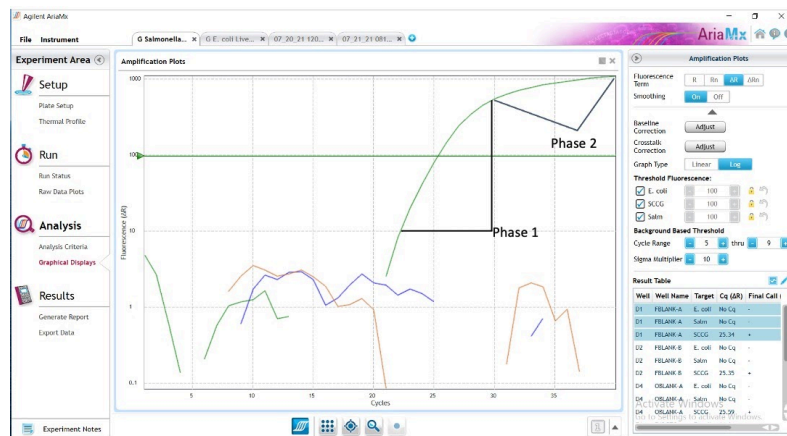


Figure 5: Example of good baseline with sigmoidal shaped curve increase in the internal cannabis control fluorescence.

3. Check the amplification plots for a semi-logarithmic curve with two distinct phases that crosses the threshold:
 - a. Select the ΔR Fluorescence Term and look at the shape of the amplification plot. The graph should have the classic semi-logarithmic shape with two visually distinct phases.
 - b. Phase 1 shows exponential growth in the fluorescence, in a span of around 5 cycles.
 - c. Phase 2 is a plateau where the amplification signal growth ends but remains level.



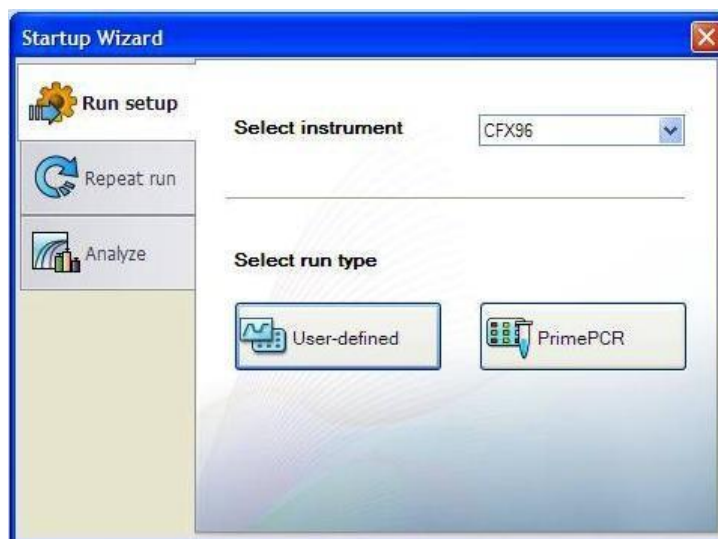
- b. An “absence” or passing result for the unknown *Listeria* target.
 - i. Internal Cannabis Control (IC), on the HEX Fluorophore, has a Cq value ≤ 35 for flower samples, ≤ 40 for all other matrices.
 - ii. No Cq value for the FAM Fluorophore.
 - iii. Visually confirm that there is no curve on the graph.

Running the BioRad CFX96

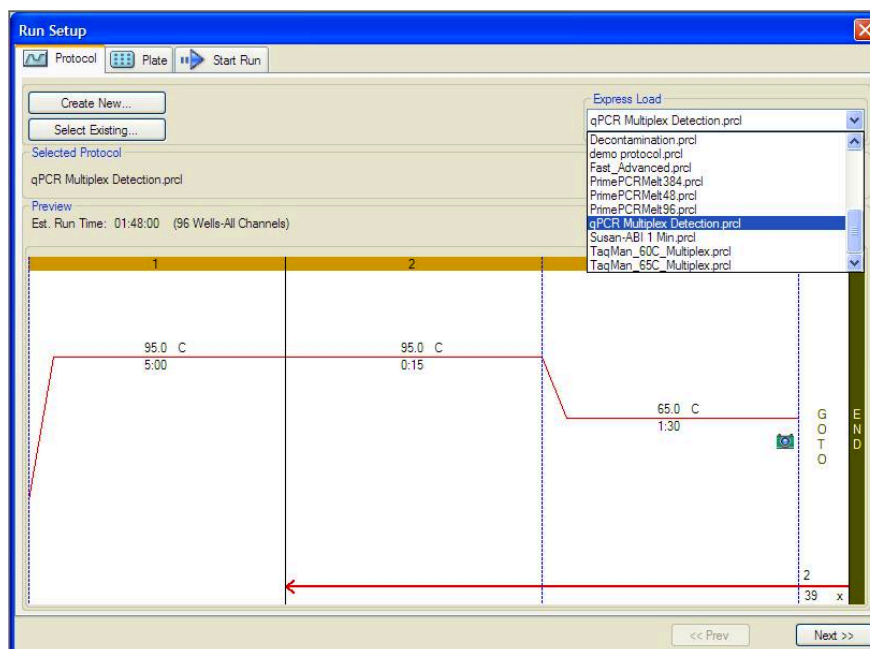
The following targets will be detected on the following fluorophores:

Target	Optical Channel
<i>Listeria</i>	FAM
Internal Cannabis Control	HEX

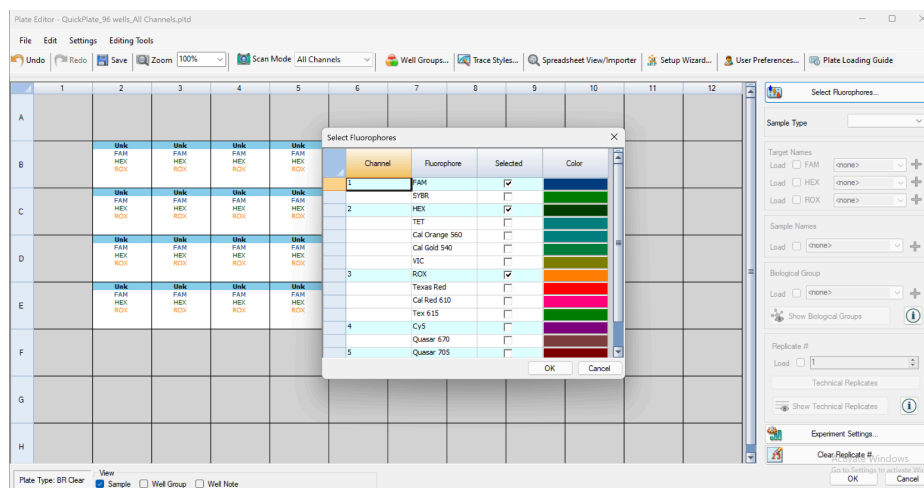
1. Start the qPCR Cycling program
 - a. Select User-Defined in the Startup Wizard under Run setup



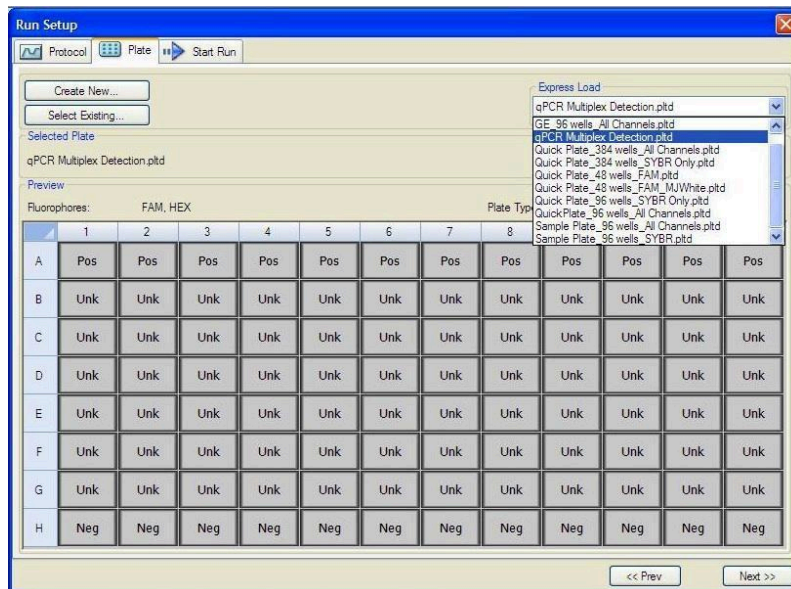
2. Use the Express Load dropdown menu to pick the qPCR Multiplex Detection Program and click “Next”.
 - a. If not already pre-programmed, create a cycling program with the following specifications and save as “qPCR Multiplex Detection”
 - b. Hot start at 95 °C for 5 minutes, followed by 40 cycles of 95 °C for 15 seconds and 65 °C for 90 seconds.



3. Design your plate under the plate tab in the Run Setup
 - a. Select the qPCR Multiplex Detection from the dropdown menu. If not already present, click “Create New”
 - b. Plate editor window will appear. Choose FAM and HEX Fluorophores and click “OK”.



- c. If the plate layout was previously saved, click “Edit Selected” to move to the Plate Editor Screen.



- d. On the Plate Editor Screen, change the Sample Type to correlate with your specific plate setup.

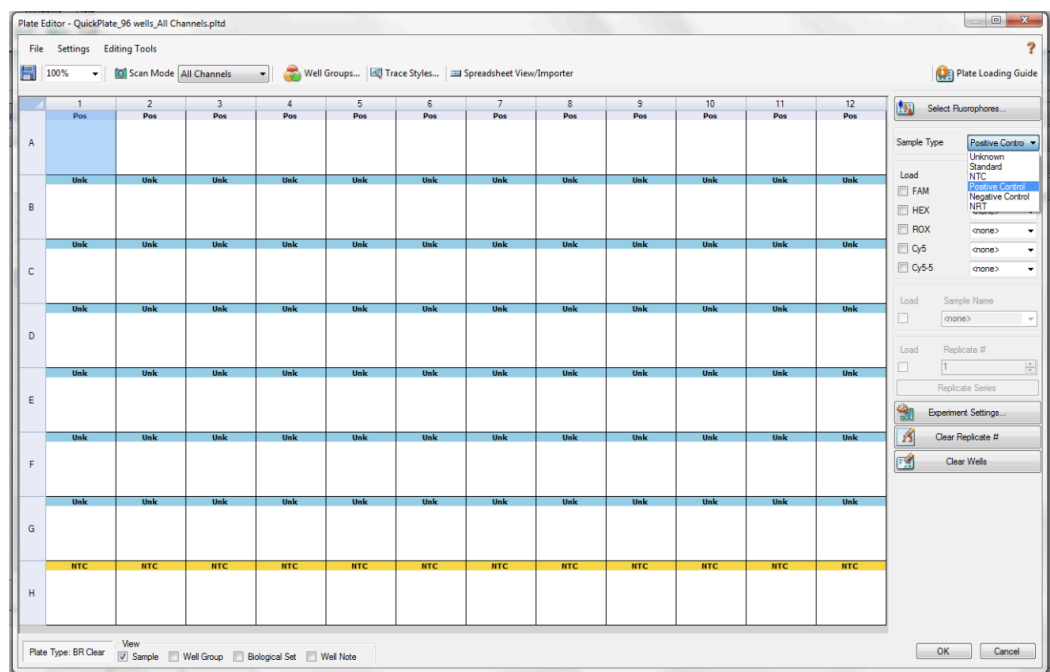
NOTE: To select the Sample Type, highlight the wells you would like to define, then choose from the dropdown menu one of three types:

Unknown

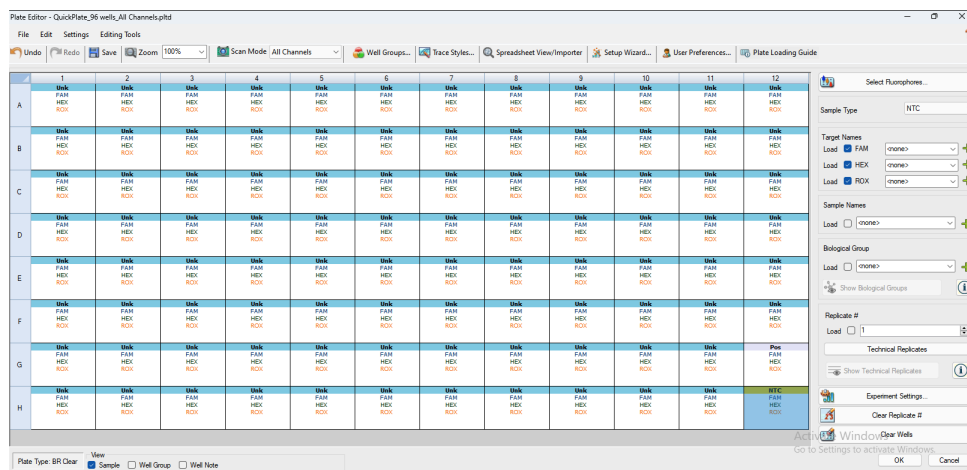
Positive Control

Negative Control

- e. Make sure All Channels is selected from the dropdown menu at the top.



- f. Attach the fluorophores to the wells being used.
 - i. Highlight all the wells being used
 - ii. For Listeria Assay, **Highlight the well locations and click on FAM and HEX.**
 - g. When the plate is designed correctly, click OK.
4. Click “Yes” to save your plate. If creating plate layout for the first time, save as “qPCR Multiplex Detection”. If you do not save the plate, it will return to the default plate.



Note: Saving will override the template (that is fine).

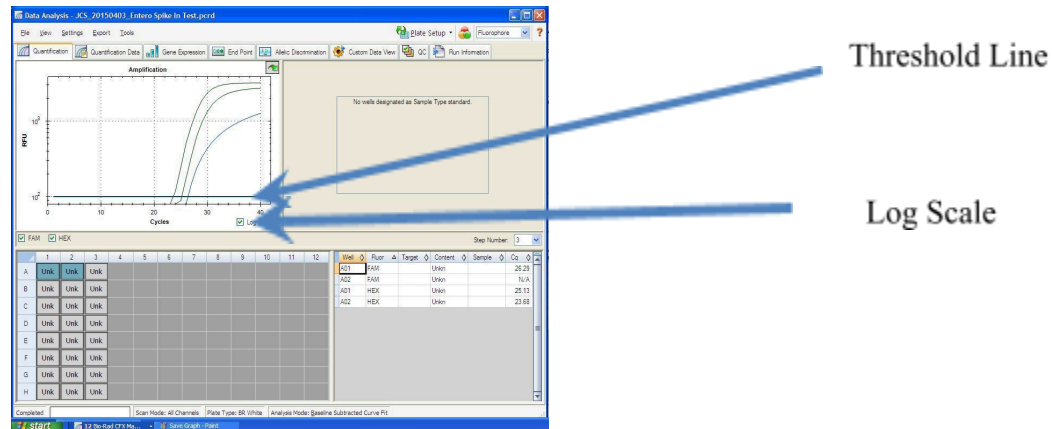
5. Close the lid and click Start Run.
6. Save the experiment with the [User] and [date].
7. When the run is complete, immediately dispose of the plate after qPCR. Do not remove the plate seal or tube caps after the run to avoid contamination in the lab.

PathoSEEK® Listeria Data Analysis Quick Reference Table

Table 3: All Matrices

PathoSEEK Assay	Cq Value	Fluor	Negative Control (Cq)	CFU threshold (CFU/g)
<i>Listeria</i>	≤ 40	FAM	No Cq	Presence/Absence
Internal Cannabis Control (IC)*	≤35	HEX	*Internal cannabis control verifies the presence or absence of cannabis DNA	
Assay Positive Control	≤35	FAM/ROX		

1. The Data Analysis window will open automatically when the run is complete.
2. Highlight the well of interest.
 - a. The graph will appear above.
 - b. The Cq values will appear to the right.
3. To analyze the results
 - a. Start by turning the graph to Log Scale and manually moving the threshold to 10² for all fluorophores.
 - i. To turn the graph to Log Scale, click on the box at the bottom right of the graph.
 - ii. To adjust the threshold, click on the horizontal lines, and move them to the specified value mentioned above on the y-axis.
 - iii. Alternatively, set an exact threshold value by selecting a single fluorophore beneath the graph, then Settings > Baseline Threshold. In the next window select User Defined and enter 100.



4. Controls

- a. Assay-specific Positive Controls on the FAM fluorophore have a Cq value ≤ 35 .
 - i. Visually confirm with the curve on the graph.
- b. Assay-specific Negative Controls on the FAM fluorophore have no Cq value.
 - i. Visually confirm with the curve on the graph.

5. Unknown *Listeria* Targets:

- a. A “presence” or failing result for the unknown *Listeria* targets.
 - i. Any Cq value for the FAM fluorophore ≤ 40 .
 - a. Visually confirm with the curve on the graph. It is very important to confirm with the amplification curve when a presence result occurred. Sometimes the background amplification will give a false positive reading.
 - ii. A Cq of ≤ 40 on the FAM channel indicates the presence of *Listeria*.
- b. An “absence” or passing result for the unknown *Listeria* target
 - i. Internal Cannabis Control (IC), on the HEX Fluorophore, has a Cq value ≤ 35 for flower samples, ≤ 40 for all other matrices.
 - ii. No Cq value for the FAM fluorophore.
 - iii. Visually confirm no curve on the graph.

Troubleshooting Guide

Symptom	Reason	Solution
Internal control failure	IC not spiked into sample/MaGiC lysis when applicable	Repeat MaGiC Lysis and PathoSEEK by following the protocol.
	Sample not added	Repeat MaGiC Lysis and PathoSEEK by following the protocol.
	Mix up in Reaction Setup	Repeat the qPCR by following the protocol.
	Missing Fluorophore on plate set up	Check plate setup in qPCR platform data file and ensure the correct fluorophores were chosen for the assay being run.
	Optional IC spike not performed for flower samples for Aspergillus Assays	The IC spike into flower is not mandatory when running the Aspergillus MaGiC lysis process. However if samples are low quality, it may be necessary
	qPCR inhibition	Dilute MaGiC Lysates 1:10 with nuclease free water and re-perform qPCR
Amplification of the Internal control is not expected in the assay positive or negative control wells. No Cq or a Cq of more than 35 is acceptable. Any Cq lower than 35 constitutes a rerun	Cannabis DNA or IC contamination in a reagent	Troubleshoot which reagent was contaminated; use new reagents, thoroughly clean all pipettes and bench areas with 10% bleach solution.
	qPCR bench too close to MaGiC lysis area	Designate separate benches, pipettes etc. for lysis and qPCR setup
Positive Negative Control	Small Cq value <15	Visually confirm that there is a true amplification curve. If not, this is a low level background and can be ignored
	Contamination	Repeat the qPCR by following the protocol.
	Insufficient pre-setup bleaching	Wipe down the lab workspace and all equipment with 10% Bleach. Repeat qPCR.

Glossary and Definitions

Deoxyribonucleic acid (DNA) is a [molecule](#) that encodes the [genetic](#) instructions used in the development and functioning of all known living [organisms](#).

Polymerase Chain Reaction (PCR) is a technology in molecular biology used to amplify a single copy or a few copies of a piece of DNA across several orders of magnitude, generating thousands to millions of copies of a particular DNA sequence.

A **fluorophore** is a fluorescent chemical compound that can re-emit light upon light excitation.

The **Negative Controls** are the reactions where no Cq is expected. It helps to ensure that all Assay-specific reactions are clean of contaminants.

The assay-specific **Positive Controls** are the reactions where a Cq is expected. It helps ensure that all Assay-specific reactions are working correctly. The Assay specific Positive Control is targeting the pathogen using the FAM Fluorophores.

Amplification of the **Internal Cannabis Control** or the microbial target of interest is expected in every reaction containing DNA isolated from a cannabis sample. It ensures the DNA isolation procedure was successful or the presence of microbial contamination. The internal cannabis control targets the cannabis genome, using the HEX Fluorophore.

Listeria Broth (LB) is a broth used to grow *Listeria* species more effectively.

References

1. FDA (2022) *Listeria* (Listeriosis) <https://www.fda.gov/food/foodborne-pathogens/listeria-listeriosis> (Accessed October 2024)
2. Ramaswamy, V., Cresence, V. M., Rejitha J. S., Lekshmi, M. U., Dharsana, K. S., Prasad, S. P., Vijila, H. M. (2007). *Listeria - review of epidemiology and pathogenesis*. Journal of Microbiology, Immunology and Infection. 2007 Feb;40(1):4-13.

Revision History

Version	Date	Description
v1	October 2024	PathoSEEK® Listeria Detection Assay v2 with MaGiC Lysis Method Launch
v2	April 2025	Amplification Mix packaging update Update to Listeria Enrichment Broth Packaging Info

DISCLAIMER

This test was developed, and its performance characteristics determined by Medicinal Genomics Company, for laboratory use. Any deviations from this protocol are not supported by MGC.

This test has not been validated on remediated (irradiated, ozone treated, acid treated, hydrogen peroxide treated, etc.) samples. Samples that have undergone remediation may cause discordant results between plating methods and PathoSEEK methods. When remediated samples produce a result above the action limit on qPCR, we recommend confirming viability with an approved plating method.

Results may vary based on laboratory conditions. Altitude and humidity are among factors known to affect the growth of bacterial and fungal species.

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