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Pathogen Detection and Quantification for the Beef Industry

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Current Industry Workflows - Beef



Salmonella Detection in Beef

For beef trim, 60 pounds was sourced from a local butcher for sampling. MicroTally[™] manual sampling devices (MSD) were removed from the sample bag, unfolded and firmly used to swab the beef trim. Once both sides of the cloth were used, swabs were folded back to the original dimensions with an additional horizontal fold per the manufacturer's instructions and placed into the original sample bag. The MSD swabs (n=15) were then spiked with various levels of *Salmonella* (1-10,000 CFU), homogenized in the proper enrichment medium and incubated. After 3-6 hours, samples were analyzed using the **BAX® System Real-Time PCR Assay for Salmonella**. All inoculated samples were detected after 4, 5, and 6 hours of enrichment and all 30 were confirmed with culture.

Salmonella Quantitation

When repeated for ground beef and beef trim (and MicroTally MSD swabs) using contamination levels of 10 CFU/g, the **BAX System Real-Time PCR Assay for Salmonella** detected the presence of the organism at 4-5 hours of enrichment. In addition, after a 6-hour enrichment, Salmonella could be quantified from 1-10,000 CFU/g when following the SalQuant[™] approach. This allows beef processors to not only identify the presence of Salmonella, but also to identify which lots of ground beef or beef trim contain higher levels of Salmonella, allowing for rapid action to reduce risk of exposure to consumers and to improve



BAX System Q7

food safety processes internally. (Similar results were also obtained for poultry.)

AOAC Validation Summary

AOAC-RI *PTM*SM Level 2 Modification for Hygiena[™] BAX System SalQuant Utilizing BAX System Real-Time PCR Assay for *Salmonella* for 7 Matrix Extensions

Process control and final product decisions based only on prevalence have shown limitations reducing consumer risk. Therefore, adoption of validated quantification methodologies with low error and wide enumerable ranges should be utilized to make data-driven food safety decisions.



This certification provides the poultry, beef, and pork industries with an accurate, reliable, and validated quantification tool to

reduce product hold-times, verify corrective actions, monitor process control, and promote faster datadriven diversion decisions which ultimately reduces consumer risk in animal protein products.

Validation Methods

- The evaluation consisted of 7 matrix studies to extend the methods' claims.
- Three distinct levels with a range of 1.0 Log CFU/mL(g) are established by the AOAC committee based upon enumeration capabilities of the candidate method.
- Each level has 5 unpaired individual samples that are tested for each level with the mean and error of each level utilized for comparison to the reference method.
- The candidate method must be within +/- 0.5 Log CFU/mL(g) of the MLG MPN 2.05 reference method for each level and be within the 90% confidence interval.

Matrix	Sample Size	Incubation Conditions	Enumerable Range
Poultry Rinsate	30 mL	42°C for 6 h	1 – 10,000 CFU/mL
Ground Beef	375 g	42°C for 6 h	1 – 10,000 CFU/g
Ground Pork	375 g	42°C for 6 h	1 – 10,000 CFU/g
Beef Trim	375 g	42°C for 6 h	1 – 10,000 CFU/g
Pork Trim	375 g	42°C for 6 h	1 – 10,000 CFU/g
MicroTally - Beef Trim	1 Swab	42°C for 6 h	1 – 10,000 CFU/mL
MicroTally - Pork Trim	1 Swab	42°C for 6 h	1 – 10,000 CFU/mL

Validation Results

- The Level 2 modification to the BAX System Real-Time PCR Assay for Salmonella, BAX System SalQuant (Certification No. 081201) was evaluated and approved by the AOAC Research Institute Performance Tested MethodsSM Program on January 12, 2022.
- Results of the validation study showed the SalQuant demonstrated comparable performance to that of the USDA-FSIS MPN reference methods for estimating *Salmonella* spp.

Application Highlights

- One enrichment, one sample prep, one assay; no additional equipment, consumables, or steps.
- Widest enumerable range across all matrices to facilitate contamination levels observed in sample types taken from farm to final product.
- Lowest level of enumeration (1 CFU/g(mL)) in order to truly quantify consumer risk.
- Largest data generation (>100,000 tests) to develop, verify, and validate on real industry samples across matrices, locations, facilities, instruments, and users for robustness and integrity of results.

AOAC-RI *PTM*SM Level 2 Modification for Hygiena BAX System SalQuant Utilizing BAX System Real-Time PCR Assay for *Salmonella* for Beef Matrix Extensions







Quantification Options



Limit of Quantification (LOQ)

If SalQuant samples are negative, but positive at prevalence, the result should be \leq enumerable range. (i.e., a poultry rinse was negative at the 6 h timepoint. The sample underwent continued incubation and was tested for prevalence. The prevalence test was positive; therefore with a negative quantification test, but a positive prevalence test, the result for quantification would be < 1 CFU/mL).

Application	Industry	Segment	Matrix	Tiimepoint	Limit of Quantification
SalQuant	Beef	Primary Production	Feces	8 h	10 CFU/g
SalQuant	Beef	Primary Production	Feces (high*)	0 h	100,000 CFU/g
SalQuant	Beef	Processing	Lymph Nodes (small)	6 h	10 CFU/Lymph Nodes
SalQuant	Beef	Processing	Lymph Nodes (medium)	6 h	10 CFU/Lymph Nodes
SalQuant	Beef	Processing	Cecal Swabs	8 h	10 CFU/mL
SalQuant	Beef	Processing	Cecal Contents	8 h	1 CFU/g
SalQuant	Beef	Processing	Cecal Contents (high*)	0 h	100,000 CFU/g
SalQuant	Beef	Final Product	Ground Beef	6 h	1 CFU/g
SalQuant	Beef	Final Product	Trim	6 h	1 CFU/g
SalQuant	Beef	Final Product	MicroTally	6 h	1 CFU/mL
SalQuant	Environmental	Environmental	Swabs	6 h	1 CFU/mL

* high = high concentration of Salmonella

Limit of Detection (LOD)

LOD is utilized as a limits approach or threshold testing. No calculations are utilized to determine LOD, only the timepoint and detection of bacteria indicate the limit of detection has been met. (i. e., ground beef LOD at 5 h is 10 CFU, therefore if the sample is positive at 5 hours, the results would be \geq 10 CFU/g)

Application	Industry	Segment	Matrix	Tiimepoint	Limit of Detection
SalQuant	Beef	Final Product	Ground Beef	5 h	≥ 10 CFU/g
SalQuant	Beef	Final Product	Trim	4 h	≥ 10 CFU/g
SalQuant	Beef	Final Product	MicroTally	4 h	≥ 10 CFU/mL

Beef Primary Production

Feces Enrichment & PCR Procedure

Add 10 g of beef feces to 90 mL of pre-warmed 42°C BAX MP + 0.5 mL/L Quant Solution as primary enrichment. Homogenize by hand for 60 seconds.

Transfer 10 mL of primary enrichment into a sterile container with 30 mL of pre-warmed (42°C) BAX MP media containing 0.5 mL/L of BAX Quant Solution. Hand massage for 15-30 seconds for homogenization.

Samples with enumerable ranges from 10 - 10,000 CFU/g require incubation at 42°C for 8 h. Samples with greater than 100,000 CFU/g do not require incubation, proceed directly to PCR.

Following enrichment for quantification, perform PCR analysis with the BAX System Real-Time PCR Assay for *Salmonella*. Collect results from the BAX System Q7 and utilize the Excel[®] Calculator or Online Software to quantify results.

After transferring aliquot for quantification enrichment, incubate the remaining sample in primary enrichment at $37 \pm 1^{\circ}$ C for 18-24 h for prevalence testing using the BAX System Real-Time PCR Assay for *Salmonella*.



Lymph Nodes Enrichment & PCR Procedure

Weigh and process lymph nodes into small (<10 g) or medium (>10 g) size category.

For small nodes, add 40 mL of pre-warmed (42°C) BAX MP media and for medium nodes, 80 mL of pre-warmed (42°C) BAX MP media as primary enrichment. Homogenize at 230 RPM for 30 seconds.

Incubate sample at $42 \pm 1^{\circ}$ C for 6 h.

Following enrichment for quantification or LOD, perform PCR analysis with the BAX System Real-Time PCR Assay for *Salmonella*. Collect results from the BAX System Q7 and utilize the Excel Calculator or Online Software to quantify results.

After quantification enrichment, incubate the remaining primary enrichment at $37 \pm 1^{\circ}$ C for 18-24 h for prevalence testing using the BAX System Real-Time PCR Assay for *Salmonella*.



Cecal Swabs Enrichment & PCR Procedure

Swab beef ceca with a 25 mL pre-moistened BPW swab and combine with 50 mL of prewarmed (42°C) BAX MP media containing 1 mL/L of BAX Quant Solution as the primary enrichment. Homogenize at 230 RPM for 30 seconds.

Incubate sample at $42 \pm 1^{\circ}$ C for 8 h.

Following enrichment for quantification, perform PCR analysis with the BAX System Real-Time PCR Assay for *Salmonella*. Collect results from the BAX System Q7 and utilize the Excel Calculator or Online Software to quantify results.

After quantification enrichment, incubate the remaining primary enrichment at $42 \pm 1^{\circ}$ C for 18-24 h for prevalence testing using the BAX System Real-Time PCR Assay for *Salmonella*.



Cecal Contents Enrichment & PCR Procedure

Add 10 g of beef cecal contents to 90 mL of BAX MP with + 0.5 mL/L Quant Solution as the primary enrichment. Homogenize at 230 RPM for 30 seconds.

Transfer 10 mL of the primary enrichment into a sterile container with 10 mL of pre-warmed (42°C) BAX MP media containing 0.5 mL/L of BAX Quant Solution. Hand massage for 15-30 seconds for homogenization.

Samples with enumerable ranges from 10 - 10,000 CFU/g require incubation at 42°C for 8 h. Samples with greater than 100,000 CFU/g do not require incubation, proceed directly to PCR.

Following enrichment for quantification, perform PCR analysis with the BAX System Real-Time PCR Assay for *Salmonella*. Collect results from the BAX System Q7 and utilize the Excel Calculator or Online Software to quantify results.

After transferring aliquot for quantification enrichment, incubate the remaining primary enrichment at $37 \pm 1^{\circ}$ C for 18-24 h for prevalence testing using the BAX System Real-Time PCR Assay for *Salmonella*.



Trim Enrichment & PCR Procedure

Add 375 g of beef trim to 1,500 mL of pre-warmed (42°C) BAX MP media as primary enrichment. Homogenize at 230 RPM for 30 seconds

Incubate sample at $42 \pm 1^{\circ}$ C for 4 h for LOD10 or 6 h for LOD1 and 1 - 10,000 CFU/g enumerable range.

Following enrichment for quantification or LOD, perform PCR analysis with the BAX System Real-Time PCR Assay for *Salmonella*. Collect results from the BAX System Q7 and utilize the Excel Calculator or Online Software to quantify results.

After quantification enrichment, incubate the remaining primary enrichment at $37 \pm 1^{\circ}$ C for 18-24 h for prevalence testing using the BAX System Real-Time PCR Assay for *Salmonella*.



MicroTally Enrichment & PCR Procedure

Add 1 MicroTally to 200 mL of pre-warmed (42°C) BAX MP media as primary enrichment. Homogenize at 230 RPM for 30 seconds

Incubate sample at $42 \pm 1^{\circ}$ C for 4 h for LOD10 or 6 h for LOD1 and 1 - 10,000 CFU/g enumerable range.

Following enrichment for quantification or LOD, perform PCR analysis with the BAX System Real-Time PCR Assay for *Salmonella*. Collect results from the BAX System Q7 and utilize the Excel Calculator or Online Software to quantify results.

After quantification enrichment, incubate the remaining primary enrichment at $37 \pm 1^{\circ}$ C for 18-24 h for prevalence testing using the BAX System Real-Time PCR Assay for *Salmonella*.



Beef Final Product

Ground Beef Enrichment & PCR Procedure

Add 375 g of ground beef to 1,500 mL of pre-warmed (42°C) BAX MP media as primary enrichment. Homogenize at 230 RPM for 30 seconds.

Transfer 30 mL of primary enrichment into a sterile container with 30 mL of pre-warmed (42°C) BAX MP media containing 1 mL/L of Quant Solution. Hand massage for 15-30 seconds for homogenization.

Incubate sample at $42 \pm 1^{\circ}$ C for 5 h for LOD10 or 6 h for LOD1 and 1 - 10,000 CFU/g enumerable range.

Following enrichment for quantification or LOD, perform PCR analysis with the BAX System Real-Time PCR Assay for *Salmonella*. Collect results from the BAX System Q7 and utilize the Excel Calculator or Online Software to quantify results.

After transferring aliquot for quantification enrichment, incubate the remaining primary enrichment at $37 \pm 1^{\circ}$ C for 18-24 h for prevalence testing using the BAX System Real-Time PCR Assay for *Salmonella*.



Environmental Monitoring

Swab - D/E Broth Enrichment & PCR Procedure

Add 1 environmental swab to 50 mL of pre-warmed (42°C) BPW media as the primary enrichment. Homogenize at 230 RPM for 30 seconds.

Incubate sample at $42 \pm 1^{\circ}$ C for 6 h.

Following enrichment for quantification, perform PCR analysis with the BAX System Real-Time PCR Assay for *Salmonella*. Collect results from the BAX System Q7 and utilize the Excel Calculator or Online Software to quantify results.

After quantification enrichment, incubate the remaining primary enrichment at $42 \pm 1^{\circ}$ C for 18-24 h for prevalence testing using the BAX System Real-Time PCR Assay for *Salmonella*.



Quantification Workflow Comparison Beef Trim

BAX [®] System SalQuant [™]
375 g beef trim
1,500 mL BAX MP
Homogenize/stomach
Incubation at 42° C for 5 – 6 h
Transfer 5 µL into lysis solution
Heat at 37°C for 10 min
Heat at 95°C for 20 min
Cool in cold block for 5 min
Hydrate BAX System Real-Time PCR Assay <i>Salmonella</i> with 30 µL of lysate
Initialize and run the BAX System (75 min)
Utilize the BAX Cycle Threshold (CT) in Excel spreadsheet or BAXQuant Online Software
True quantification results available



GENE-UP® Quant Salmonella

100 g beef trim
5X PBS or other media
Homogenize/stomach
No incubation
Transfer 40 mL of sample into 50 mL tube
Centrifuge for 10 min into a 500 g pellet of debris
Transfer 25 mL of supernatant to clean tube
Centrifuge for 10 min at 4300 g, to concentrate Salmonella
Decant supernatant
Resuspend pellet with 600 µL of Promega Nuclei Lysis Solution
Vortex for 10 seconds
Transfer 600 μ L of sample into 1.5 mL tube
Incubate at 80°C for 5 min
Cool on ice for 20 min
Resuspend pellet with 600 µL of Promega Nuclei Lysis Solution
Add 200 µL of Promega Protein Precipitation Buffer
Vortex sample solution
Incubate on ice for 5 min
Centrifuge 3 min at 16,000 g
Prepare 600 μL of 95% ethanol in 1.5 mL tubes and preheat to 80°C
Transfer 600 μL of supernatant into prepared solution above (be careful to avoid precipitate)
Invert to mix solutions
Centrifuge 3 min at 16,000 g
Pipette 50 μL of solution plus 50 μL of DNA Resuspension Buffer into 1.5 mL tubes and warm in heat block
Remove PCR tubes from freezer to thaw in centrifuge rack
Decant alcohol from tubes and tap on absorbent paper
Resuspend pellet in 80 μL of prewarmed DNA suspension buffer from above step
Vortex 5 seconds
Transfer 5 μL from sample to thawed VeriPro Salmonella qPCR tube
Briefly centrifuge to settle
Load plate and initialize GENE-UP System

True quantification results available

Quantification Workflow Comparison MicroTally for Beef Trim

BAX[®] System SalQuant[™]

1 Swab	
200 mL BAX MP	
Homogenize/stomach	
Incubation at 42°C for 4 - 6 h	
Transfer 5 μ L into lysis solution	
Heat at 37°C for 10 min	
Heat at 95°C for 20 min	
Cool in cold block for 5 min	
Hydrate BAX System Real-Time PCR Assay <i>Salmonella</i> with 30 μL of lysate	
Initialize and run the BAX System (75 min)	
Utilize the BAX Cycle Threshold (CT) in Excel	

spreadsheet or BAXQuant Online Software

True quantification results available



GENE-UP® Quant Salmonella

1 Swab
5X PBS or other media
Homogenize/stomach
No incubation
Transfer 40 mL of sample into 50 mL tube
Centrifuge for 10 min into a 500 g pellet of debris
Transfer 25 mL of supernatant to clean tube
trifuge for 10 min at 4300 g to concentrate Salmonella
Decant supernatant
Resuspend pellet with 600 µL of Promega Nuclei Lysis Solution
Vortex for 10 seconds
Transfer 600 μ L of sample into 1.5 mL tube
Incubate at 80°C for 5 min
Cool on ice for 20 min
Resuspend pellet with 600 µL of Promega Nuclei Lysis Solution
Add 200 µL of Promega Protein Precipitation Buffer
Vortex sample solution
Incubate on ice for 5 min
Centrifuge 3 min at 16,000 g
Prepare 600 μL of 95% ethanol in 1.5 mL tubes and preheat to 80°C
fer 600 μL of supernatant into prepared solution above (be careful to avoid precipitate)
Invert to mix solutions
Centrifuge 3 min at 16,000 g
tte 50 μ L of solution plus 50 μ L of DNA Resuspension

Remove PCR tubes from freezer to thaw in centrifuge rack

Decant alcohol from tubes and tap on absorbent paper

Resuspend pellet in 80 µL of prewarmed DNA suspension buffer from above step

Vortex 5 seconds

Transfer 5 µL from sample to thawed VeriPro Salmonella qPCR tube

Briefly centrifuge to settle

Load plate and initialize GENE-UP System

True quantification results available