

**Preparation Guidance for the detection of *Salmonella* in Papaya**  
**Updated: 8/18/2017**

\*\*Please note, this guidance is intended to provide supplemental general information to private laboratories on how to perform the initial sample preparation for papayas. This guidance document **does not** outline all of the analytical method or worksheet requirements for packages being submitted for FDA review. \*\*

Please refer to the current FDA Private Laboratory Guidance for comprehensive information on private laboratory package requirements and the review process:  
<https://www.fda.gov/downloads/ScienceResearch/FieldScience/LaboratoryManual/UCM092191.pdf>

**Preparation Guidance:**

- Samples should consist of 10 sub samples for official analysis. An additional sub sample of papaya will be required if a matrix spike is being performed (refer to Matrix Validation/Spike section below).

**Analytical Protocol:**

- Samples consist of 10 sub samples. Analyze subs individually.

**Preparation Modified Buffered Peptone Water**

Peptone	10.0 g
Sodium Chloride	5.0 g
Disodium Phosphate (Na <sub>2</sub> HPO <sub>4</sub> )	7.0 g
Monopotassium Phosphate (KH <sub>2</sub> PO <sub>4</sub> )	3.0 g
Distilled water	1000 mL

pH to 7.2±0.2

Note: commercial Buffered Peptone Water (BPW; BD catalog number 218105) may be substituted for the above formulation. If a commercial preparation is used, then add 3.5 g of Na<sub>2</sub>HPO<sub>4</sub> and 1.5 g of KH<sub>2</sub>PO<sub>4</sub> to 20 g commercial BPW in 1000 ml distilled water.

Sterilize by autoclaving 15 min at 121°C.

### **Sub-sample soak preparation:**

For each individual sub sample (e.g. one papaya ), place contents into a sterile plastic bag (Biopro Sample Bag, 12 x 18 inches, catalog number BP-41218, available from International Bioproducts, phone 800-729-7611 or equivalent). Add a volume of modified buffered peptone water (mBPW) that is needed to allow the papayas to float. Normally this volume of mBPW is 1.5 times the weight of the papaya. Allow the open-end flap of the plastic bag to “fold over” so as to form a secure, but not airtight, closure during incubation.

### **Sample preparation/method:**

- Incubate each, individual papaya sub sample at 35<sup>0</sup>± 2<sup>0</sup> C for 22 ± 2 hours. Sub samples should be analyzed individually (**not** wet composited)
  - It is acceptable for laboratories to use any AOAC Official Method for *Salmonella*.

**For rapid method kits, selectively enrich as instructed by the kit manufacturer.**

### **Confirmation:**

- Samples found positive are confirmed as directed in the BAM Online.  
<https://www.fda.gov/Food/FoodScienceResearch/LaboratoryMethods/ucm070149.htm>

### **Matrix Validation/Spiking:**

- Laboratories must demonstrate successful detection of *Salmonella* for the methodology being utilized by analyzing a spiked matrix concurrently with the sample through confirmation.
- Matrix spike should consist of an inoculum of 30 cells or less of *Salmonella* added to a papaya control sample. A negative matrix spike will invalidate the analysis.
- Matrix spike details (CFU or colonies/gram) must be included with the analytical package.
- Once a laboratory can demonstrate that their papaya spikes have yielded at least

seven positive and no negative matrix spikes or a >95% confidence level (19 of 20 samples positive), the matrix can be considered validated for the method being used. The laboratory can discontinue performing matrix spikes on subsequent papaya samples analyzed with that method, **but must submit documentation of the matrix validation data with each subsequent papaya sample analytical package submitted for FDA review.**

**Quality Assurance:**

- The Quality Assurance information for the media and reagents used in the analysis must be submitted with the analytical package (e.g. pH check, autoclave run time/temp documentation, performance and sterility of media, etc.).
- Laboratory must follow the methodology specified in the private laboratory package submission. Any method modifications or deviations to the cited method must be explained and validation must be documented.