IT 1-2-3™ SWIPE
Sample Purification Kit

Instruction Booklet

Manufactured by
BioFire Defense, LLC
79 West 4500 South, Suite 14
Salt Lake City, Utah 84107 USA
<table>
<thead>
<tr>
<th>CUSTOMER AND TECHNICAL SUPPORT</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Reach Us on the Web</strong></td>
</tr>
<tr>
<td><strong>Reach Us by E-mail</strong></td>
</tr>
<tr>
<td><a href="mailto:support@biofiredefense.com">support@biofiredefense.com</a></td>
</tr>
<tr>
<td><strong>Reach Us by Fax</strong></td>
</tr>
</tbody>
</table>
# Table of Contents

Abbreviations and Acronyms .......................................................... 5

General Safety Precautions ................................................................. 7

Laboratory Procedures and Precautions .............................................. 7

Introduction ....................................................................................... 8

IT 1-2-3 Sample Purification Kits ....................................................... 8

Purifying DNA or RNA ................................................................. 9

Inhibition Controls ................................................................. 9

Sample Purification Procedures .......................................................... 10

IT 1-2-3 SWIPE Sample Purification Kit for Nasal/Pus Swab, Surface Swab, Lymph Node Aspirates, Powder and Culture Samples .......... 10

Optimal Protocol Definition ....................................................... 10

Alternative Protocol Definition ...................................................... 10

Nasal/Pus Swab Protocol (DNA) .................................................. 12

Lymph Node Aspirate Protocol (DNA) ......................................... 14

Surface Swab Protocol (DNA) ..................................................... 16

Surface Swab Protocol (RNA) ....................................................... 18

Powder Protocol (DNA) ............................................................ 20

Powder Protocol (RNA) ............................................................ 22

Culture Protocol (Including Blood Culture; DNA) ....................... 24

Culture Protocol (RNA) ............................................................. 26

Water/PBS Protocol (DNA) ......................................................... 28

Water/PBS Protocol (RNA) ......................................................... 30

Troubleshooting ........................................................................... 32

Ordering Information .................................................................. 33

Sample Purification Kits and Supplies ............................................. 33
Abbreviations and Acronyms

BFDf...............................BioFire Defense
BMBL...............................Biosafety in Microbiological and
Biomedical laboratories
BW...............................Biological Warfare
C...............................Celsius
DFU...............................Dry Filter Unit
DHHS...............................Department of Health and Human Services
DNA...............................Deoxyribonucleic acid
\( g \)...............................Gravity (= RCF)
LOD...............................Limit of Detection
mL...............................Milliliter
MSDS...............................Material Safety Data Sheet
PBS...............................Phosphate-buffered saline
RCF...............................Relative Centrifugal Force ( = g)
RNA...............................Ribonucleic acid
\( \mu L \)...............................Microliter (0.000001 Liters)
General Safety Precautions

Laboratory Procedures and Precautions

• To avoid contamination, we recommend using filter tips on pipettes when working with any liquid solution.
• A Biosafety cabinet should be used when a potentially infectious material is used in procedures where there is potential for creating aerosols or splashes.
• When working with potentially harmful samples or chemicals always wear the appropriate personal protective equipment (lab coat, gloves, and eye protection).
• Avoid exposure to any potentially infectious samples or harmful chemicals. Exposure can occur by inhalation, ingestion or skin absorption.
• For more information on kit components consult the appropriate SDS provided by BioFire Defense.

Precautions


Handling of Biohazard Wastes

Use universal precautions when handling human blood and body fluids. Dispose of used reagent vials in accordance with good laboratory practices. Before disposal, waste from possible biohazardous samples should be inactivated using appropriate procedures.

For more information refer to DHHS (NIOSH) Publication No. 88-119 Guidelines for Protecting the Safety and Health of Health Care Workers (section 6, Hazardous Waste Disposal).
Introduction

IT 1-2-3 Sample Purification Kits

For a complete overview of matrices and kits, see the table at the end of this booklet.

BioFire Defense offers additional kits for purification of DNA and RNA:

- **IT 1-2-3 Platinum Path Sample Purification Kit**
  for magnetic bead purification of biological, environmental, and food samples
- **IT 1-2-3 FLOW Sample Purification Kits**
  for purification of blood, air, water, and food samples
- **IT 1-2-3 SWIPE Sample Purification Kit**
  for purification of nasal/pus and surface swabs, lymph node aspirates, live culture and powder samples
- **IT 1-2-3 SCOOP Sample Purification Kit**
  for purification of stool and soil samples
- **IT 1-2-3 VIBE Sample Purification Kit**
  for purification of sputum samples and RNA from blood, nasal, and throat swab samples
- **IT 1-2-3 DNA Sample Purification Kit**
  for purification of surface swab, powder, culture, and water/PBS samples

All kits contain protocols that are performed manually and have been simplified and clearly defined to reduce the risk of operator error. They have been designed to purify samples for analysis with BioFire Defense’s R.A.P.I.D.® and JBAIDS detection systems.

The IT 1-2-3 SWIPE Kit is an upgraded version of the IT 1-2-3 RAPID DNA Purification Kit.
**Purifying DNA or RNA**

DNA and RNA are present in bacterial or viral pathogens that can be found in biological, environmental or food samples. Such samples include nasal and environmental swabs, water, blood, etc. There are many inhibitors present in some samples that need to be removed with purification protocols for downstream sample analysis. Inhibitors include cellular debris, chemicals, enzymes and other naturally occurring inhibitors that either degrade DNA or RNA or hinder downstream analysis. DNA and RNA extraction and purification from a complex sample is usually necessary before identification and/or quantification steps can be successfully performed. The protocols in this kit employ the steps described below.

Sample Purification involves four steps:

1. DNA/RNA is **Extracted** from the sample (e.g. cells or spores) through lysis. This is achieved by physical agitation and chemical disruption of the cells with bead-beating or heat (heat is adequate for RNA viruses).
2. DNA/RNA is **Bound** and concentrated on the filter (i.e. **Buffer 1**).
3. The DNA/RNA on the filter is **Washed** to remove inhibitors (i.e. **Buffer 2**).
4. The DNA/RNA is **Eluted** from the filter (i.e. **Buffer 3**).

Other steps are often added to increase total DNA/RNA recovered from various sample matrices. For example, a protease step is added to the blood protocol to break down unwanted proteins.

**Inhibition Controls**

Inhibition controls included in a test ensure that a given sample does not inhibit the downstream analysis and that a negative result is real. The troubleshooting section describes what to do if an inhibition control shows that a sample is inhibited. Contact BFDf to see if inhibition controls are available for your assay.
Sample Purification Procedures

IT 1-2-3 SWIPE Sample Purification Kit for Nasal/Pus Swab, Surface Swab, Lymph Node Aspirates, Powder and Culture Samples

This kit and protocols are designed to purify deoxyribonucleic acids (DNA) or ribonucleic acids (RNA) from bacteria or viruses in the following matrices:

<table>
<thead>
<tr>
<th>Sample</th>
<th>DNA protocol</th>
<th>RNA protocol</th>
</tr>
</thead>
<tbody>
<tr>
<td>Biological</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Nasal/Pus Swab</td>
<td>X</td>
<td></td>
</tr>
<tr>
<td>Lymph Node Aspirates</td>
<td>X</td>
<td></td>
</tr>
<tr>
<td>Culture, including blood culture</td>
<td>X</td>
<td>X</td>
</tr>
<tr>
<td>Environmental</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Surface Swab</td>
<td>X</td>
<td>X</td>
</tr>
<tr>
<td>Powder</td>
<td>X</td>
<td>X</td>
</tr>
<tr>
<td>Water/PBS</td>
<td>X</td>
<td>X</td>
</tr>
</tbody>
</table>

PBS = phosphate buffered saline

Approximate time to process up to 12 samples is 1-2 hours. Ambient temperature is defined as 18-30°C.

Optimal Protocol Definition

Protocols with this symbol were optimized and verified for the detection of pathogens at low levels with the R.A.P.I.D. and JBAIDS detection systems.

Alternative Protocol Definition

Alternative protocols or tips given with this symbol have been tested, but not optimized, to add more flexibility for users. These protocols may not remove all inhibitors and may not achieve desired sensitivity.
**Equipment Required**

- Microcentrifuge capable of spinning 1.5 mL tubes 7,200-16,000 x g (RCF)
- Disruptor Genie (SI-D237) [or Vortex-Genie 2T (SI-T236) with Turbo Mix attachment] and adapter for 2.0 mL tubes (SI-0562) (Scientific Industries).
- Pipettors (Required range: 5-200 μL) and tips
- **RNA protocols only**: Heat block for 1.5 mL tubes
- **RNA protocols only**: Freezer (non frost-free, -20°C) (for Carrier RNA storage, Carrier RNA purchased separately in RNA Module)

**IT 1-2-3 SWIPE Kit Contents**

- Sample Swabs for surface, powder and culture sampling (40)
- Swab Snippers (2) and Wipes (40)
- Small Bead Tubes with beads (40)
- Small Spin Filters (40)
- Small Receiver Tubes (120)
- Small Transfer Pipets (for measuring 450 μL) (120)
- **Buffer 1** (Binding Buffer)
- **Buffer 2** (Wash Buffer)
- **Buffer 3** (Elution Buffer)
- SWIPE Instruction Booklet

**Additional Items Needed**

- **RNA protocols only**: Carrier RNA, provided in the IT 1-2-3 RNA Module (BFDf Part Number ASAY-ASY-0501)
- Blood culture only: Conical tube (10 mL or larger); PBS or other phosphate buffer (pH range 7.2-7.4), PBS packets: BFDf Part Number ASAY-ASY-0061 (water not included); water
Nasal/Pus Swab Protocol (DNA)

This protocol describes how to purify DNA from a nasal or pus swab sample. (A protocol to process RNA from a nasal swab is included in the IT 1-2-3 VIBE Kit.)

**Acceptable sample:** Nasal or pus sample on a swab and stored in a dry tube (Copan 168C rayon swab with twisted wire or equivalent is recommended).

**Note:** Calcium alginate swabs with aluminum shafts should not be used.

**Preparation:**
1. Remove swab from its tube. With provided Swab Snippers, cut off the end of the swab into a Small Bead Tube approximately ¼ inch above the head.
2. Discard the remaining part of the swab.
3. Clean Snippers with alcohol wipe.

**Lysis:**
4. Place Bead Tubes into 2 mL tube holder on the Disruptor Genie (or Vortex Genie with Turbo Mix) and Bead Beat on the highest setting for 5 minutes with lid down to disrupt cells or viruses and release nucleic acids.

**Bind filter:**
5. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 1 (Binding Buffer) to Small Bead Tube containing bead beaten sample. Keep transfer pipet for next step.
6. Use same transfer pipet to transfer liquid from Small Bead Tube to Small Spin Filter (packaged in a Small Receiver Tube), avoiding beads in the Bead Tube.
7. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

**Wash filter:**
9. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 2 (Wash Buffer) to Small Spin Filter.
10. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
12. Centrifuge for 3 minutes at maximum speed (min. 7,200 g) to remove residual ethanol.
Elute purified sample:

14. Use a Pipettor set to 200 µL to add **Buffer 3** (Elution Buffer) to the spin filter.
15. Incubate for 2 minutes at ambient temperature.
16. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
17. To enhance the recovery of nucleic acids, use the Pipettor set to 200 µL and repeat elution by pipetting the eluate (spin-through) from the bottom of the Receiver Tube back onto the Small Spin Filter.
18. Incubate for 2 minutes at ambient temperature.
19. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
20. Keep Receiver Tube containing the purified sample and test as soon as possible.

Storage and downstream analysis:

If the purified sample will not be tested within 30 minutes, it is recommended to store it at 2-8ºC and using it for downstream analysis within 4 hours. If testing does not occur within 4-8 hours, sensitivity may be affected due to degradation of target template.

<table>
<thead>
<tr>
<th>Shorter protocol</th>
<th>Steps 17-19 (re-elution) may be skipped if optimal sensitivity is not required.</th>
</tr>
</thead>
</table>
| No Pipettor option | Steps 14 and 17 can be replaced with:  
  “Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL **Buffer 3** (Elution Buffer)” or to repeat elution (if re-elution is necessary, the same transfer pipet must be used for both steps).  
  This option can be used (1) if optimal sensitivity is not required or (2) if using Freeze-dried reagents without a Reconstitution Buffer that allows hydration of reagents with sample only. |
Lymph Node Aspirate Protocol (DNA)

This protocol describes how to purify DNA from a lymph node aspirate sample.

Acceptable sample: Lymph node aspirate sample (approximately 20 µL) (can be collected on a swab and stored in a dry tube (Copan 168C rayon swab with twisted wire or equivalent is recommended). Note: Calcium alginate swabs with aluminum shafts should not be used.) If the sample does not contain liquid and consists of solid tissue larger than the equivalent of 20 µL, the solid tissue should be minced, mashed, or drawn through a needle in order to create a slurry of loose cells that can be picked up with a swab (see step 1 below.)

Preparation:

1. Use a Sample Swab to absorb liquid portion of the sample. If no liquid is present, touch sample with the swab in order to collect as much material as possible from the outside of the solid sample. A small amount of solid sample can be picked up with the swab (only enough to cover the lower quarter of the swab tip, do not pick up a large amount of solid sample or filter will clog).

2. If sample is stored on a swab: With provided Swab Snippers, cut off the end of the swab into a Small Bead Tube approximately ¼ inch above the head. Discard the remaining part of the swab. Clean Snippers with alcohol wipe, if used.

Lysis:

3. Place Bead Tubes into 2 mL tube holder on the Disruptor Genie (or Vortex Genie with Turbo Mix) and Bead Beat on the highest setting for 5 minutes with lid down to disrupt cells or viruses and release nucleic acids.

Bind filter:

4. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 1 (Binding Buffer) to Small Bead Tube containing bead beaten sample. Keep transfer pipet for next step.

5. Use same transfer pipet to transfer liquid from Small Bead Tube to Small Spin Filter (packaged in a Small Receiver Tube), avoiding beads in the Bead Tube.

6. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).


Wash filter:

8. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 2 (Wash Buffer) to Small Spin Filter.
9. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
11. Centrifuge for 3 minutes at maximum speed (min. 7,200 g) to remove residual ethanol.

**Elute purified sample:**

13. Use a Pipettor set to 200 µL to add Buffer 3 (Elution Buffer) to the spin filter.
14. Incubate for 2 minutes at ambient temperature.
15. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
16. To enhance the recovery of nucleic acids, use the Pipettor set to 200 µL and repeat elution by pipetting the eluate (spin-through) from the bottom of the Receiver Tube back onto the Small Spin Filter.
17. Incubate for 2 minutes at ambient temperature.
18. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
19. Keep Receiver Tube containing the purified sample and test as soon as possible.

**Storage and downstream analysis:**

If the purified sample will not be tested within 30 minutes, it is recommended to store it at 2-8°C and using it for downstream analysis within 4 hours. If testing does not occur within 4-8 hours, sensitivity may be affected due to degradation of target template.

<table>
<thead>
<tr>
<th>Shorter Protocol</th>
<th>Steps 16-18 (re-elution) may be skipped if optimal sensitivity is not required.</th>
</tr>
</thead>
<tbody>
<tr>
<td>No Pipettor Option</td>
<td>Steps 13 and 16 can be replaced with: “Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 3 (Elution Buffer)” or to repeat elution (if re-elution is necessary, the same transfer pipet must be used for both steps). This option can be used (1) if optimal sensitivity is not required or (2) if using Freeze-dried reagents without a Reconstitution Buffer that allows hydration of reagents with sample only.</td>
</tr>
</tbody>
</table>
Surface Swab Protocol (DNA)

This protocol is designed to purify DNA from pathogens present on surface swab samples.

Acceptable sample: Sample Swabs are included in the kit for surface swabbing. The swab sampling protocol is listed below. To maximize recovery of pathogens/nucleic acid from the swab, the swab is broken into the tube for bead beating and purified.

Surface Swab Sampling:
1. Wet a Sample Swab with liquid in a Small Bead Tube.
2. Thoroughly wipe a 2-inch x 2-inch surface area with the Swab.
3. Break the Sample Swab off in the Bead Tube at the swab breakpoint.

Lysis:
4. Place Bead Tubes into 2 mL tube holder on the Disruptor Genie (or Vortex Genie with Turbo Mix) and Bead Beat on the highest setting for 5 minutes with lid down to disrupt cells or viruses and release nucleic acids.

Bind filter:
5. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 1 (Binding Buffer) to Small Bead Tube containing bead beaten sample. Keep transfer pipet for next step.
6. Use same transfer pipet to transfer liquid from Small Bead Tube to Small Spin Filter (packaged in a Small Receiver Tube), avoiding beads in the Bead Tube.
7. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

Wash filter:
9. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 2 (Wash Buffer) to Small Spin Filter.
10. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
12. Centrifuge for 3 minutes at maximum speed (min. 7,200 g) to remove residual ethanol.
Elute purified sample:

14. Use a Pipettor set to 200 µL to add **Buffer 3** (Elution Buffer) to the spin filter.
15. Incubate for 2 minutes at ambient temperature.
16. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
17. To enhance the recovery of nucleic acids, use the Pipettor set to 200 µL and repeat elution by pipetting the eluate (spin-through) from the bottom of the Receiver Tube back onto the Small Spin Filter.
18. Incubate for 2 minutes at ambient temperature.
19. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
20. Keep Receiver Tube (fourth) containing the purified sample and test as soon as possible.

Storage and downstream analysis:

If the purified sample will not be tested within 30 minutes, it is recommended to store it at 2-8°C and using it for downstream analysis within 4 hours. If testing does not occur within 4-8 hours, sensitivity may be affected due to degradation of target template.

<table>
<thead>
<tr>
<th>Shorter protocol</th>
<th>Steps 17-19 (re-elution) may be skipped if optimal sensitivity is not required.</th>
</tr>
</thead>
</table>
| No Pipettor option | Steps 14 and 17 can be replaced with:  
  “Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL **Buffer 3** (Elution Buffer)” and to repeat elution (if re-elution is necessary, the same transfer pipet must be used for both steps).  
  This option can be used if (1) optimal sensitivity is not required or (2) if using Freeze-dried reagents without a Reconstitution Buffer that allows hydration of reagents with sample only. |
| Other swabs | If surface swabs are taken with a swab other than a Sample Swab, cut the head of the swab into the Small Bead Tube with the Swab Snippers. (See Nasal swab protocol.) |
| Direct testing: Bead beat only | Some surface swab samples do not contain downstream inhibitors and can be tested directly after bead beating. In this case, follow the protocol through step 4 and proceed to downstream analysis. Inhibition controls can be used with this protocol. |
Surface Swab Protocol (RNA)

This protocol is designed to purify RNA from pathogens present on surface swab samples. **Acceptable sample:** Sample Swabs are included in the kit for surface swabbing. The swab sampling protocol is listed below. To maximize recovery of pathogens/nucleic acid from the swab, the swab is broken into the tube for bead beating and purified.

**Surface Swab Sampling:**
1. Wet a Sample Swab with liquid in a Small Bead Tube.
2. Thoroughly wipe a 2-inch x 2-inch surface area with the Swab.
3. Break the Sample Swab off in the Bead Tube at the swab breakpoint.

**Preparation:**
4. Add 16 µL of **Carrier RNA** to each Small Bead Tube containing a swab. Cap Tube.

**Lysis:**
5. Place Bead Tubes into 2 mL tube holder on the Disruptor Genie (or Vortex Genie with Turbo Mix) and Bead Beat on the highest setting for 5 minutes with lid down to disrupt cells or viruses and release nucleic acids.

**Bind filter:**
6. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL **Buffer 1** (Binding Buffer) to Small Bead Tube containing bead beaten sample. Keep transfer pipet for next step.
7. Use same transfer pipet to transfer liquid from Small Bead Tube to Small Spin Filter (packaged in a Small Receiver Tube), avoiding beads in the Bead Tube.
8. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

**Wash filter:**
10. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL **Buffer 2** (Wash Buffer) to Small Spin Filter.
11. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
13. Centrifuge for 3 minutes at maximum speed (min. 7,200 g) to remove residual ethanol.

15. To remove more residual ethanol (important for RNA), incubate the uncapped tube with the Small Spin Filter inside, for 5 minutes in a dry heat block set to 88-95ºC.

**Elute purified sample:**

16. Use a Pipettor set to 200 µL to add **Buffer 3** (Elution Buffer) to the spin filter.

17. Incubate for 2 minutes at ambient temperature.

18. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

19. To enhance the recovery of nucleic acids, use the Pipettor set to 200 µL and repeat elution by pipetting the eluate (spin-through) from the bottom of the Receiver Tube back onto the Small Spin Filter.

20. Incubate for 2 minutes at ambient temperature.

21. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

22. Keep Receiver Tube (fourth) containing the purified sample and test as soon as possible.

**Storage and downstream analysis:**

If the purified sample will not be tested within 30 minutes, it is recommended to store it at 2-8ºC and using it for downstream analysis within 4 hours. If testing does not occur within 4-8 hours, sensitivity may be affected due to degradation of target template.

---

<table>
<thead>
<tr>
<th><strong>Shorter protocol</strong></th>
<th>Steps 19-21 (re-elution) may be skipped if optimal sensitivity is not required.</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Reduce Pipettor option</strong></td>
<td>Steps 16 and 19 can be replaced with: “Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL <strong>Buffer 3</strong> (Elution Buffer) and to repeat elution (if re-elution is necessary, the same transfer pipet must be used for both steps), if optimal sensitivity is not required.”</td>
</tr>
<tr>
<td><strong>Other swabs</strong></td>
<td>If surface swabs are taken with a swab other than a Sample Swab, cut the head of the swab into the Small Bead Tube with the Swab Snippers. (See Nasal swab protocol.)</td>
</tr>
<tr>
<td><strong>DNA and RNA from same sample</strong></td>
<td>The RNA protocol given here for surface swab samples can be used to purify both DNA and RNA from the same sample. Optimal sensitivity is not guaranteed for DNA detection using this RNA protocol.</td>
</tr>
</tbody>
</table>
Powder Protocol (DNA)

This protocol is designed to purify DNA from pathogens present in powder samples. **Acceptable sample:** A small amount (~ 1 mg) of powder picked up with a Sample Swab (see sampling protocol below).

**Powder Swab Sampling:**
1. Wet a Sample Swab with liquid in a Small Bead Tube.
2. Touch the powder with only the very tip of the moist Swab. (Note: Large quantities of powder can clog the filter in succeeding steps).
3. Break the Sample Swab off in the Bead Tube at the swab breakpoint.

**Lysis:**
4. Place Bead Tubes into 2 mL tube holder on the Disruptor Genie (or Vortex Genie with Turbo Mix) and Bead Beat on the highest setting for 5 minutes with lid down to disrupt cells or viruses and release nucleic acids.

**Bind filter:**
5. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 1 (Binding Buffer) to Small Bead Tube containing bead beaten sample. Keep transfer pipet for next step.
6. Use same transfer pipet to transfer liquid from Small Bead Tube to Small Spin Filter (packaged in a Small Receiver Tube), avoiding beads in the Bead Tube.
7. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

**Wash filter:**
9. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 2 (Wash Buffer) to Small Spin Filter.
10. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
12. Centrifuge for 3 minutes at maximum speed (min. 7,200 g) to remove residual ethanol.

**Elute purified sample:**
14. Use a Pipettor set to 200 µL to add Buffer 3 (Elution Buffer)
to the spin filter.

15. Incubate for 2 minutes at ambient temperature.

16. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

17. To enhance the recovery of nucleic acids, use the Pipettor set to 200 µL and repeat elution by pipetting the eluate (spin-through) from the bottom of the Receiver Tube back onto the Small Spin Filter.

18. Incubate for 2 minutes at ambient temperature.

19. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

20. Keep Receiver Tube (fourth) containing the purified sample and test as soon as possible.

Storage and downstream analysis:

If the purified sample will not be tested within 30 minutes, it is recommended to store it at 2-8°C and using it for downstream analysis within 4 hours. If testing does not occur within 4-8 hours, sensitivity may be affected due to degradation of target template.

<table>
<thead>
<tr>
<th>Shorter protocol</th>
<th>Steps 17-19 (re-elution) may be skipped if optimal sensitivity is not required.</th>
</tr>
</thead>
<tbody>
<tr>
<td>No Pipettor option</td>
<td>Steps 14 and 17 can be replaced with: “Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 3 (Elution Buffer)” and to repeat elution (if re-elution is necessary, the same transfer pipet must be used for both steps). This option can be used if (1) optimal sensitivity is not required or (2) if using Freeze-dried reagents without a Reconstitution Buffer that allows hydration of reagents with sample only.</td>
</tr>
<tr>
<td>Direct testing: Bead beat only</td>
<td>Some powder samples do not contain downstream inhibitors and can be tested directly after bead beating. In this case, follow the protocol through step 4 and proceed to downstream analysis. With powder samples (usually highly concentrated) it is advisable to prepare a ten-fold dilution of the bead-beaten sample in a Small Receiver Tube with Buffer 3 (for example 40 µL sample plus 360 µL Buffer 3). Inhibition controls can be used with this protocol.</td>
</tr>
</tbody>
</table>
**Powder Protocol (RNA)**

This protocol is designed to purify RNA from pathogens present in powder samples. **Acceptable sample:** A small amount (~1 mg) of powder picked up with an Sample Swab (see sampling protocol below).

**Powder Swab Sampling:**
1. Wet a Sample Swab with liquid in a Small Bead Tube.
2. Touch the powder with only the very tip of the moist Swab. (Note: Large quantities of powder can clog the filter in proceeding steps).
3. Break the Sample Swab off in the Bead Tube at the swab breakpoint.

**Preparation:**
4. Add 16 µL of **Carrier RNA** to each Small Bead Tube containing a swab. Cap Tube.

**Lysis:**
5. Place Bead Tubes into 2 mL tube holder on the Disruptor Genie (or Vortex Genie with Turbo Mix) and Bead Beat on the highest setting for 5 minutes with lid down to disrupt cells or viruses and release nucleic acids.

**Bind filter:**
6. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL **Buffer 1** (Binding Buffer) to Small Bead Tube containing bead beaten sample. Keep transfer pipet for next step.
7. Use same transfer pipet to transfer liquid from Small Bead Tube to Small Spin Filter (packaged in a Small Receiver Tube), avoiding beads in the Bead Tube.
8. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

**Wash filter:**
10. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL **Buffer 2** (Wash Buffer) to Small Spin Filter.
11. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
13. Centrifuge for 3 minutes at maximum speed (min. 7,200 g) to remove residual ethanol.
(fourth). Discard old Tube.

15. To remove more residual ethanol (important for RNA), incubate the uncapped tube with the Small Spin Filter inside, for 5 minutes in a dry heat block set to 88-95°C.

**Elute purified sample:**

16. Use a Pipettor set to 200 µL to add **Buffer 3** (Elution Buffer) to the spin filter.

17. Incubate for 2 minutes at ambient temperature.

18. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

19. To enhance the recovery of nucleic acids, use the Pipettor set to 200 µL and repeat elution by pipetting the eluate (spin-through) from the bottom of the Receiver Tube back onto the Small Spin Filter.

20. Incubate for 2 minutes at ambient temperature.

21. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

22. Keep Receiver Tube (fourth) containing the purified sample and test as soon as possible.

**Storage and downstream analysis:**

If the purified sample will not be tested within 30 minutes, it is recommended to store it at 2-8°C and using it for downstream analysis within 4 hours. If testing does not occur within 4-8 hours, sensitivity may be affected due to degradation of target template.

<table>
<thead>
<tr>
<th>Shorter protocol</th>
<th>Steps 19-21 (re-elution) may be skipped if optimal sensitivity is not required.</th>
</tr>
</thead>
<tbody>
<tr>
<td>Reduce Pipettor option</td>
<td>Steps 16 and 19 can be replaced with: “Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL <strong>Buffer 3</strong> (Elution Buffer)” and to repeat elution (if re-elution is necessary, the same transfer pipet must be used for both steps), if optimal sensitivity is not required.</td>
</tr>
<tr>
<td>DNA and RNA from same sample</td>
<td>The RNA protocol given here for powder samples can be used to purify both DNA and RNA from the same sample. Optimal sensitivity is not guaranteed for DNA detection using this RNA protocol.</td>
</tr>
</tbody>
</table>
Culture Protocol (Including Blood Culture; DNA)

Use the alternative optimized protocol for processing plated colonies intended for use with in vitro diagnostic products.

Organisms cultured on solid (plates) or in liquid media generally produce highly concentrated samples. Sample volumes for this protocol are therefore small. **Acceptable sample:** Bacterial cultures or viral cell culture with appropriate culture media.

1. **For plated colonies or plaques:** touch a Sample Swab to the colony or plaque and break off the swab at the breakpoint into a Small Bead Tube.

   **For liquid culture:** dip a Sample Swab in culture media, allow it to soak for 3-5 seconds and break off the swab at the breakpoint in a Small Bead Tube. (Alternatively, pipette 40-50 µL the liquid culture directly into a bead tube.)

   **For blood culture:** First dilute the blood culture media by adding 50 µL of blood culture media to 5 mL of PBS in a conical tube (or equivalent). Mix well. Transfer 50 µL of the diluted blood culture media into a Small Bead Tube.

**Lysis:**

2. Place Bead Tubes into 2 mL tube holder on the Disruptor Genie (or Vortex Genie with Turbo Mix) and Bead Beat on the highest setting for 5 minutes with lid down to disrupt cells or viruses and release nucleic acids.

**Bind filter:**

3. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL **Buffer 1** (Binding Buffer) to Small Bead Tube containing bead beaten sample. Keep transfer pipet for next step.

4. Use same transfer pipet to transfer liquid from Small Bead Tube to Small Spin Filter (packaged in a Small Receiver Tube), avoiding beads in the Bead Tube.

5. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).


**Wash filter:**

7. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL **Buffer 2** (Wash Buffer) to Small Spin Filter.

8. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).


10. Centrifuge for 3 minutes at maximum speed (min. 7,200 g) to remove residual ethanol.

**Elute purified sample:**

12. Use a Pipettor set to 200 µL to add **Buffer 3** (Elution Buffer) to the spin filter.
13. Incubate for 2 minutes at ambient temperature.
14. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
15. To enhance the recovery of nucleic acids, use the Pipettor set to 200 µL and repeat elution by pipetting the eluate (spin-through) from the bottom of the Receiver Tube back onto the Small Spin Filter.
16. Incubate for 2 minutes at ambient temperature.
17. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
18. Keep Receiver Tube (fourth) containing the purified sample and test as soon as possible.

**Storage and downstream analysis:**

If the purified sample will not be tested within 30 minutes, it is recommended to store it at 2-8°C and using it for downstream analysis within 4 hours. If testing does not occur within 4-8 hours, sensitivity may be affected due to degradation of target template.

---

### Alternative Optimized Protocol

| Direct testing: Bead beat only | Some bacterial colonies can be tested directly after bead beating. Follow the protocol through step 2 and immediately prior to testing:  
1) Add 5 µL bead-beaten sample to a Small Receiver Tube  
2) Add 500 µL Buffer 3 and proceed with downstream analysis  
Use of inhibition controls with this protocol is recommended. |
<table>
<thead>
<tr>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>Shorter protocol</td>
<td>Steps 15-17 (re-elution) may be skipped if optimal sensitivity is not required.</td>
</tr>
</tbody>
</table>
| No Pipettor option          | Steps 12 and 15 can be replaced with:  
   “Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL **Buffer 3** (Elution Buffer)” and to repeat elution (if re-elution is necessary, the same transfer pipet must be used for both steps).  
   This option can be used if (1) optimal sensitivity is not required or (2) if using Freeze-dried reagents without a Reconstitution Buffer that allows hydration of reagents with sample only. |
**Culture Protocol (RNA)**

Organisms cultured on solid (plates) or in liquid media generally produce highly concentrated samples. Sample volumes for this protocol are therefore small.

**Acceptable sample:** Viral cell culture with appropriate culture media.

1. **For plated colonies or plaques:** touch a Sample Swab to the plaque and break off the swab at the breakpoint into a Small Bead Tube.

   **For liquid culture:** dip a Sample Swab in culture media, allow it to soak for 5 seconds and break off the swab at the breakpoint in a Small Bead Tube. (Alternatively, pipette 40-50 µL the liquid culture directly into a bead tube.)

**Preparation:**

2. Add 16 µL of **Carrier RNA** to each Small Bead Tube containing a swab. Cap Tube.

**Lysis:**

3. Place Bead Tubes into 2 mL tube holder on the Disruptor Genie (or Vortex Genie with Turbo Mix) and Bead Beat on the highest setting for 5 minutes with lid down to disrupt cells or viruses and release nucleic acids.

**Bind filter:**

4. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL **Buffer 1** (Binding Buffer) to Small Bead Tube containing bead beaten sample. Keep transfer pipet for next step.

5. Use same transfer pipet to transfer liquid from Small Bead Tube to Small Spin Filter (packaged in a Small Receiver Tube), avoiding beads in the Bead Tube.

6. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).


**Wash filter:**

8. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL **Buffer 2** (Wash Buffer) to Small Spin Filter.

9. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).


11. Centrifuge for 3 minutes at maximum speed (min. 7,200 g) to remove residual ethanol.

13. To remove more residual ethanol (important for RNA), incubate the uncapped tube with the Small Spin Filter inside, for 5 minutes in a dry heat block set to 88-95ºC.

Elute purified sample:

14. Use a Pipettor set to 200 µL to add **Buffer 3** (Elution Buffer) to the spin filter.
15. Incubate for 2 minutes at ambient temperature.
16. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
17. To enhance the recovery of nucleic acids, use the Pipettor set to 200 µL and repeat elution by pipetting the eluate (spin-through) from the bottom of the Receiver Tube back onto the Small Spin Filter.
18. Incubate for 2 minutes at ambient temperature.
19. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
20. Keep Receiver Tube (fourth) containing the purified sample and test as soon as possible.

**Storage and downstream analysis:**
If the purified sample will not be tested within 30 minutes, it is recommended to store it at 2-8ºC and using it for downstream analysis within 4 hours. If testing does not occur within 4-8 hours, sensitivity may be affected due to degradation of target template.

<table>
<thead>
<tr>
<th>Shorter protocol</th>
<th>Steps 17-19 (re-elution) may be skipped if optimal sensitivity is not required.</th>
</tr>
</thead>
<tbody>
<tr>
<td>Reduce Pipettor option</td>
<td>Steps 14 and 17 can be replaced with: “Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL <strong>Buffer 3</strong> (Elution Buffer)” and to repeat elution (if re-elution is necessary, the same transfer pipet must be used for both steps), if optimal sensitivity is not required.</td>
</tr>
</tbody>
</table>
Water/PBS Protocol (DNA) - Alternative Protocol

This protocol is an alternative to the Water and PBS protocols in the IT 1-2-3 FLOW Kit. It processes less liquid, therefore sensitivity of detection is not as good with this protocol, but it is a good option for users with only small equipment (smaller centrifuge, etc.). This protocol has not been used for sensitivity verification with the RAPID or JBAIDS detection systems.

Acceptable sample: Water or PBS (with or without Triton X-100).

1. Pipette 100 µL Water or PBS into a Small Bead Tube (alternatively, for 40 µL, dip a Sample Swab into the liquid, allow it to soak for 3-5 seconds and break off the swab at the breakpoint into a Small Bead Tube).

Lysis:
2. Place Bead Tubes into 2 mL tube holder on the Disruptor Genie (or Vortex Genie with Turbo Mix) and Bead Beat on the highest setting for 5 minutes with lid down to disrupt cells or viruses and release nucleic acids.

Bind filter:
3. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 1 (Binding Buffer) to Small Bead Tube containing bead beaten sample. Keep transfer pipet for next step.
4. Use same transfer pipet to transfer liquid from Small Bead Tube to Small Spin Filter (packaged in a Small Receiver Tube), avoiding beads in the Bead Tube.
5. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

Wash filter:
7. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 2 (Wash Buffer) to Small Spin Filter.
8. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
10. Centrifuge for 3 minutes at maximum speed (min. 7,200 g) to remove residual ethanol.

Elute purified sample:
12. Use a Pipettor set to 200 µL to add Buffer 3 (Elution Buffer) to the spin filter.
13. Incubate for 2 minutes at ambient temperature.

14. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

15. To enhance the recovery of nucleic acids, use the Pipettor set to 200 µL and repeat elution by pipetting the eluate (spin-through) from the bottom of the Receiver Tube back onto the Small Spin Filter.

16. Incubate for 2 minutes at ambient temperature.

17. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

18. Keep Receiver Tube (fourth) containing the purified sample and test as soon as possible.

Storage and downstream analysis:

If the purified sample will not be tested within 30 minutes, it is recommended to store it at 2-8ºC and using it for downstream analysis within 4 hours. If testing does not occur within 4-8 hours, sensitivity may be affected due to degradation of target template.

<table>
<thead>
<tr>
<th>Shorter protocol</th>
<th>Steps 15-17 (re-elution) may be skipped if optimal sensitivity is not required.</th>
</tr>
</thead>
<tbody>
<tr>
<td>No Pipettor option</td>
<td>Steps 12 and 15 can be replaced with:</td>
</tr>
<tr>
<td></td>
<td>“Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 3 (Elution Buffer)” and to repeat elution (if re-elution is necessary, the same transfer pipet must be used for both steps).</td>
</tr>
<tr>
<td></td>
<td>This option can be used if (1) optimal sensitivity is not required or (2) if using Freeze-dried reagents without a Reconstitution Buffer that allows hydration of reagents with sample only.</td>
</tr>
<tr>
<td>Direct testing: Bead beat only</td>
<td>Some samples do not contain downstream inhibitors and can be tested directly after bead beating. In this case, follow the protocol through step 2 and proceed to downstream analysis. To dilute out inhibitors, prepare a ten-fold dilution of the bead-beaten sample in a Small Receiver Tube with Buffer 3 (for example 40 µL sample plus 360 µL Buffer 3). Inhibition controls can be used with this protocol.</td>
</tr>
</tbody>
</table>
Water/PBS Protocol (RNA) Alternative Protocol

This protocol is an alternative to the Water and PBS protocols in the IT 1-2-3 FLOW Kit. It processes less liquid, therefore sensitivity of detection is not as good with this protocol, but it is a good option for users with only small equipment (smaller centrifuge, etc.). This protocol has not been used for sensitivity verification with the RAPID or JBAIDS detection systems.

Acceptable sample: Water or PBS (with or without Triton X-100).

1. Pipette 100 µL Water or PBS into a Small Bead Tube (alternatively, for 40 µL, dip a Sample Swab into the liquid, allow it to soak for 3-5 seconds and break off the swab at the breakpoint into a Small Bead Tube).

Preparation:
2. Add 16 µL of Carrier RNA to each Small Bead Tube containing a swab. Cap Tube.

Lysis:
3. Place Bead Tubes into 2 mL tube holder on the Disruptor Genie (or Vortex Genie with Turbo Mix) and Bead Beat on the highest setting for 5 minutes with lid down to disrupt cells or viruses and release nucleic acids.

Bind filter:
4. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 1 (Binding Buffer) to Small Bead Tube containing bead beaten sample. Keep transfer pipet for next step.
5. Use same transfer pipet to transfer liquid from Small Bead Tube to Small Spin Filter (packaged in a Small Receiver Tube), avoiding beads in the Bead Tube.
6. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

Wash filter:
8. Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 2 (Wash Buffer) to Small Spin Filter.
9. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).
11. Centrifuge for 3 minutes at maximum speed (min. 7,200 g) to remove residual ethanol.

13. To remove more residual ethanol (important for RNA), incubate the uncapped tube with the Small Spin Filter inside, for 5 minutes in a dry heat block set to 88-95ºC.

**Elute purified sample:**

14. Use a Pipettor set to 200 µL to add Buffer 3 (Elution Buffer) to the spin filter.

15. Incubate for 2 minutes at ambient temperature.

16. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

17. To enhance the recovery of nucleic acids, use the Pipettor set to 200 µL and repeat elution by pipetting the eluate (spin-through) from the bottom of the Receiver Tube back onto the Small Spin Filter.

18. Incubate for 2 minutes at ambient temperature.

19. Centrifuge for 2 minutes at maximum speed (min. 7,200 g).

20. Keep Receiver Tube (fourth) containing the purified sample and test as soon as possible.

**Storage and downstream analysis:**

If the purified sample will not be tested within 30 minutes, it is recommended to store it at 2-8ºC and using it for downstream analysis within 4 hours. If testing does not occur within 4-8 hours, sensitivity may be affected due to degradation of target template.

<table>
<thead>
<tr>
<th>Shorter protocol</th>
<th>Steps 17-19 (re-elution) may be skipped if optimal sensitivity is not required.</th>
</tr>
</thead>
<tbody>
<tr>
<td>Reduce Pipettor option</td>
<td>Steps 14 and 17 can be replaced with: “Use a Small Transfer Pipet (fill to bottom of bulb) to add 450 µL Buffer 3 (Elution Buffer)” and to repeat elution (if re-elution is necessary, the same transfer pipet must be used for both steps) if optimal sensitivity is not required.</td>
</tr>
<tr>
<td>DNA and RNA from same sample</td>
<td>The RNA protocol given here for water/PBS samples can be used to purify both DNA and RNA from the same sample. Optimal sensitivity is not guaranteed for DNA detection using this RNA protocol.</td>
</tr>
</tbody>
</table>
## Troubleshooting

<table>
<thead>
<tr>
<th>Symptom</th>
<th>Resolution</th>
</tr>
</thead>
<tbody>
<tr>
<td>Spin Filter Clogging</td>
<td>If the Small Spin Filter clogs, follow protocol up to the addition of Buffer 2. Before adding Buffer 2, pour liquid off of the top of the filter and continue with the protocol.</td>
</tr>
<tr>
<td>Inhibited Sample</td>
<td>Sometimes unknown inhibitors of downstream analysis are not adequately removed from a purified sample. If amplification does not occur as expected or an inhibition control shows inhibitors are present, perform a ten-fold dilution of that sample in Reagent Grade Water or Buffer 3. A ten-fold dilution is usually adequate to remove the effects of the inhibitors, however sensitivity is decreased. Note: If inhibition controls are used, refer to the reagent product insert for specific directions.</td>
</tr>
</tbody>
</table>
### Sample Purification Kits and Supplies

<table>
<thead>
<tr>
<th>Item</th>
<th>Contents</th>
<th>Part No.</th>
</tr>
</thead>
<tbody>
<tr>
<td>IT 1-2-3 DNA Sample Purification Kit</td>
<td>Sample Purification and Extraction Kit for minimally trained technicians to extract DNA from environmental sources</td>
<td>3800</td>
</tr>
<tr>
<td>IT 1-2-3 SWIPE Sample Purification Kit*</td>
<td>Sample Purification Kit for purification of nasal swab, surface swab, live culture and powder samples</td>
<td>ASAY-ASY-0005</td>
</tr>
<tr>
<td>IT 1-2-3 FLOW Sample Purification Kit**</td>
<td>Sample Purification Kit for purification of blood, air, water, food and body fluid samples</td>
<td>ASAY-ASY-0004</td>
</tr>
<tr>
<td>IT 1-2-3 VIBE Sample Purification Kit</td>
<td>Sample Purification Kit for purification of sputum samples and RNA from blood and nasal swab samples</td>
<td>ASAY-ASY-0500</td>
</tr>
<tr>
<td>IT 1-2-3 SCOOP Sample Purification Kit</td>
<td>Sample Purification Kit for purification of stool and soil samples</td>
<td>ASAY-ASY-0502</td>
</tr>
<tr>
<td>IT 1-2-3 QFLOW DNA Sample Purification Kit*</td>
<td>Sample Purification Kit for purification of blood, air, water, food and body fluid samples</td>
<td>ASAY-ASY-0503</td>
</tr>
<tr>
<td>IT 1-2-3 QFLOW RNA Sample Purification Kit</td>
<td>Sample Purification Kit for purification of blood, air, water, food and body fluid samples</td>
<td>ASAY-ASY-0504</td>
</tr>
<tr>
<td>IT 1-2-3 Platinum Path Sample Purification Kit</td>
<td>Sample Purification Kit for magnetic bead purification of biological, environmental, and food samples</td>
<td>ASAY-ASY-0120</td>
</tr>
<tr>
<td>IT 1-2-3 RNA Module</td>
<td>An accessory to the SWIPE, FLOW, VIBE, and QFLOW DNA purification kits and contains items for the purification of RNA</td>
<td>ASAY-ASY-0501</td>
</tr>
<tr>
<td>Filtered Blender Bags</td>
<td>1 Bag of 10 (Brinkmann)</td>
<td>ASAY-ASY-0060</td>
</tr>
<tr>
<td>PBS packets</td>
<td>Phosphate Buffered Saline, pH 7.4 PBS powder, 10 packets per box (each packet makes 1 L PBS) (Sigma-Aldrich)</td>
<td>ASAY-ASY-0061</td>
</tr>
<tr>
<td>Triton X-100</td>
<td>100 mL Triton (Sigma-Aldrich)</td>
<td>ASAY-ASY-0062</td>
</tr>
<tr>
<td>Stool Collection Vials (Cary Blair)</td>
<td>Box of 20 (PROTOCOL-Fisher)</td>
<td>ASAY-ASY-0063</td>
</tr>
<tr>
<td>Stool Concentrator Filters</td>
<td>Box of 30 (PARAPAK, Meridian Diagnostics))</td>
<td>ASAY-ASY-0064</td>
</tr>
</tbody>
</table>

*These purification kits require the RNA Module (ASAY-ASY-0501) for some preparation applications.

**Large bead tube adapter (P/N PREP-ASY-0001) is required for this kit, but not included.
<table>
<thead>
<tr>
<th>Sample Type</th>
<th>Platinum Path</th>
<th>SWIPE</th>
<th>VIBE</th>
<th>SCOOP</th>
<th>FLOW</th>
<th>RNA Module</th>
</tr>
</thead>
<tbody>
<tr>
<td>Whole Blood</td>
<td>X</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Nasal/Pus Swabs</td>
<td>X</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Culture</td>
<td>X</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Stool</td>
<td>X</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Gastric Wash</td>
<td>X</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Lymph Node Aspirates</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Cerebral Spinal Fluid</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Nasopharyngeal Swab and Throat Swab</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Air (PBS)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Surface Swabs</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Powder</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Water</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Soil</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Milk</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Mixed Greens</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Ground Beef</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Tuna Salad</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

* = Alternative protocols
For additional information regarding our products and applications, please contact our Customer Service Department at BioFire Defense.