

# Sample Sizes for Surveillance and Suspected Contamination Sampling

Surveillance is the routine collection of information used to characterize risk with clearly established intervention points used to monitor and maintain animal health. Surveillance also utilizes thresholds to dictate further action either through sampling, implementing mitigation strategies, or a combination of both. If surveillance thresholds have been met or exceeded, it is time to transition to suspected contamination sampling. Pre-determined sample sizes for surveillance sampling with thresholds (Table 1) and suspected contamination sampling with return to surveillance sampling thresholds can be found at the end of this resource (Table 2). This resource will detail how to interpret those pre-determined sample sizes for surveillance and suspected contamination. If electing to calculate sample sizes yourself, consult the additional resource titled "Calculating Sample Sizes and Thresholds" or if needing more information on how to coordinate the transition to suspected contamination sampling, consult the additional resource titled "Transitioning from Surveillance Sampling to Suspected Contamination Sampling."

The pre-determined sample sizes depends on two factors, the probability of feed serving as a source for pathogen of interest and the severity of the pathogen in regards to species of interest. There are spectrums, from high to very low, within these two factors that will determine sample size.

## ***Probability of feed serving as a source for pathogen of interest***

Probability of feed serving as a source for pathogen of interest takes in account the potential feed ingredients and mitigation strategies already implemented at the feed mill.

- **High probability**
  - High probability indicates that there is immediate danger that the hazard will occur.
  - If there are no mitigation techniques in place at a feed mill, then this is the proper designation.
- **Medium probability**
  - Medium probability indicates that the hazard will probably occur if not controlled.
  - If a feed mill utilizes only point-in-time mitigation techniques, this is the appropriate designation.
    - Examples of point-in-time mitigation techniques include quarantining or holding ingredients, thermally processing feed, implementation of feed batch sequencing, or implementation of flushes after manufacturing certain diets.
    - These techniques can only guarantee that potential contamination has been reduced or infectivity of pathogen reduced, but doesn't prevent recontamination.
  - If a feed mill has or utilizes rendered ingredients for diets, this is the appropriate designation.
    - Rendered ingredients are manufactured at a temperature range of 240-290°F for at least 40-90 minutes which has been shown to reduce pathogen contamination (Hamilton, 2006). However, this temperature range does not prevent recontamination during further feed manufacturing or delivery.
    - Transportation of these ingredients from rendering facilities also has a risk of pathogen introduction to a feed mill (Lowe et al., 2014) while these types of ingredients have been

shown to better support pathogen survival when compared to plant based ingredients (Dee et al., 2018)

- **Low probability**

- Low probability indicates that it's possible for hazard to occur if not controlled.
- If a feed mill utilizes a chemical feed additive as a means to reduce pathogen contamination or infectivity, this is the appropriate designation.
  - Chemical feed additives have been shown to reduce pathogens in feed at time of application and remain active throughout the feed supply chain (Stewart et al., 2020).

- **Very low probability**

- Very low probability indicates that it's unlikely for the hazard to occur and an assumption that the hazard will not occur is warranted.
- If a feed mill utilizes point-in-time mitigation techniques in combination with a chemical feed additive, this is the appropriate designation.

### ***Severity of the pathogen of interest in regards to species of interest***

The severity of the pathogen of interest in regards to species of interest is based on the consequences of the pathogen of interest if introduced into the production system via the feed supply chain. The type of production system served by the feed mill and the production system's definition of mortality and morbidity will influence the designated severity.

- **High severity**

- Pathogen of interest would cause high mortality and high morbidity if introduced into the production system.

- **Medium severity**

- Pathogen of interest would cause high mortality and low morbidity if introduced into the production system.

- **Low severity**

- Pathogen of interest would cause low mortality and high morbidity if introduced into the production system.

- **Very low severity**

- Pathogen of interest would cause low mortality and low morbidity if introduced into the production system.

**Table 1:** Recommendations for surveillance sample size and thresholds based on severity of pathogen of interest and probability of pathogen being introduced through feed.

<div>Severity</div> <div>Probability</div>	<b>HIGH</b> Pathogen of interest would cause high mortality and high morbidity	<b>MEDIUM</b> Pathogen of interest would cause high mortality and low morbidity	<b>LOW</b> Pathogen of interest would cause low mortality and high morbidity	<b>VERY LOW</b> Pathogen of interest would cause low mortality and low morbidity
<b>HIGH</b> Immediate danger that the hazard will occur.	<u>75 samples/week:</u> 10 feed samples 65 environmental samples <b>Threshold = 1 positive</b>	<u>75 samples/week:</u> 10 feed samples 65 environmental samples <b>Threshold = 1 positive</b>	<u>15 samples/week:</u> 5 feed samples 10 environmental samples <b>Threshold = 1 positive</b>	<u>5 samples/week:</u> 1 feed sample 4 environmental samples <b>Threshold = 1 positive</b>
<b>MEDIUM</b> Hazard will <u>probably</u> occur if not controlled.	<u>75 samples/week:</u> 5 feed samples 70 environmental samples <b>Threshold = 1 positive</b>	<u>15 samples/week:</u> 2 feed samples 13 environmental samples <b>Threshold = 2 positives</b>	<u>8 samples/week:</u> 2 feed samples 6 environmental samples <b>Threshold = 2 positives</b>	<u>5 samples/week:</u> 0 feed samples 5 environmental samples <b>Threshold = 2 positives</b>
<b>LOW</b> It's <u>possible</u> for hazard to occur if not controlled.	<u>25 samples/week:</u> 1 feed sample 24 environmental samples <b>Threshold = 1 positive</b>	<u>15 samples/week:</u> 1 feed sample 14 environmental samples <b>Threshold = 2 positives</b>	<u>8 samples/week:</u> 1 feed sample 7 environmental samples <b>Threshold = 3 positives</b>	<u>4 samples/week:</u> 0 feed samples 4 environmental samples <b>Threshold = 2 positives</b>
<b>VERY LOW</b> It's unlikely for the hazard to occur and can assume that hazard will not occur.	<u>25 samples/week:</u> 0 feed samples 25 environmental samples <b>Threshold = 1 positive</b>	<u>15 samples/week:</u> 0 feed samples 15 environmental samples <b>Threshold = 2 positives</b>	<u>8 samples/week:</u> 0 feed samples 8 environmental samples <b>Threshold = 3 positives</b>	<u>3 samples/week:</u> 0 feed samples 3 environmental samples <b>Threshold = 2 positives</b>

**Table 2:** Recommendations for suspected contamination sample size and when to return to surveillance sampling based on severity of pathogen of interest and probability of pathogen of interest introduced through feed.

<div>Severity</div> <div>Probability</div>	<b>HIGH</b> Pathogen of interest would cause high mortality and high morbidity	<b>MEDIUM</b> Pathogen of interest would cause high mortality and low morbidity	<b>LOW</b> Pathogen of interest would cause low mortality and high morbidity	<b>VERY LOW</b> Pathogen of interest would cause low mortality and low morbidity
<b>HIGH</b> Immediate danger that the hazard will occur.	<u>300 samples:</u> 102 feed samples 198 environmental samples <b>Return to surveillance: no more than 3 positive samples</b>		<u>100 samples:</u> 25 feed samples 75 environmental samples <b>Return to surveillance: no more than 3 positive sample</b>	
<b>MEDIUM</b> Hazard will <u>probably</u> occur if not controlled.				
<b>LOW</b> It's <u>possible</u> for hazard to occur if not controlled.	<u>100 samples:</u> 15 feed sample 85 environmental samples <b>Return to surveillance: no more than 3 positive sample</b>		<u>60 samples:</u> 5 feed samples 55 environmental samples <b>Return to surveillance: no more than 3 positive sample</b>	
<b>VERY LOW</b> It's unlikely for the hazard to occur and can assume that hazard will not occur.				

## References

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# Calculating Sample Sizes and Thresholds

A challenge when trying to implement sampling programs within a feed mill is defining proper calculation of the necessary sample size to detect pathogens of interest. If interested in pre-determined sample size recommendations and thresholds, consult the additional resource titled “Sample Size for Surveillance and Suspected Contamination.” However, if a production system or feed mill has a general idea of prevalence rate for the pathogen of interest and would like to defer from the general recommendations, this resource aims to explain how to calculate sample size, basis for sample size, and how to set thresholds for sampling feed mills. If interested in how to transition from surveillance to suspected contamination sampling, consult the additional resource titled “Transitioning from Surveillance Sampling to Suspected Contamination Sampling.”

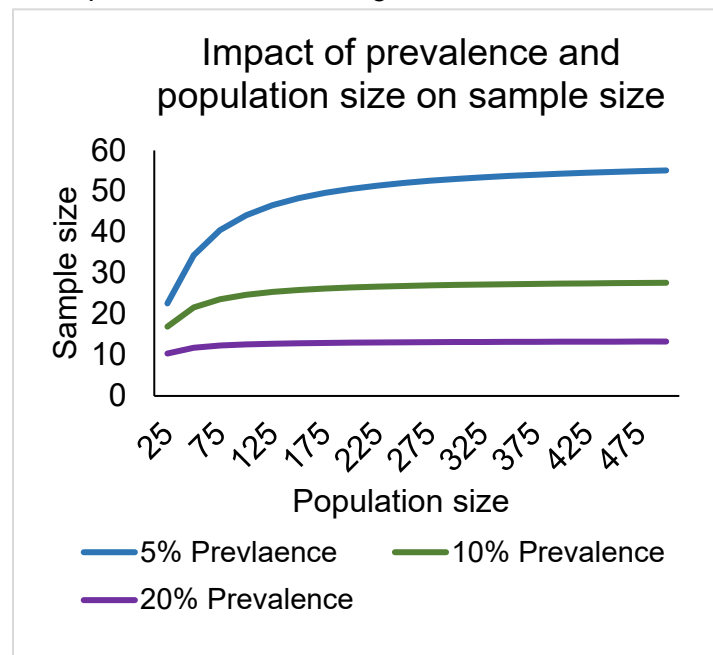
## Sample size formula

The formula to determine sample size is the same as that which is used for sampling to detect disease in animals (Dahoo et al., 2014). This formula is used to determine the sample size necessary to have confidence in the outcome while minimizing interpretation error. It takes in to account the number of animals from within a population that must be sampled from a population to have a given level of confidence that at least one sample would be positive based on a given prevalence level. The sample size,  $n$ , is determined by the confidence interval ( $\alpha = 1 - \text{confidence level}$ ), the population size ( $N$ ), and estimated minimum number of diseases animals in the group ( $D = \text{estimated prevalence population size}$ ):

$$n = \left[ \left( 1 - \alpha^{1/D} \right) \times \left( N - \frac{D - 1}{2} \right) \right]$$

However, when considering feed mills and the presence of a pathogen of interest, the feed mill is more concerned about detecting the pathogen of interest within feed or feed mill environment and not clinical disease within an animal. So in this instance, the sample size could be thought of as the number of total samples to take at the feed mill, the population could be thought of as the possible number of samples to take either in feed or in the environment, and the prevalence as the perceived prevalence of the pathogen of interest within the feed mill. When interpreting the formula in a scenario like this, it can be inferred that the number of possible samples to be collected approaches infinity because samples could be taken per ton, per pound, or per gram.

As shown in the graph on the right, when utilizing the sampling to detect disease formula, the sample size for a desired prevalence level will plateau. This is due to there being minimal changes within the sample sizes as the population size increases.



So if looking for a desired prevalence level within a population, identify where the sample size plateaus for the prevalence level, and utilize the sample size corresponding with the start of the plateau. The graph on the right also illustrates that as prevalence rate increases, the sample size decreases which is because the pathogen of interest is present at higher percentages and thus, requires a smaller sample size to detect it within the population. When consulting the pre-determined sample sizes for surveillance, the basis for sample size recommendations was based on this same principle.

### **Sample size references**

When trying to set sample sizes for feed and environmental samples, there are some peer-reviewed works to reference for naturally contaminated complete feed, feed ingredients, or feed mill environments which can be used as reference ranges. These sources evaluated the prevalence, or the number of samples containing detectable pathogen of interest divided by the total number of samples collected, for some pathogens within feed mills. Feed samples are defined as samples pulled directly from the source (either from feed ingredients or complete feed) while environmental samples are samples from surfaces that are sampled with their respective materials. The data from these published works give reference ranges for prevalence of pathogens and can help guide surveillance sampling. It is important to note, that when considering natural contaminated feed, these papers found that on average, 14% of feed samples tested positive for the pathogen of interest while on average, 22% of environmental samples tested positive for the pathogen of interest (Table 1). These prevalence rates are lower than when compared to experimentally inoculated studies indicating that surveillance sampling should reflect this trend when considering natural contamination.

### **Thresholds**

Thresholds are action points to designate further action because the prevalence rate has increased to an unwanted rate during surveillance sampling. Thresholds can be adjusted to be stricter or more lenient with increased prevalence rates depending upon the production system and how risk averse they choose to be.

### **References**

Dahoo, I., Martin, W., and Stryhm H. 2014. Sampling. In: S Marget McPike, editor, Veterinary Epidemiologic Research. VER Inc. Charlottetown, Prince Edward Island, Canada. p. 33-56.

**Table 1.** Prevalence rates from available published data where pathogens were naturally present in feed ingredients, complete feed, or feed mill environments.

Item	Reference doi	Pathogen <sup>1</sup>	Prevalence, %	Total Number of Samples Collected
Environmental Samples:				
Elijah et al., 2022	10.54846/jshap/1250	PDCoV	2.33	86
Elijah et al., 2022	10.54846/jshap/1250	PEDV	2.33	86
Garrido-Mantillo et al., 2022	10.1111/tbed.14354	PEDV	37.50	8
Gebhardt et al., 2021	10.1111/tbed.14335	ASFV	0.73	2186
Magossi et al., 2019	10.1002/mbo3.711	Salmonella sp.	66.24	237
Magossi et al., 2019	10.1002/mbo3.711	<i>Salmonella enterica</i>	19.75	157
Feed Samples:				
Gebhardt et al., 2021	10.1111/tbed.14335	ASFV	0.70	142
Leme et al., 2019	10.1111/tbed.13215	SVA	25.93	27
Wu et al., 2021	10.1111/tbed.14209	PEDV	14.29	77
<b>Environmental sample summary:</b>		<b>Feed sample summary:</b>		
Minimum prevalence: 0.73%		Minimum prevalence: 0.70%		
Average prevalence: 21.48%		Average prevalence: 13.64%		
Maximum prevalence: 66.24%		Maximum prevalence: 25.93%		

<sup>1</sup>Abbreviations defined as: African swine fever virus (ASFV), Porcine deltacoronavirus (PDCoV), Porcine epidemic diarrhea virus (PEDV), Seneca valley virus (SVA)

# Transitioning from Surveillance Sampling to Suspected Contamination Sampling

When surveillance sampling results meet or exceed pre-established thresholds, it's time to transition into suspected contamination sampling. The purpose of suspected contamination sampling is to identify areas within the feed mill or feed delivery that are contributing to the increased prevalence of the pathogen of interest. Feed mills can also implement mitigation strategies while undergoing suspected contamination sampling to gauge how successful these mitigation strategies are for the pathogen of interest. This resource focuses on how to transition from surveillance to suspected contamination sampling through changes in sample size and sampling frequency. If there are questions regarding sample size and thresholds, refer to the additional resource titled "Calculating Sample Sizes and Thresholds." If there are questions regarding areas of focus, refer to the additional resource titled "Interpreting Sample Results."

## *Changes in sampling size*

Surveillance sample size is based on the probability of feed serving as a source of the pathogen of interest and the severity of the pathogen of interest. However in the case of suspected contamination, the thresholds have been met or exceeded, indicating that the pathogen of interest may be present more frequently or greater than originally perceived. Therefore, to maximize the ability to detect the pathogen of interest, increase the sample size, or decrease the estimated prevalence rate. During suspected contamination sampling, sample sizes might be larger than the sample sizes commonly used for surveillance sampling. To accommodate the larger sampling size, adjust the threshold, or in this instance, the amount of samples that need to be negative in order to return back to surveillance sampling.

## *Changes in sampling frequency*

Suspected contamination samplings need to occur more frequently than surveillance sampling if surveillance thresholds are met or exceeded. For example, the pre-determined surveillance sample sizes are based on monthly prevalence rates and divided across weeks to make sample taking more manageable. However, when thresholds are met or exceeded in surveillance sampling that is indicative that the set prevalence rates for a monthly basis have already been met or exceeded in a week. Therefore, to understand the source of suspected contamination, for the next week, the feed mill will transition to suspected contamination. The feed mill will transition back to the normal surveillance schedule if suspected contamination thresholds are not met or exceeded. If a production system or feed mill chooses to sample more or less frequently than suspected contamination sampling will need to be adjusted accordingly.

# Type of Samples

There are two types of samples to take in a feed mill: feed or environmental samples. This resource will help explain the most current and scientifically backed methodology for sampling complete feed, feed ingredients, and feed mill environments. To access information on how to take feed or environmental samples, please refer to the standard operating procedures titled “Collecting Feed Samples” and “Collecting Environmental Samples.” If requiring information on how to prepare for sampling, please refer to the standard operating procedure titled “Preparing for Sampling of Viral Pathogens.”

## Feed samples

Sampling feed intended for livestock species can offer a way to assess potential contamination in either complete feed or feed ingredients. However, sampling feed is challenging since potential contamination may not be evenly distributed within the feed or ingredient, sometimes referred to as “hot spots” of contamination. To account for this type of distribution, the Association of American Feed Control Officials (AAFCO) feed inspector’s manual offers different solutions on how to sample feed: utilizing sleeved feed probes or single tube triers or sampling via cut stream (AAFCO, 2020). Sampling with sleeved feed probes has been the only methodology to be validated for viral pathogens while the other methodologies have yet to be validated (Jones et al., 2020; Elijah et al., 2021, Dee et al., 2022). Sampling via cut stream could be a solution if the sampling person can’t get to a location to look down into the storage or transport container like what is done with sleeved feed probes. While sampling with single tube triers offers a solution if interested in sampling bagged feed ingredients or complete feed. Table 1 offers a summary of the three methodologies for feed sampling.

All methodologies rely on collecting 10 subsamples per load or lot of complete feed or feed ingredients and combining the 10 subsamples for a single composite sample for submission. The AAFCO feed inspector’s manual recommends a minimum of 10 subsamples so that the sampling methodology can account for any potential of unevenly distributed contamination. If a feed mill is trying to identify potential contamination within a specific batch of feed, taking 10 subsamples within a single load of feed answers the question of potential contamination before delivery. However, if a feed mill is busier than normal, like during times of harvest, collecting 10 subsamples per truck load can be challenging. In this case, if the feed mill is interested in potential contamination throughout the day, each load of bulk ingredient could be considered a subsample, one subsample pulled from each load, and then 10 subsamples from 10 loads could be combined as a composite sample for the bulk ingredients received that day. Depending on the question, the minimum of 10 subsamples can be manipulated to account for different sampling scenarios.

**Table 1.** Methodologies for sampling feed ingredients or complete feed.

Methodology	Used When?	Materials Needed	Minimum number of sub-samples <sup>1</sup>
Sampling with Sleeved Probes <sup>2</sup>	<ul style="list-style-type: none"> <li>• Can get an overview of the sampling container.</li> <li>• Sample container deep enough for the double tube feed probe.</li> <li>• Used for bulk feed ingredients or complete feed.</li> </ul>	<ul style="list-style-type: none"> <li>• Sleeved feed probe</li> <li>• Plastic storage bag</li> <li>• Permanent marker</li> <li>• Disinfectant wipes</li> </ul>	10
Sampling via Cut Stream <sup>3</sup>	<ul style="list-style-type: none"> <li>• If unable to sample feed with sleeved feed probes.</li> <li>• Used for bulk feed ingredients or complete feed.</li> </ul>	<ul style="list-style-type: none"> <li>• 8 ounce cup</li> <li>• Plastic storage bag</li> <li>• Permanent marker</li> <li>• Disinfectant wipes</li> </ul>	10
Sampling with Single Tube Trier <sup>4</sup>	<ul style="list-style-type: none"> <li>• Used for bagged feed ingredients or complete feed.</li> </ul>	<ul style="list-style-type: none"> <li>• Single tube trier</li> <li>• Plastic storage bag</li> <li>• Permanent marker</li> <li>• Disinfectant wipes</li> </ul>	10

<sup>1</sup>Sub-samples refers to the number of samples, or pulls, from the intended sample container that will go into the composite sample.

<sup>2</sup>Sleeved feed probes have an internal and external compartment. Insert the sleeved probe with compartments closed, open compartments once probe is inserted into the feed ingredient or complete feed, shake the probe to fill, close the probe, then withdraw from feed ingredient or complete feed.

<sup>3</sup>Cut stream is the terminology used to describe when sampling relies on a stream of feed ingredients or complete feed and the sampling container passes through the stream and fills the sampling material to obtain a sub-sample.

<sup>4</sup>Single tube trier has an open sampling compartment with a handle. Single tube triers are rotated so sampling material is collected into the open compartment.

## Environmental Samples

Sampling the environment of the feed mill can offer a way to understand the directionality, or spread, of pathogens of interest or monitor the biosecurity practices in place. To help with this, environmental samples are classified into zones based on the surface and what that surface comes into contact with. Environmental samples can be classified into the following zones:

- Feed contact zones: these surfaces have direct contact with feed ingredients or complete feed.
- Non-feed contact zones: these surfaces have a fixed location and are close or next to feed contact zones.
- Transient zones: these surfaces do not have a fixed location and can move within the feed mill environment or feed delivery.

An example of these surfaces within the zones can be found in the “Sampling Locations” additional resource. Understanding which zone each sample was taken from can help guide strategies on how to reduce potential contamination.

Based on the pathogen of interest, there are methodologies that have been shown to maximize the potential of finding the pathogen on feed mill surfaces. Table 2 offers a summary of the methodologies based on pathogen.

**Table 2.** Methodologies for environmental sampling based on pathogen.

Pathogen	Methodology <sup>1</sup>	Sampling Material <sup>2</sup>	Pre-Moistening Solution <sup>3</sup>	Size of Sampling Area	Number of Passes of Sampling Area <sup>4</sup>
Bacteria	Hand Sampling	3M Sponge Sticks	Buffered Peptone Water	8 × 8 in.	10 horizontal pushes and pulls 10 vertical pushes and pulls
Virus	Hand Sampling	4 × 4 in. Cotton Gauze	Phosphate Buffered Solution 0.9% NaCl Sterile Saline	8 × 8 in.	10 horizontal pushes and pulls 10 vertical pushes and pulls
Virus	Extension Set Sampling	Synthetic Paint Roller Cover	Phosphate Buffered Solution 0.9% NaCl Sterile Saline	--	10 horizontal pushes and pulls 10 vertical pushes and pulls

<sup>1</sup>Methods of collecting environmental samples can rely on hand sampling or usage of an extension set to sample hard to reach areas for viral pathogens. The surface of interest will determine which methodology will work best. At this time, hand sampling is the only method for bacterial pathogens.

<sup>2</sup>Sampling material refers to the material that will pass over the surface of interest. The 3M sponge sticks have been shown to be most effective for bacterial pathogens (Moore and Griffith, 2002; FDA, 2021) while cotton gauze is the most effective for viral pathogens (Stewart et al., 2019). Synthetic paint roller covers is the material of choice and shown to work the best (Wu et al., 2021; Elijah et al., 2022) but if unable to acquire synthetic paint roller covers, cotton paint roller covers are an acceptable substitute given the data to support cotton as a suitable material for viral pathogens.

<sup>3</sup>Pre-moistening solution refers to the solution that moistens the material before sampling. By pre-moistening the material, the ability of the sample material to pick up potential pathogen is maximized (Moore and Griffith, 2002). For gram negative bacteria, buffered peptone water is the pre-moistening solution of choice. For viruses, phosphate buffered solution (1X concentration, pH=7.4) is the pre-moistening solution of choice but recent research has shown that 0.9% NaCl sterile saline is an acceptable solution if unable to acquire phosphate buffered solution (Rodino et al., 2020).

<sup>4</sup>Number of passes refers to the number of times the sampling material should pass over the sampling area to pick up the pathogen of interest.

## References

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# Sampling Locations

Collecting environmental samples is a way to proactively monitor for pathogens of interest within a feed mill and surfaces associated with feed delivery. Previous research has demonstrated that objects or people involved feed mill or feed delivery contribute to the spread of pathogens like African swine fever virus, porcine deltacoronavirus, or porcine epidemic diarrhea virus (Gebhardt et al., 2021; Elijah et al., 2022). There are a multitude of surfaces that could be sampled within a feed mill so it can be overwhelming to decide where to focus sampling efforts. Therefore, this factsheet aims to provide a list of surfaces based on zone that have been shown to potentially harbor pathogen of interest.

## Feed Contact Surfaces

Feed contact surfaces have direct contact with feed ingredients or complete feed. These surfaces are associated with feed manufacturing, storage, and delivery. If these surfaces are positive for pathogen of interest, its origin may have been from a contaminated feed ingredient.

Surfaces associated with the feed mill	Surfaces associated with feed delivery
Corn cleaner	Interior of feed truck compartments
Receiving pit grates	Interior of feed truck boom
Fat intake inlet or hose	
Interiors of feed bins	
Load out auger or sock	
Interiors of bucket elevators	

## Non-Feed Contact Surfaces

Non-feed contact surfaces are surfaces with a fixed location that are close or next to feed contact surfaces. These surfaces are either covered by dust from feed manufacturing or have a lot of foot traffic. If these surfaces are positive for pathogen of interest, its origin may have been through a contaminated feed ingredient dust generated during feed manufacturing or transient surfaces contributing to the spread of the contamination.

Surfaces associated with the feed mill	Surfaces associated with feed delivery
Floor of load out bay	Exterior of feed truck compartments
Control room floor	Exterior of feed truck boom
Floor mat by main entrance	
Receiving floors	
Manufacturing floors <ul style="list-style-type: none"> <li>Floors around hand add ports</li> <li>Areas near sample ports</li> </ul>	
Warehouse floors	
Exterior of pellet mill	
Pellet mill air intake	
Inside dust collection system	

## **Transient Surfaces**

Transient surfaces are surfaces with a non-fixed location and can move within the feed mill or during feed delivery. These surfaces have intermittent contact with other surfaces that could potentially have exposure to pathogen of interest. If these surfaces are positive for pathogen of interest, its origin may have been through employees introducing or spreading the contamination.

<b>Surfaces associated with the feed mill</b>	<b>Surfaces associated with feed delivery</b>
Fork lift tires	Feed truck steps
Broom	Feed truck floor mat and pedals
Shovels	Feed truck tires
Worker shoes	Workers shoes
Worker clothing	Workers clothing

## **References**

- Elijah CG, Harrison OL, Blomme AK, Woodworth JC, Jones CK, Paulk CB, and Gebhardt JT. (2022). Understanding the role of feed manufacturing and delivery within a series of porcine deltacoronavirus investigations. *J Swine Health Prod.* 30(1):17-23.
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# Diagnostic Tests for Samples

Once samples have been collected, the samples should be sent to a veterinary diagnostic laboratory for viral or bacterial analysis. Veterinary diagnostic laboratories can take a variety of sample types and have the personnel trained to appropriately handle the samples for analysis. A list of available accredited veterinary diagnostic laboratories can be found [here](#). While the veterinary diagnostic laboratories have a wide variety of tests available, this resource will focus on the tests, assays, or processes that are typically conducted on environmental or feed samples.

## Bacterial Pathogens

Before running any tests, instruct the diagnostic laboratory to enrich the sample. The purpose of the enrichment step is to grow the contamination present in the samples to detectable levels. Enriching samples increases the sensitivity of any intended tests to be run on the bacterial samples. If electing to run tests without enrichment, understand that there is the potential for the diagnostic tests to return as false negative since the contamination might not be present at detectable levels.

After enrichment, samples can be submitted to either bacterial culture or PCR analysis. Bacterial culture is considered the less expensive option but tends to be more time consuming and can identify genus of bacteria but not specific types of bacteria. While PCR analysis has the ability to identify specific bacterial genus and species, the sample will still have to be cultured to increase the sensitivity of the PCR assay. An example of when a production system or feed mill would elect PCR analysis over bacterial culture is if the production system or feed mill is concerned about *Salmonella* in the feed intended for livestock consumption. If concerned about *Salmonella* in feed for livestock species, consult the Food and Drug Administration guide on *Salmonella* in food for animals (FDA, 2013).

Another potential option for bacterial analysis is the use of near infrared spectroscopy (NIR). This is a relatively new technology to detect bacterial contamination within feed that offers a quick turnaround time and is less expensive than PCR (Tian et al., 2021). It offers the same results as PCR but since this is a more recently developed technology, the sample will still require bacterial culture and in some instances, NIR might be less precise than PCR assay (Tian et al., 2021). Consult with the diagnostic laboratory at time of sample submission if this is a test that they are offering and would recommend.

## Viral Pathogens

Before running any tests, request that the diagnostic laboratory centrifuge all samples. Research suggests that centrifugation of samples before laboratory analysis increases the sensitivity of PCR (Elijah et al., 2021). Request the samples to be centrifuged at 4000 × g for 10 minutes (Khanal et al., 2022). If desiring to centrifuge environmental samples before submission, refer to the standard operating procedure titled “Centrifugation of Environmental Samples for Viral Pathogens” and notify the veterinary diagnostic laboratory that the environmental samples were centrifuged prior to submission.

Some diagnostic laboratories can offer multiplex PCR assays that will look for multiple viruses within the sample at once. Generally, multi-plex assays are for enteric viral pathogens since it is difficult to distinguish the different enteric viruses based on clinical signs. If a production system or feed mill is concerned about enteric viral pathogens, then a multi-plex PCR assay offers the ability to test for multiple viral pathogens at once. Diagnostic laboratories also have single-plex PCR assays which look for one pathogen within the sample. There are a wide variety of options for evaluating for viral pathogens with single-plex PCR assays. However, single-plex PCR assays can be more

expensive if concerned about two different viral pathogens since the samples will have to be run twice for different tests.

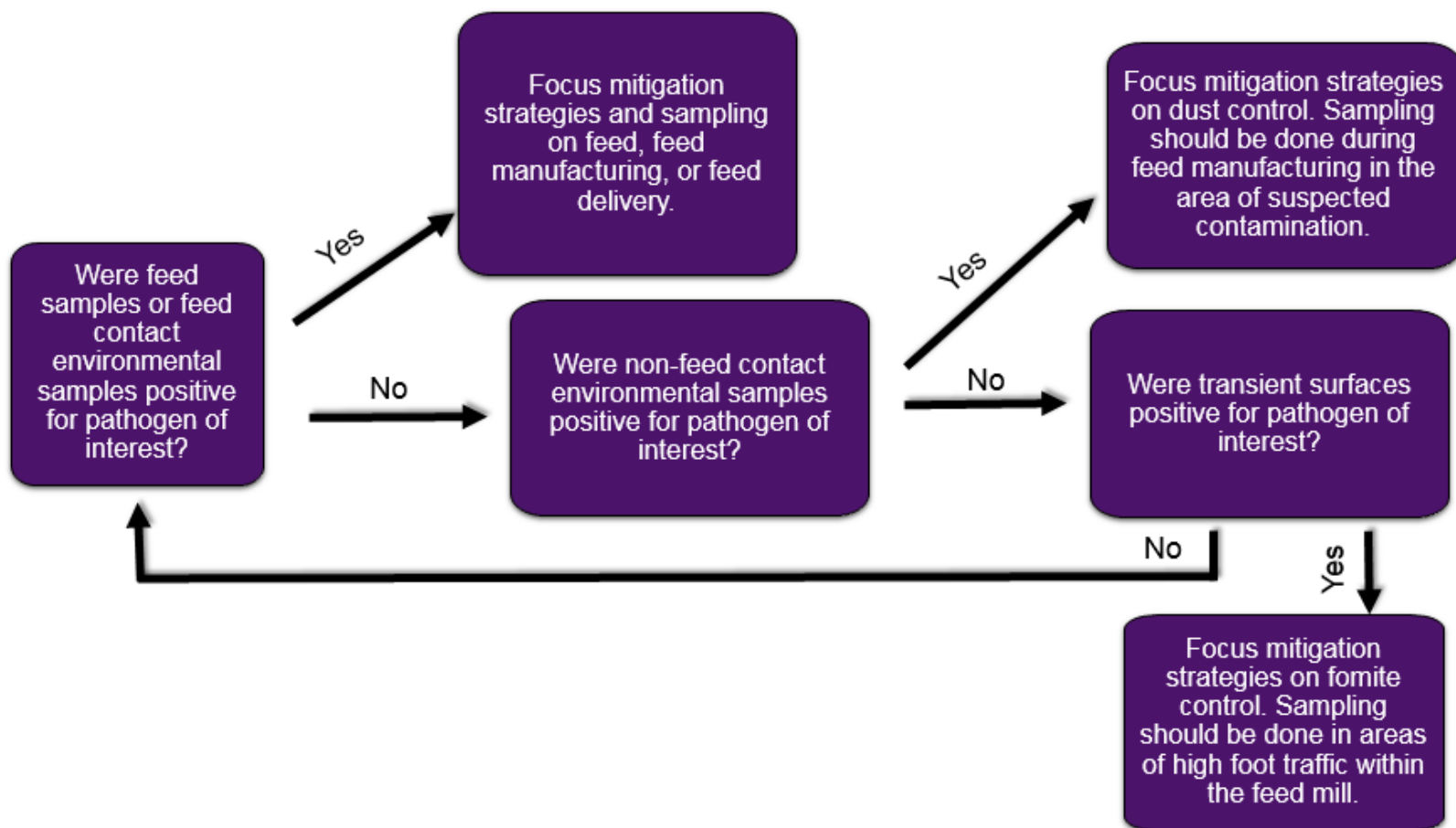
Once deciding on the PCR assay for samples, ask the diagnostic laboratory to run the PCR assay to maximum number of cycles of 45. By running the PCR assay to the maximum number of cycles, this practice increases the diagnostic sensitivity, or the ability to detect positive samples. However, by increasing the diagnostic sensitivity, this increases the risk of false positives thereby reducing the diagnostic specificity or the ability to detect negative samples. By running to the maximal number of cycles, the assay repeatability is also reduced. An example of when to run to the maximal number of cycles would be when the objective of the sampling is to identify lapses in biosecurity and want to maximize the sensitivity of the assay to detect the pathogen. While an example of when to utilize the lower cut off value would be when a feed mill is conducting confirmatory testing to prove pathogen presence.

## References

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# Interpreting Sample Results

After samples have been collected, submitted, and analyzed, communicating sample results to the appropriate groups is essential. Interpretation of the samples will guide future mitigation techniques but also areas of focus for sampling since these areas may be challenging to maintain or remain contamination free. Consult the flow chart below to understand how to interpret the sample results from surveillance or suspected contamination sampling.



# Strategies to Reduce Contamination

When samples from surveillance or suspected disease contamination come back as positive for the pathogen of interest, there are strategies that can be implemented to reduce contamination. These strategies can be implemented at any time that a feed mill or production system is concerned about potential pathogen contamination. This resource will discuss potential risk mitigation techniques and examples of these techniques.

- **Limiting entry of potential pathogens into the receiving pit.**

- Utilizing receiving pit covers
  - Constructing a pit cover that lays flat while not receiving bulk ingredients but can be lifted up and constructed into a funnel shape to help reduce shrink during unloading of bulk feed ingredients.
  - Covering the receiving pit with a rubber mat, or something similar, when not in use.
- Discarding spilled feed into the trash instead of adding it back into the receiving pit.

- **Chemical feed additives**

- Addition of chemical feed additives to feed intended for livestock has been shown to potentially decrease the risk of cross-contamination during feed manufacturing or feed delivery.
  - Common chemical additives include organic acids, formaldehyde, essential oils, medium chain fatty acids, or dietary acidifiers (Huss et al., 2018).
  - More information on chemical feed additives can be found [here](#).

- **Implementing point-in-time mitigation techniques.**

- Point-in-time mitigation techniques are strategies implemented during a time point of feed manufacturing. These techniques do not prevent the possibility that feed may become contaminated again during further feed manufacturing or delivery.
  - Thermal Processing
    - Addition of heat to the feed manufacturing process to reduce potential infectivity of the pathogen of interest (Huss et al., 2018).
    - For livestock feed, pelleting is considered the traditional method of thermal processing.
  - Feed Batch Sequencing
    - Requires the order of production, storage, and distribution to be planned to reduce the carryover of high-risk ingredients to sensitive diets (Huss et al., 2018).
  - Flushing
    - Consists of running an ingredient, usually with abrasive material, through the system between batches to flush out any residual material (Huss et al., 2018).
  - Holding or quarantining feed ingredients
    - Consists of storing ingredients in a low foot traffic areas for a specified amount of time between manufacture and use to give an opportunity for viral contamination to naturally degrade so as not to be infectious.
    - More information on how to calculate holding times can be found [here](#).

- **Zoning**

- Restricting employees to certain locations within a feed mill to limit the spread of pathogen

- **Implementing downtime**

- Requiring a specified amount of time before employees return to the feed mill that have had recent contact with animals outside of the feed mill or when making deliveries to certain production sites.

- **Limiting the amount of contaminated objects**

- Requiring shoe covers for feed truck drivers and ensuring they wear and change them during deliveries.
- Providing feed mill specific uniforms.
- Scheduling or restricting deliveries to certain production sites on certain days.
- Requiring showers before entering or exiting the feed mill.

- **Implementing usage of disinfectants**

- Liquid or dry boot baths at the entrances or exits into the feed mill.
- Applying disinfectants to semi-truck cabs after deliveries.
- Combining disinfectant application with heat treatment like baking trailers after power washing with disinfectant.
- More information regarding disinfectants can be found [here](#).

## **References**

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