

final report

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Process Risk Models – Continued Development 13/14 and 14/15

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Background

Previously a process risk model was developed to utilise existing data from MLA projects and the wider literature, and place it into a risk context which could be used as a research tool to better understand risks and identify areas within the program (particularly the pathogen and microbial contamination area) requiring further investigation. The model allows for analysis of data in a descriptive and mathematical manner, and is useful within the pathogen program plan as a predictive tool to ensure MLA and the industry stays "ahead of the play" rather than just "reacting/responding" to food safety concerns. Modelling was also used to understand contamination of cartons of manufacturing beef.

The maintenance and further development of the existing risk model is important to MLA for a number of reasons. Firstly, using data collected for a pathogen known to currently pose a food safety problem, the model can be used to predict prevalence and concentration of those pathogens which are foreseen to cause problems, but for which, little data exists. In addition, the model can be used to identify particular steps throughout the processing chain that present significant risk, thus providing direction as to what areas require further investigation and data collection. Carton beef models can be used to support Australia's testing and control of pathogens such as *E. coli* O157

Project Objectives

The maintenance of the Process Risk Model will involve:

- 1. Documenting and explaining the model to maintain transparency and accessibility to MLA and MLA's scientific risk management panel
- 2. Identifying parts of the existing model which may need improvement / updating
- 3. Identifying areas within existing data, where there may be incomplete data, and a need for additional collection
- 4. Specifying the data requirements and allow for data obtained from a wide range of different projects within the program to be fed into the model for evaluation
- 5. Contributing to the development of experimental and survey design for projects related to the model
- 6. Identifying areas within the processing chain which may be more important from a risk viewpoint, and therefore require a greater degree of investigation / knowledge
- 7. Assisting in the development of recommendations for complete risk assessments, and performing risk assessment, when required
- 8. Assisting in the development of risk management options, based on outcomes from the use of the process risk model
- 9. Interacting with MLA's scientific risk management panel, as required.

Results and Discussion

The following is a summary of the work undertaken as part of this project between the period of 1 July 2013 and 30 September 2013.

Risk assessment of E. coli O157 in burgers made from Australian beef trim

A risk assessment for *E. coli* O157 in burgers made from Australian beef trim, which was initiated in project A.MFS.0222, has been further developed and documented. This risk assessment uses data on the contamination of cartons of Australian beef trim¹ and subsequently models the *E. coli* O157 concentration in beef patties through retail storage, transport to the home, home storage, cooking and consumption. A manuscript reporting the risk assessment and results is still being prepared for scientific publication in *Risk Analysis*.

The key results from this risk assessment have been presented at the Annual Conference of the International Association for Food Protection in Charlotte, North Carolina from 28-31 July 2013 (Attachment 1).

Abattoir visits with Sam Rogers

As part of MLA project G.MFS.0294 "Statistical Process Control – Hygiene and Hazards" Dr Kiermeier visited five abattoirs as part of Sam Rogers' industry familiarisation. Abattoirs included the Northern Co-operative Meat Company, Oakey Abattoir, Greenmountain Meats and JBS Australia (Dinmore) on 29 & 30 August 2013, and Thomas Foods International on 27 September 2013.

MINTRAC QA Manager's Network meeting

Dr Kiermeier delivered an MLA presentation entitled "*Does your performance measure up?*" at the Adelaide MINTRAC QA Manager's Network meeting on 21 August 2013 (Attachment 2).

MINTRAC Conference

Dr Kiermeier contributed to the development and in the delivery of an MLA presentation entitled *"Increasing your certainty of meeting market requirements"* at the MINTRAC 2013 Meat Inspection and Quality Assurance Conference in Melbourne on 11 & 12 September 2013 (Attachment 3).

Discussion document on aspects related to sampling beef trim for E. coli O157

Exporters of manufacturing beef undertake *E. coli* O157 testing of each lot using the 'Robust N-60' sampling plan. An argument which is frequently made in the USA is that testing smaller lots results in higher probabilities of detecting contamination. Subsequently, there has been pressure on establishments to reduce the lot size from a maximum of 700 cartons to 350 or even 175 cartons. A document to address some of the issues related to *E. coli* O157 testing has previously been produced and this document has been revised following discussions with Dr Mansour Samapour at the Annual Conference of the International Association for Food Protection in Charlotte in July 2013. The most recent version of this document is included with this report (Attachment 4).

¹ Kiermeier, A., Mellor, G., Barlow, R. & Jenson, I. (2011). Assumptions of Acceptance Sampling and the Implications for Lot Contamination: *Escherichia coli* O157 in Lots of Australian Manufacturing Beef. *Journal of Food Protection* 74(4): 539-544.

Conclusions and Recommendations

This project has provided MLA with a flexible mechanism to address a variety of statistical and modelling issues. As of 4 October 2013, Dr Andreas Kiermeier will no longer be working for SARDI. Because of this loss of key scientific capability to deliver this research project, it is recommended that this project is terminated.

A. Kiermeier (SARDI) J. Sumner (MLA) I. Jenson (MLA)

Risk Assessment of *E. coli* O157 in Hamburgers Made from Australian Beef Trim



MEAT & LIVESTOCK AUSTRALIA

Background

- Australia exports large amounts of lean beef destined for grinding to the US
- All lots are tested for *E. coli* O157 using N-60: Detections are infrequent
- Microbial status of Australian beef has been well documented
 - 4 baseline studies

Aim

To assess the risk posed by *E. coli* O157 from the consumption of burgers made only from Australian beef trim and to compare the effects of sampling and cooking interventions on illness estimates.





Simulation

- Based on export sampling *E.coli* O157 not found frequently
 - Estimate that only about 0.5% of AU lots are contaminated
- Simulate only contaminated lots (10,000)
 - 700 cartons per lot each carton 27.2kg
 - 190,400 hamburger patties (100g) per lot

Carton Contamination

- Five contaminated lots (A-E)
- Identified through standard N-60 sampling
- The 12 original cartons were sampled again
- 75 surface slice per carton 900 samples per lot – tested individually for *E. coli* O157

Carton Contamination

- Lots A, B, C: could not isolate *E. coli* O157
 Assume one carton was contaminated
 - Estimate <0.0013 org/cm²
- Lot D: 1 piece of meat from 1 carton
 Estimate 0.0014 org/cm²
- Lot E: 27 pieces of meat from 2 cartons
 Estimate 0.019 and 0.093 org/cm²

Grinding and Patty Forming

- Individual cartons
- No cross contamination between cartons
- Organisms in a carton end up in patties

Storage & transport

- Quick service restaurants
 Stored frozen ⇒ No growth
- Retail, transport to home and home storage
 - Based on FSIS 2001 RA approach
 - Using updated EcoSure (2008) data

Cooking & Consumption

- Quick Service Restaurant
 - Cooked to internal temperature of constant 68°C (154.4 °F)
- Home
 - Variable internal endpoint temperature based EcoSure (2008) data
- Cassin's Dose-Response model



Results – All AU lots

- Baseline no testing undertaken (!)
- Take non-contaminated lots into account
- Through the home variable cooking: 3.0 illnesses per 10,000,000 hamburgers
- Through QSR cooking to 68°C (154.4 °F)
 7.3 illnesses per 100,000,000,000 hamburgers

Results – All Australian lots

Scenario	Rate of illness per burgers consumed
Home cooking (variable temp.)	3.0 per 10,000,000
Sampling N6o + home cooking	1.5 per 10,000,000
Sampling N120 + home cooking	1.1 per 10,000,000

Results

Location	Patties consumed	Illnesses estimated
Home	232,500,000	34.10
QSR	2,085,750,000	0.07
Total	2,425,000,000	34.17

Conclusions

- Sampling removes the most contaminated lots
- Less contaminated lots are harder to detect, but they also cause less illness
- Law of diminishing returns

Conclusions

• Cooking is (still) the best way to reduce risk



Conclusion

Australia's contribution

- About 10% of raw materials for hamburger production in the US
- About 0.2% of illnesses



Thank You!





























Establishments 1, 2 and 3						
E) ca are	xample of some ind arcase immediately ea	creases - contamin after hide remova	ation from hide to I – sponging a large			
		Mean Log TVC (cfu/sample) [~ cfu/cm ₂]	Prevalence E. coli			
	Est 1	4.20 [1.6]	82% positive			
	Est 2	3.75 [1.15]	40% positive			
	Est 3	4.39 [1.79]	27% positive			







						mla
We start with a sterile carcase under the hide	+ c fr (i	ium of Contamination rom various places increases)	-	Sum of treatment to remove contaminatior (removal)		The performance objective that contributes to neeting the Food Safety Objective
H_{o}	+	ΣI	-	∑R	≤	PO (FSO)
0 sterile	+	1.6 carcase 0.2 RI	-	0.5 trim	\leq	2.0
0	+	1.8	-	0.5	≤	2.0





- Australian meat industry/MLA since 1990's
- Predictive microbiology/risk assessment, also expert panels
- Given us tools which are routinely used
 - Refrigeration Index
 - EHECs in fermented meats
 - Shelf life/spoilage of VP meats



They've done this for E. coli.















mla

Increasing your certainty of meeting market requirements

Andreas Kiermeier, Casey Smith, Ian Jenson, John Sumner, Mandy Smart, Sam Rogers, Teresa Hore, SARDI E C Throsby MLA MLA Midfield International SARDI Southern Meats





mla

Need to deal with these requirements in a consistent way

- No control over requirements
- · We have to meet them
- We have to manage our systems
- We need to be confident that our product meets requirements
- Best if we can use a common approach to thinking about how we meet requirements

Traditional process control

Outline

- New approaches to meeting requirements
- Investigating your operations





MENT & LONGTOCK AUSTRALIA	Plant A: MHA history
sut	Plant A: MITA history
Data analysis indicates that pathogens can occur even though our existing systems indicate that things are 'in control'	























G.MFS.0314 Final Report - Attachment 3





Three examples	
 Tail flick at hide pulling – Casey Smith Throsby 	, EC
 Dropping socks on smallstock line – T Hore, Southern Meats 	eresa
 Cleaning chillers using ozone – Mandy Midfield International 	y Smart,









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Objective

If we drop socks with an airknife how will contamination compare with current procedure using a conventional knife?

If it does we can save an operator.









тус			E. coli	T	
Summary	Knife (log) Air	knife (log)	Summary	EC (Knife)	EC (Air Knife)
Mean	1.70	1.72	Detect	3	2
St. Dev.	0.45	0.43	n	25	25
n	25	25	Prev	12.0%	8.0%
Conf level	95%		Conf level	g)5%
CI Lower	1.51	1.54	CI Lower	3.5%	1.2%
CI Upper	1.89	1.90	CI Upper	31.0%	26.3%
Significance	Not significa	ant	Significance	Not si	gnificant



Using ozone to clean chiller walls ma

Objective

Determine if ozonation of a chiller over a 2h period will result in lower levels of TVC on the wall

Theory

Ozone is a powerful oxidizing agent which by the action of UV light and atmospheric electrical discharge has inhibitory microbiological effects

Experimental

A grid pattern was taped to the chiller wall. Twenty-five sites were sampled using a press plate. After ozonation, sites adjacent to the original 25 were sampled.













STEC Sampling

This document discusses sampling and testing using an enrichment test. While we have used testing for *E. coli* O157 as the example, the approach applies equally to the big 6 STEC, or any other bacteria tested in a similar way.

Background

In 2007, FSIS commenced testing of beef trim destined for grinding using surface slice and "Robust N-60" sampling¹ at USA processing establishments. Subsequently, Australian establishments were required to also sample and test all lots of beef trim destined for the USA using an N-60 protocol (Australian Meat Notice 2007/17). The meat notice stipulated a maximum lot size of 700 cartons (a container equivalent) and that a lot would consist of product packed on a given packing line and based on Sanitation SOPs and/or determined by the establishment, based on the implementation of a statistically based sampling program to distinguish between segments of production. For each lot, five 5-10g samples – surface slices or small grab samples – were to be collected from a minimum of 12 randomly selected cartons, to a total sample weight of at least 375g.

Initially, sampling was undertaken on cartons of fresh meat prior to carton sealing and freezing, though collection of frozen samples was an option. However, it wasn't until implementation of Australian Meat Notice 2008/9 that testing of frozen samples became more established throughout the industry, with many establishments testing at load-out or shortly before. This was possible because there was no longer a requirement to define lots through "Sanitation SOPs" but only through Robust N-60 sampling and testing and hence could be confined to a single container load (or less). That is, lots that were tested separately could be deemed independent and a detection of *E. coli* O157 in one lot did not trigger a rejection of other lots provided they had been tested separately – even if they had been produced during the same production period.

In comparison to the USA, Australian lot sizes were large – up to 20t (700 cartons or a container load) compared with 5 combo bin lots (total of 5t). The trend in the USA has been toward smaller lots and 1 combo bin lots (2000lb or 1t) are now common. In contrast, Australian lot sizes remained largely unchanged, until recently. Some establishments have now reduced lot size and decided to have two port marks (sometimes even four) per container and have tested each of these port marks as separate lots. Reducing lot size has become more popular since the introduction of the requirement for DAFF verification testing for six additional Shiga toxigenic *E. coli* (STECs), in addition to *E. coli* 0157, in June 2012 (Australian Meat Notice 2012/01). In addition, information obtained from the USA has prompted some establishments to believe that testing an increased number of port

¹ FSIS (2011) National Prevalence Estimate of Pathogens in Domestic Beef Manufacturing Trimmings (Trim): December 2005 – January 2007,

http://www.fsis.usda.gov/PDF/Baseline_Data_Domestic_Beef_Trimmings_Rev.pdf

marks/containers will result in a greater chance of detecting contamination in Australia and a reduced likelihood that a lot will test positive at Port of Entry.

Unfortunately, some of the advice provided from the USA is not appropriate for the Australian situation and this document has been prepared to provide information on how lot size and testing frequency impact on sampling and likely sampling results.

The situation in the USA

Let's make an assumption that an establishment typically produces lots in which 1% of meat pieces are contaminated with *E. coli* O157. In this case, randomly collecting 60 samples, i.e. robust N-60 sampling, would result in a probability of detecting *E. coli* O157 of 45.3% (see Appendix).² That is, on average about half of all lots contaminated at this level (1%) would have O157 detected, while in the other half the contamination would not be detected and these lots would progress through to commerce.

Information from the USA indicates that processors create lots by combining several combo bins, possibly from different days of production. These lots are then tested by selecting an equal number of samples from each combo bin, while maintaining the overall N-60 sampling plan (Table 1). For example, if a lot contains four combo bins, then each bin would be sampled by selecting 60/4 = 15 pieces of meat. Based on an average 1% contamination rate (and assumptions detailed in the Appendix), the probability of each one of these combo bins yielding a detection is 14.0%. Because all 60 meat samples are pooled into one analytical sample, which is enriched, the whole lot of four combo bins is rejected when O157 is detected (even if the contamination originates from just from one bin). As it turns out, testing N-60 drawn from all four combo bins is equivalent, in terms of probabilities, to testing each combo bin separately using n=15 and rejecting the whole lot when only one of them results in an O157 detection. The fact that all four combo bins are rejected is important as it results in an overall probability of 45.3%³ rejecting for the whole lot – the same probability we calculated above using a pooled sample and n=60.

	AU sampling	US sampling
Lot units	Carton (27.2 kg)	Combo Bin (1t)
Predominant lot size	175/350/700 cartons	1-5 combo bins
Number of samples	N-60	N-60
Samples are collected	5 samples (5-10g) from	Equal number from each combo bin
how?	each of 12 randomly	to give N-60, e.g. 15 samples (5-10g)
	selected cartons	from each of 4 combo bins in lot

Table 1: Summary of Australian and USA sampling schemes for E. coli O157.

As indicated above, it is possible that the four combo bins originate from different days of production. Let's assume that one of these days was a "High Event Period", that is, a day

² Throughout this document we use the approach and statistical assumptions detailed in the Appendix for the calculation of probabilities.

 $^{^{3}}$ 1-(1-0.14)⁴ = 0.453 = 45.3%

with worse than average O157 contamination, and that the combo bin from that day has 4% of pieces contaminated. Let's also assume that the other three combo bins have no contamination (0%) as this gives us an average contamination rate of 1% (same as above). The probability of rejecting this lot remains at 45.3%, despite three combo bins not being contaminated (they have a 0% probability of rejection). However, the highly contaminated combo bin has a 45.3% probability of detection, although it is only sampled n=15. And this is where the concern in the USA comes from, namely that (a) lots are created from combo bins from unrelated days of production, and that (b) because of this, and the lower intensity of sampling (n=15), a highly contaminated combo bin has 'only' got a 45.3% probability of rejection. In contrast, if each combo bin was tested as a separate lot using n=60, then the heavily contaminated combo bin would have a 91.4% probability of detection while the remaining three uncontaminated combos would be cleared for commerce.

The important part to note here is that the change in probabilities – from 45.3% to 91.4% – is not because the lots size was reduced. Instead, it is a result of the things noted above, namely that

- a) lots are created by combining unrelated days of production;
- b) the contamination rate is higher; and
- c) the more contaminated lots is tested using n=60 rather than n=15.

The situation in Australia

In Australia we have lots of up to 700 cartons, although some processors are moving to smaller lots sizes, e.g. 350 or even 175 cartons. Just like in the USA we use N-60 sampling to test lots for *E. coli* O157 (Table 1). However, in contrast to the USA, we randomly select the cartons from which we collect samples, which is due to the fact that we have many more cartons than we need samples.

So, if a lot is created by taking 175 cartons from each of 4 days of production, then we don't test each day using n=15. Instead the 12 cartons are randomly selected from the 700, and hence any particular day may have any number of cartons between 0 and 12 selected for testing. This is quite different to the USA where each combo bin is tested proportionally.

While the way we collect samples is different the USA argument is frequently *interpreted* or *portrayed* incorrectly. The case seems to be made that halving a lot's size, and hence testing both halves (or port marks) using N-60, will increase the probability of detecting a contaminated lot to $1-(0.45)^2 = 0.80 = 80\%$. Furthermore, if the initial lot is divided into four port marks, then the probability of detection will increase to $1-(0.45)^4 = 0.96 = 96\%$. However, this interpretation about the probability of detecting O157 is incorrect. That's because it implies the probability of detecting O157 in every port mark, while actually it is the *probability of detecting E. coli O157 in <u>at least one</u> of the four port marks*. But each port mark is accepted or rejected independently of other port marks. This is in fact no different to the situation in the USA – if the lot size is reduced from four combo bins to just one, then a detection in one does not trigger rejection of all four.

As described above, each individual port mark still has the same probability of detection as the original lot, namely 45.3% (just like the combo bins in the USA). That's because we are using the same N-60 sampling plan and are collecting a small amount of meat (375g) from a very large lot (700 cartons = 19,040 kg and 175 cartons = 4760 kg). If we used N-240 sampling (for a port mark or a container) then the probability of detecting O157 becomes 0.91=91% when 1% of pieces are contaminated. Again, the important thing is that the probability has increased because the number of samples tested has increased, and not because the size of the lot has changed. However, this N-240 should not be confused with testing four port marks with N-60 – that's because each port mark is accepted / rejected independently.

So, what effect does this have on your sampling and testing program? To answer this question, let's also assume that 1% of pieces are contaminated in the original 700 cartons lots. Based on the above calculations, we would expect that for every two lots tested, we would reject one of these (on average), i.e. a total of 700 cartons.

Now let's divide these two lots into four port marks each, that is, a total of eight port marks are being tested. If the contamination is distributed randomly throughout each lot, then each of these eight port marks will still have 1% of its pieces contaminated. Consequently, we again expect half of these, that is, four port marks, to result in an O157 detection. Therefore, the total amount of product rejected is equal to $4 \times 175 = 700$ cartons.

On the other hand, it could be that all the contaminated pieces end up in a single port mark – this might be the case if a few highly contaminated carcases are boned and packed and the contamination is not spread widely. So in our two containers with a total of eight port marks, two will be contaminated at 4% (because all contaminated pieces are together) and the remaining six port marks will be free of contamination. Using N-60 sampling, the two contaminated port marks have a probability of detection of 91% while the remaining port marks have 0% probability of O157 detection (because they contain no contamination). In this case you would reject a total of only 350 cartons.

But which of these two extremes (or anywhere in between) we are dealing with, is anyone's guess. However, if the port marks originate on different days of production, then given the above information, it may be beneficial to test them separately and independently. It is also important to remember that not detecting the contamination is not the same as "there is no contamination" and that the above scenarios assume true independence between lots and port marks.

Conclusions

There is no statistical justification for just making lots smaller – the probability of detecting O157, and similarly STECs, does not increase simply because the lots are smaller (assuming the same prevalence or level of contamination). Practically, smaller lots may be attractive as a detection will result in less product being condemned. However, all other things being equal, smaller lots require you to test two or four times as frequently and therefore we

would expect to detect STECs two or four times as frequently, but the overall amount of meat rejected is left largely unchanged.

However, combining unrelated cartons, from different days of production, into a single port mark or lot carries some risk, because the whole lot is rejected and not just the independent portions making up the lot.

Finally, don't assume that failing to detect O157 (or STECs) is the same as O157 (or STECs) not being present in the lot.

Mathematical Appendix

You may want to discuss this topic with your customers so this appendix contains some further information for calculating lot acceptance and rejection probabilities.

To calculate the probability of lot acceptance (or rejection) FSIS utilised the commonly used binomial distribution. This approach is usually utilised when the total amount of material that can be sampled is much larger than the actual amount sampled. Under the binomial distribution, there are a number of assumptions made:

- 1. The total number of meat samples collected is fixed in advance. This is clearly the case as we take 60 samples from the lot (irrespective of lot size).
- Each individual meat sample either contains an STEC or does not contain an STEC

 the number of STECs are not important, only the presence / absence. This is also a
 reasonable assumption.

It is also assumed implicitly that any STECs present on a meat sample will be detected (i.e. no false negative results) and that compositing the 60 samples followed by enrichment is a suitable way of detect at least one STEC (and hence just as good as testing each individual piece of meat separately).

- 3. The STEC prevalence and hence the chance of a single piece being contaminated is constant throughout the lot. This assumption is harder to assess and means that there aren't known factors that increase or decrease the prevalence. For example, if the first 350 cartons in a 700 carton lot were irradiated (and hence all STECs were eliminated) then these would not yield STECs if sampled. So there is a known systematic effect that needs to be taken into account and hence these two sets of 350 cartons should be treated as separate lots (and sampled separately). However, in practice such knowledge usually doesn't exist.
- 4. Individual meat samples are independent from each other. This is not the case when samples are obtained from pieces of meat that originate from a single carcase. If parts of the carcase are contaminated with STECs then it may also be more likely to have other parts also contaminated. In many cases, meat from multiple carcases is collected in a single carton and it is not possible to discern which carcase they come from.

Under these assumptions, the probability of detecting STEC can be calculated from the binomial distribution for any given prevalence (p) and number of samples (n) as

P(detect) = P(1 or more STECs in sample)

$$= 1 - P(\text{no STECs in sample})$$

 $= 1 - (1 - p)^n$

For *p*=0.01 (=1%) and *n*=60 we get

$$P(\text{detect}) = 1 - (1 - p)^n = 1 - (1 - 0.01)^{60}$$
$$= 1 - 0.99^{60} = 1 - 0.547 = 0.453$$

The prevalence depends on the size of the sample unit collected. In the case of N-60 sampling the sample unit is 5-10g (Australian Meat Notice 2007/17), or an average of 6.25g (to give a total sample weight of 375g). Increasing the size of sample unit has the possible effect of increasing the prevalence purely by virtue that sampling more meat is more likely to result in detecting contamination (when it is present on the piece of meat or in a carton).

It should be noted that the binomial approach does not incorporate the lot size. That is because for the binomial approach to be applicable, the lot size has to be large in relation to the amount of meat sampled. Provided the total amount sampled is less than about 10% of the amount of meat in the lot, there is no impact on the probability calculations irrespective of how big the lot is (provided the assumptions above are met). Consequently, there is no impact on the probability of detection when sampling 375g from a lot weighing 5t (about 175 cartons) or 20t (about 700 cartons).