Using the ICMSF Sampling Plan Tool to assess the performance of a Microbiological Criteria

Part 2: MC for *Listeria monocytogenes* in RTE Foods not supporting growth



Leon G.M. Gorris, PhD
Food Safety Expert, The Netherlands

ICMSF Member 2001-present

Outline

- The Microbiological Criterion (MC) for Ready-to-Eat (RTE) foods that do not support growth of *Listeria* monocytogenes (Lm)
- Calculate the performance of the sampling plan with the ICMSF sampling plan tool
- Using the tool for further interpretations of the MC
 - Changing the standard deviation
 - Changing the confidence level

Microbiological criterion for ready-to-eat foods in which growth of L. monocytogenes will not occur

Point of application	Microorganism	n	с	m	Class Plan
Ready-to-eat foods from the end of manufacture or port of entry (for imported products), to the point of sale	monocytogenes	5 ª	0	100 cfu/g ^b	2 °

Where n = number of samples that must conform to the criterion; c = the maximum allowable number of defective sample units in a 2-class plan; m = a microbiological limit which, in a 2-class plan, separates acceptable lots from unacceptable lots.

Other methods that provide equivalent sensitivity, reproducibility, and reliability can be employed if they have been appropriately validated (e.g., based on ISO 16140).

^c Assuming a log normal distribution, this sampling plan would provide 95% confidence that a lot of food containing a geometric mean concentration of 93.3 cfu/g and an analytical standard deviation of 0.25 log cfu/g would be detected and rejected based on any of the five samples exceeding 100 cfu/g *L. monocytogenes*.

Such a lot may consist of 55% of the samples being below 100 cfu/g and up to 45% of the samples being above 100 cfu/g, whereas 0.002% of all the samples from this lot could be above 1000 cfu/g.

The typical actions to be taken where there is a failure to meet the above criterion would be to (1) prevent the affected lot from being released for human consumption, (2) recall the product if it has been released for human consumption, and/or (3) determine and correct the root cause of the failure.

^a National governments should provide or support the provision of guidance on how samples should be collected and handled, and the degree to which compositing of samples can be employed.

^b This criterion is based on the use of the ISO 11290-2 method.

Microbiological criterion for ready-to-eat foods in which growth of L. monocytogenes will not occur

Point of application	Microorganism	n	С	m	Class Plan
Ready-to-eat foods from the end of manufacture or port of entry (for imported products), to the point of sale	monocytogenes	5 a	0	100 cfu/g ^b	2 °

Assuming a log normal distribution, this sampling plan would provide 95% confidence that a lot of food containing a geometric mean concentration of 93.3 cfu/g and an analytical standard deviation of 0.25 log cfu/g would be detected and rejected based on any of the five samples exceeding 100 cfu/g *L. monocytogenes*.

Such a lot may consist of 55% of the samples being below 100 cfu/g and up to 45% of the samples being above 100 cfu/g, whereas 0.002% of all the samples from this lot could be above 1000 cfu/g.

Where to find the tool?



HOME ABOUT US V EVENTS IN THE NEWS V MEMBERS V PUBLICATIONS V VIDEO GALLERY V CONTACTS V LINKS





BOOKS







DOWNLOADS



ARTICLES AND PAPERS

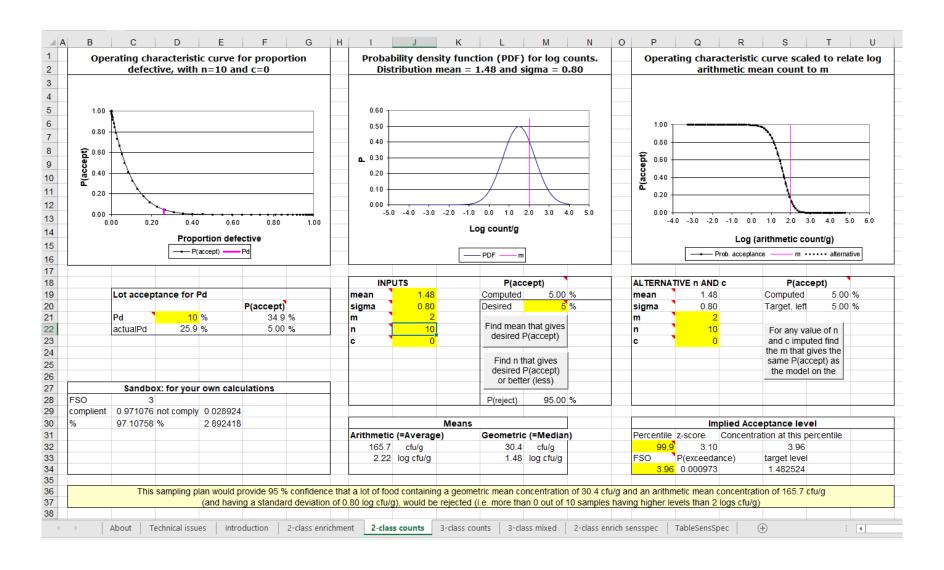
VIDEOS

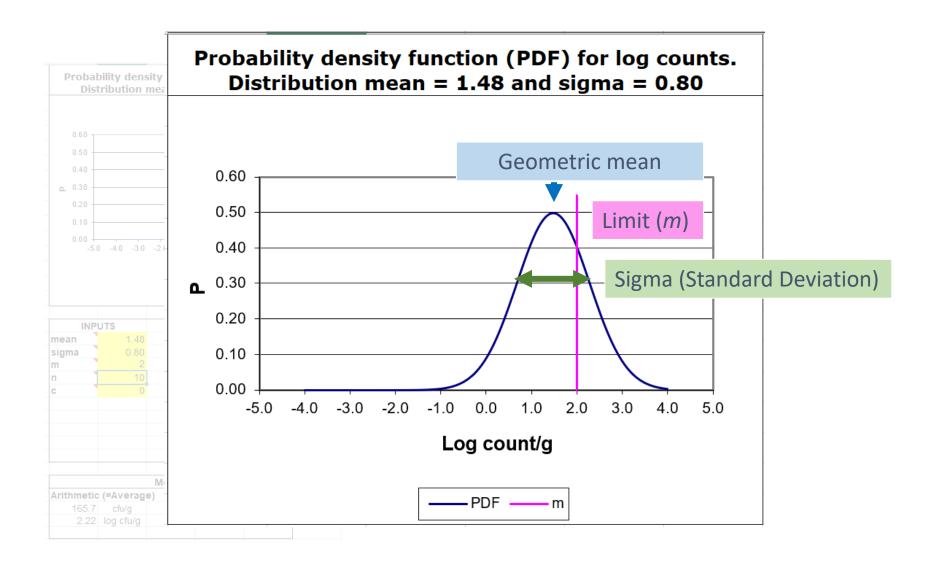
The ICMSF sampling plan tool

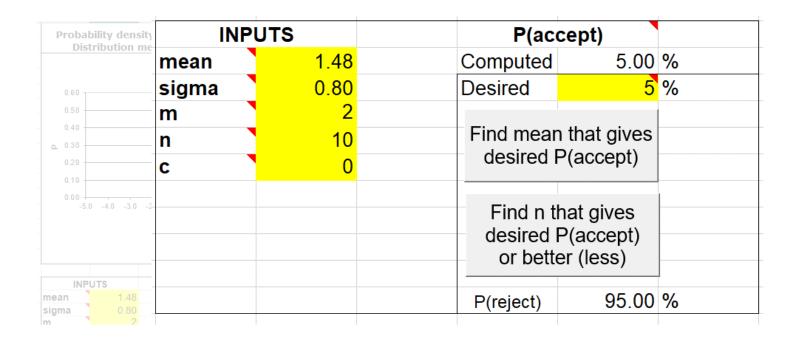


PROTECTED VIEW Be careful—files from the Internet can contain viruses. Unless you need to edit, it's safer to stay in Protected View. Enable Editing SECURITY WARNING Macros have been disabled. Enable Content Micropiological sampling plans: a tool to explore ichish recommendations Version 2.10: Copyright 1998 David Legan & Mark Vandeven 2009 Marcel Zwietering & Peter 5 DISCLAIMER X Disclaimer 5 This spreadsheet calculates probabilities The authors have taken every care to ensure that n this of acceptance for materials with different workbook is accurate. In making this tool available 7 microbial loads and population standard and understanding of the power and limitations of attributes sampling plans 8 deviations. The microbes are assumed neither the authors nor Nabisco Inc. / ICMSF accept any liability for any The authors have taken every care to ensure that the output from this 9 to be lognormally distributed. consequences, direct or indirect resulting from a decision by the user to take workbook is accurate. In making this tool available to promote 10 or not to take, any action based on an output from this workbook discussion and understanding of the power and limitations of 11 12 Version 1.01 attributes sampling plans neither the authors nor ICMSF accept any Version 2.01: Addition of enrichment calculations Version 2.02: 13 Mark Vandeven and David Legan Marcel Zwietering This version sin liability for any consequences, direct or indirect resulting from a with Marcel Zwi 14 Nabisco Research Wageningen University decision by the user to take, or not to take, any action based on an 15 R M Schaeberle Technology Center Laboratory of Food Microbiology output from this workbook. 16 200 DeForest Avenue PO Box 17 Version 2.03-2 17 East Hanover, NJ 07936-1944, USA 6700AA Wageningen The file is further 18 email: VandevenM@Nabisco.com. or The Netherlands 19 LeganD@Nabisco.com email: Marcel.Zwietering@WUR.nl Version 2.07: Version 2.08: OK 21 Background reading Version 2.09: 22 ICMSF book 2 Version 2.10: 23 Calculations for 2-class spreadsheet were checked against ICMSF book 2 Tables 2 and 3 and are in complete agreement. 24 Calculations for 3-class spreadsheet were checked against ICMSF book 2 Table 4 and are in complete agreement. 25 Relating microbiological criteria to food safety objectives and performance objectives 26 M. van Schothorst, M.H. Zwietering, T. Ross, R.L. Buchanan, M.B. Cole, International Commission on Microbiological Specifications for Foods (ICMSF) 27 Food Control 20: 967-979 (2009) 28 Microbial testing in food safety: effect of specificity and sensitivity on sampling plans — how does the OC curve move 29 M.H. Zwietering, H.M.W. den Besten. 30 Current Opinion Food Science 12, 42-51 (2016) 31 32 Acknowledgment 33 The curves showing the distribution of organisms in foods for entered mean counts and sigma values 34 were inspired by unpublished work circulated to ICMSF members by Suzanne Dahms 35 Development of this tool was inspired by Martin Cole 36 37 38 39 40 update > 9-05-20 41 update 2009-10-7: reduced further in calculations, removed empty modules, updated MZ 42 update 2009-10-12: returned to goalseek, no need for solver and analytic toolpack 43 update 2009-10-14: limited c maximally at 1e10 cfu/g in poissonlognormal 44 update 2009-10-15: limited sigma > 0.05 45 update 2009-10-27: protected help file in Introduction 46 update 2010-10-19; added log arithmetic means 47 update 2013-11-25; version 07; added boxes with performance, changed graph to arithmetic mean 48 update 2014-10-23; version 08; sensitivty and selectivity added and boxes placed 2-class counts About Technical issues Introduction 2-class enrichm 3-class mixed 2-class enrich sensspec TableSensSpec

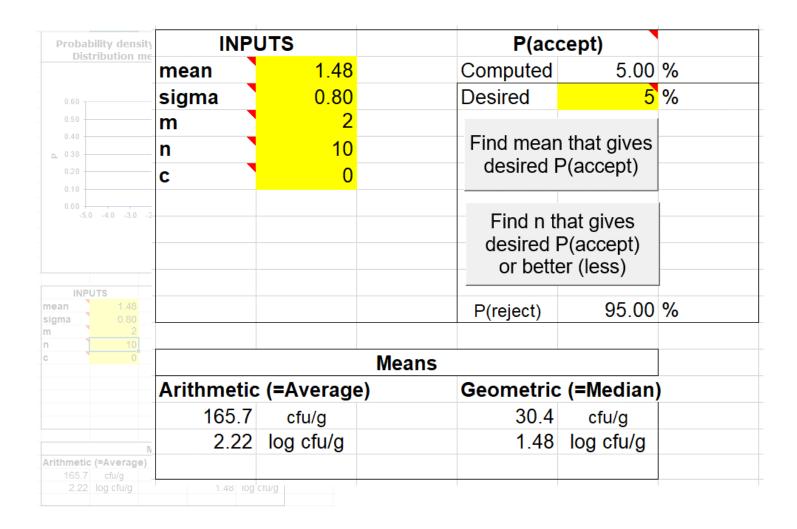




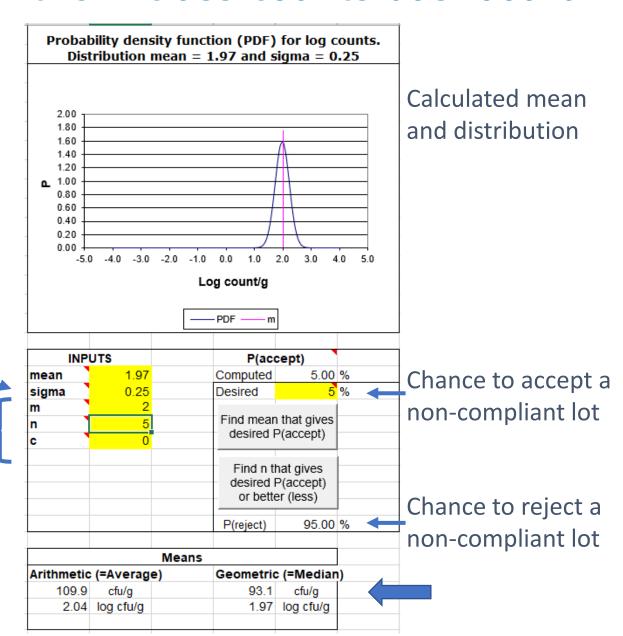




9



The Lm MC in the 2-class counts dashboard



Standard deviation

n, c, m from sampling plan

The Lm MC in the 2-class counts dashboard

INP	INPUTS		P(accept)		
mean	1.97		Computed	5.00	%
sigma	0.25		Desired	5	%
m	2				1
n	5			that gives	
С	0		desired I	P(accept)	
			Find n that gives desired P(accept) or better (less)		
			P(reject)	95.00	%
		Means			
Arithmetic	: (=Averag	e)	Geometric	: (=Median)
109.9	cfu/g		93.1	cfu/g	
2.04	log cfu/g		1.97	log cfu/g	

This sampling plan would provide 95 % confidence that a lot of food containing a geometric mean concentration of 93.1 cfu/g and an arithmetic mean concentration of 109.9 cfu/g (and having a standard deviation of 0.25 log cfu/g), would be rejected (i.e. more than 0 out of 5 samples having higher levels than 2 logs cfu/g)

This sampling plan would provide

- 95% confidence that a lot of food containing a
- geometric mean concentration of 93.1 cfu/g and an
- arithmetic mean concentration of 109.9 cfu/g
- (and having a standard deviation of 0.25 log cfu/g),
- would be rejected
- (i.e. more that 0 out of 5 samples having higher levels than 2 logs cfu/g)

The Lm MC in the 2-class counts dashboard

INP	INPUTS		P(accept)		
mean	1.97		Computed	5.00	%
sigma	0.25		Desired	5	%
m	2				1
n	5			that gives	
С	0		desired I	P(accept)	
			Find n that gives desired P(accept) or better (less)		
			P(reject)	95.00	%
		Means			
Arithmetic	: (=Averag	e)	Geometric	: (=Median)
109.9	cfu/g		93.1	cfu/g	
2.04	log cfu/g		1.97	log cfu/g	

This sampling plan would provide 95 % confidence that a lot of food containing a geometric mean concentration of 93.1 cfu/g and an arithmetic mean concentration of 109.9 cfu/g (and having a standard deviation of 0.25 log cfu/g), would be rejected (i.e. more than 0 out of 5 samples having higher levels than 2 logs cfu/g)

This sampling plan would provide

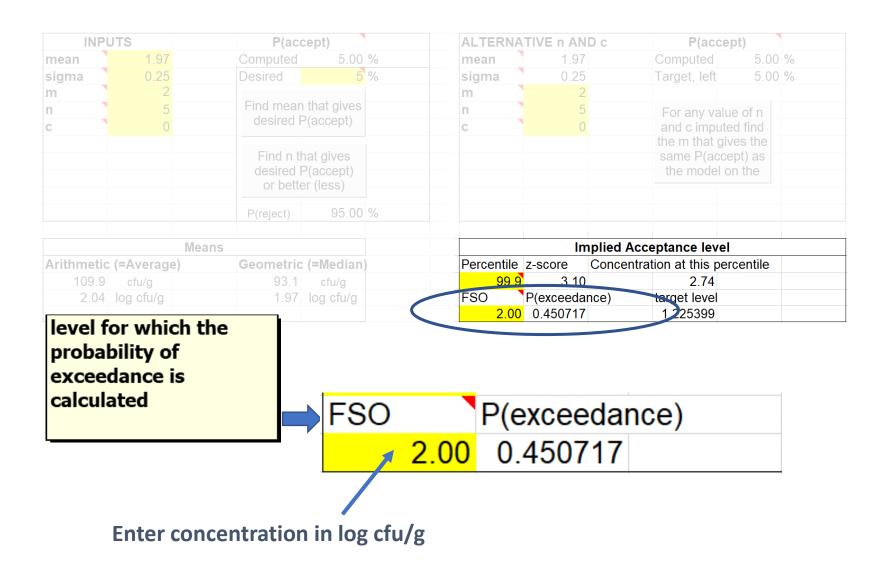
- 95% confidence that a lot of food containing a
- geometric mean concentration of 93.1 cfu/g and an
- arithmetic mean concentration of 109.9 cfu/g
- (and having a standard deviation of 0.25 log cfu/g),
- would be rejected
- (i.e. more that 0 out of 5 samples having higher levels than 2 logs cfu/g)

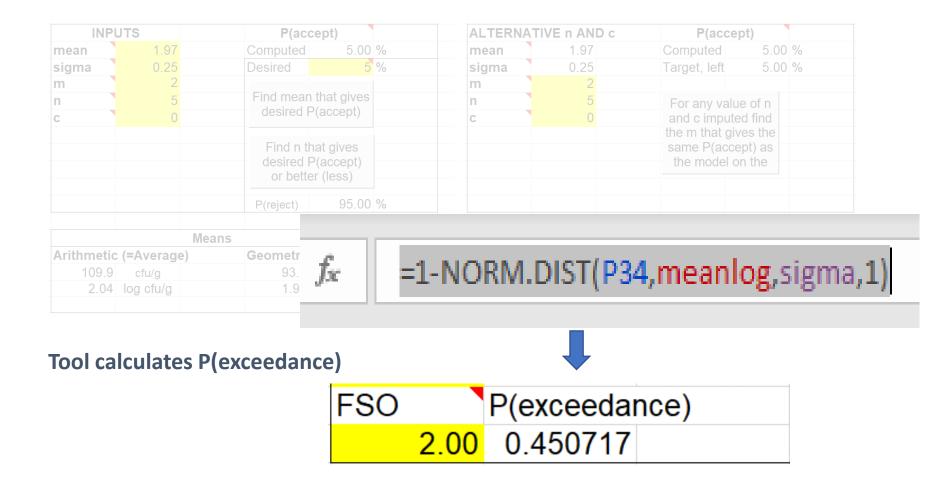
Microbiological criterion for ready-to-eat foods in which growth of L. monocytogenes will not occur

Point of application	Microorganism	n	С	m	Class Plan
Ready-to-eat foods from the end of manufacture or port of entry (for imported products), to the point of sale	monocytogenes	5 ^a	0	100 cfu/g ^b	2 °

^c Assuming a log normal distribution, this sampling plan would provide 95% confidence that a lot of food containing a geometric mean concentration of 93.3 cfu/g and an analytical standard deviation of 0.25 log cfu/g would be detected and rejected based on any of the five samples exceeding 100 cfu/g *L. monocytogenes*.

Such a lot may consist of 55% of the samples being below 100 cfu/g and up to 45% of the samples being above 100 cfu/g, whereas 0.002% of all the samples from this lot could be above 1000 cfu/g.





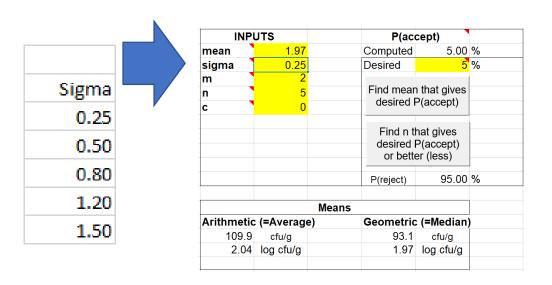
Concentration (cfu/g)	P(exceed) calculation	Proportion of food lot over chosen concentration
10	FSO P(exceedance) 1.00 0.999947	99.99 %
100	FSO P(exceedance) 2.00 0.450717	45.07 %
500	P(exceedance) 2.70 0.001729	0.17 %
1000	FSO P(exceedance) 3.00 1.86E-05	0.00186 %
5000	FSO P(exceedance) 3.70 2.2E-12	0.0000000022 %

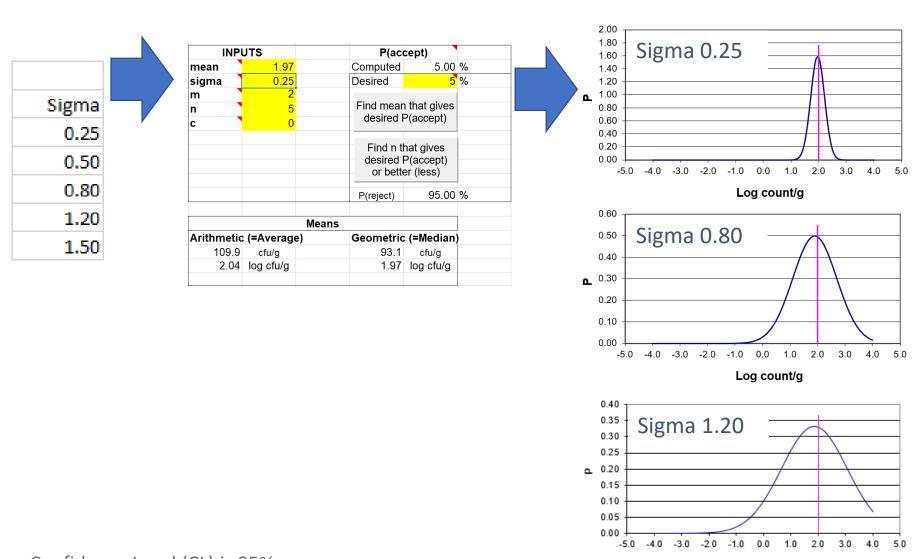
MC performance: Changing assumptions

Microbiological criterion for ready-to-eat foods in which growth of *L. monocytogenes* will not occur

n	С	m	Class Plan
5 ^a	0	100 cfu/g ^b	2 °

Standard Deviation (SD) = sigma = 0.25 log cfu/g Confidence Level (CL) = P(reject) = 95%



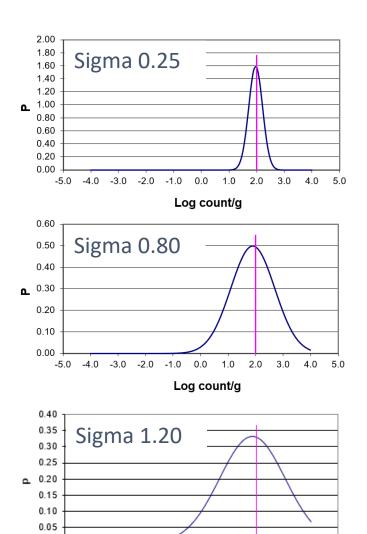


Confidence Level (CL) is 95% Sigma represents Standard Deviation (SD)

Log count/g

	Sigma	Arithme	tic mean	Geomet	ric mean
	log cfu/g	cfu/g	log cfu/g	cfu/g	log cfu/g
Code	e× 0.25			93.1	1.97
	0.50			86.7	1.94
	0.80			79.6	1.90
	1.20			71.0	1.85

Limit m = 2 log cfu/g



0.00

-4.0

-3.0

-2.0

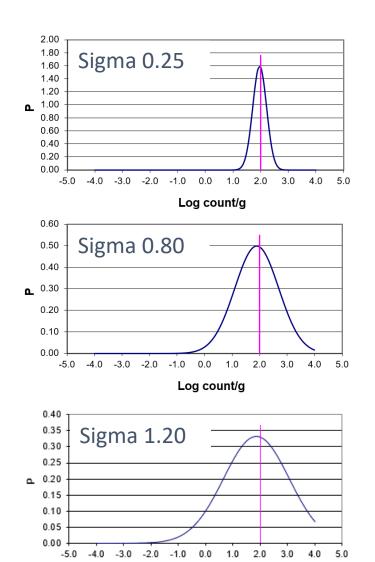
4.0

2.0 3.0

Log count/g

	Sigma	Arithme	tic mean	Geomet	ric mean
	log cfu/g	cfu/g	log cfu/g	cfu/g	log cfu/g
Cod	ex 0.25	110	2.04	93.1	1.97
	0.50	168	2.23	86.7	1.94
	0.80	434	2.64	79.6	1.90
	1.20	3230	3.51	71.0	1.85

Limit m = 2 log cfu/g



Log count/g

Impact of sigma on lot status

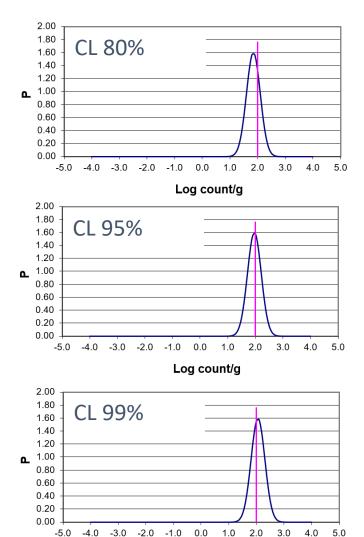
Calculate using P(exceedance)

	Sigma	Estima	Estimated Lm levels in proportions (%) of a food lot							
	(log cfu/g)	≤ 100 (2 log)	> 100 (2 log)	> 1000 (3 log)	> 10000 (4log)	> 100000 (5 log)				
Code	0.25	55 %	45 %	0.002 %	0.00000	0.00000				
	0.80	55 %	45 %	8.5 %	0.43 %	0.00536 %				
	1.20	55 %	45 %	16.9 %	3.67 %	0.435 %				

Changing assumptions: confidence level

	6 6 1	A •••	. •		•
	Confidence	Arithme	tic mean	Geomet	ric mean
	Laval (0/)	of/o	log of /a	of/a	log of /a
	Level (%)	cju/g	iog cju/g	cju/g	iog cju/g
	80.00			70.90	1.85
	00.00			70.50	1.00
	90.00			82.50	1.92
Cod	ex 95.00			93.10	1.97
	99.00			116.00	2.06
	99.90			147.00	2.17

Limit m = 2 log cfu/g



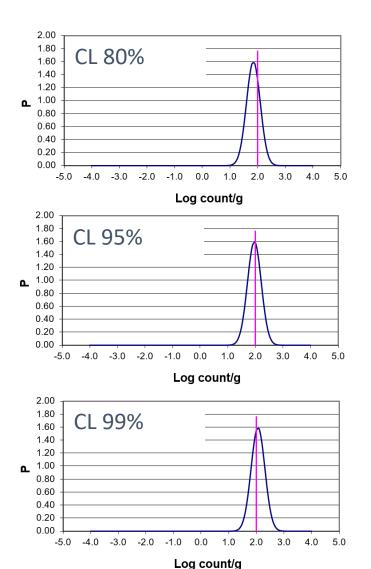
Log count/g

Standard Deviation is 0.25 log cfu/g Confidence Level (CL) represents P(reject)

Changing assumptions: confidence level

	Confidence	Arithma	tic mean	Geomet	ric maan
	Commutative	Andmile	tic illean	Geomet	iic iiicaii
	Level (%)	cfu/g	log cfu/g	cfu/g	log cfu/g
	80.00	83.70	1.92	70.90	1.85
	90.00	97.40	1.99	82.50	1.92
Cod	ex 95.00	109.90	2.04	93.10	1.97
	99.00	136.90	2.14	116.00	2.06
	99.90	173.50	2.24	147.00	2.17

Limit m = 2 log cfu/g



Standard Deviation is 0.25 log cfu/g Confidence Level (CL) represents P(reject)

Changing confidence level and standard deviation

Sigma	Confidence	Arithmetic mean		Geometric mean	
log cfu/g	Level (%)	cfu/g	log cfu/g	cfu/g	log cfu/g
0.25	80.00				
0.25	90.00				
0.25	95.00	109.90	2.04	93.10	1.97
0.80	80.00				
0.80	90.00				
0.80	95.00				
1.20	80.00				
1.20	90.00				
1.20	95.00				

Standard Deviation is 0.25 log cfu/g
Confidence Level (CL) represents P(reject)

Limit $m = 2 \log cfu/g$

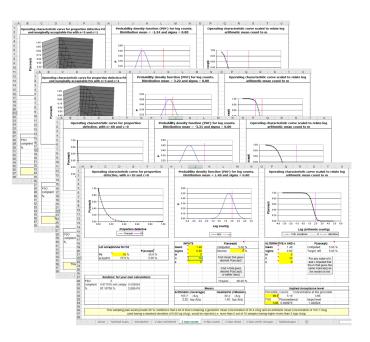
Changing assumptions: confidence level

Sigma	Confidence	Arithmetic mean		Geometric mean	
log cfu/g	Level (%)	cfu/g	log cfu/g	cfu/g	log cfu/g
0.25	80.00	83.70	1.92	70.90	1.85
0.25	90.00	97.40	1.99	82.50	1.92
0.25	95.00	109.90	2.04	93.10	1.97
0.80	80.00	181.60	2.26	33.30	1.52
0.80	90.00	294.70	2.47	54.00	1.73
0.80	95.00	434.30	2.64	79.60	1.90
1.20	80.00	873.70	2.94	19.20	1.28
1.20	90.00	1805.50	3.26	39.70	1.60
1.20	95.00	3230.80	3.51	71.00	1.85
dard Deviation	is 0.25 log cfu/g	Limit <i>m</i> =	Limit m = 2 log cfu/g		

Standard Deviation is 0.25 log cfu/g Confidence Level (CL) represents P(reject)

27

Summary



- The ICMSF sampling plan tool can be used to calculate and further interpret the performance of sampling plans included in Microbiological Criteria
- The robustness of key assumptions should be understood and choices should be well informed.
- See our other clips to understand how the ICMSF Sampling plan tool can be used to assess and interpret the performance of these MCs

