

# Modeling Salmonella Growth for Small and Very Small Processors with Limited Data

**Moderator:** Dennis Seman, DL Seman Consulting, L.L.C.

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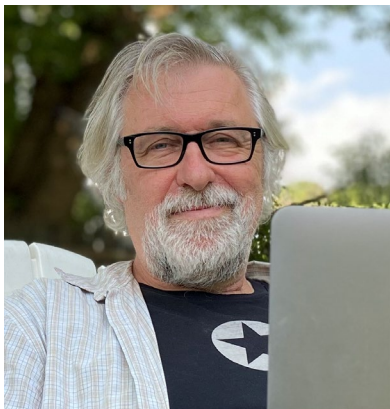
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# Today's Panelists



Dr. Dickson is currently a Professor in the Department of Animal Science and the Inter-Departmental Program in Microbiology at Iowa State University. Dr. Dickson's research focuses on the control of bacteria of public health significance in foods of animal origin. Prior to his appointment at Iowa State University in 1993, he was employed by USDA-ARS as a Research Food Technologist and lead scientist of the Meat Safety Assurance Program, located at the Roman L. Hruska U.S. Meat Animal Research Center, Clay Center, NE. Dr. Dickson was employed in the food industry for three years before joining USDA-ARS. He is a Fellow in the American Academy of Microbiology, and is a Past President of the International Association for Food Protection. He is also active in the American Society for Microbiology and a Certified Food Technologist with the Institute of Food Technologists. Dr. Dickson is currently serving on the National Advisory Committee for Microbiological Specifications for Foods and chairs the U.S. mirror committee for ISO that addresses food safety standards (ISO 22000 series).



Dr. Donald W. Schaffner is Extension Specialist in Food Science and Distinguished Professor at Rutgers University. He is the current Chair of the Rutgers Department of Food Science. His research interests include handwashing, cross-contamination and quantitative microbial risk assessment. He is a Fellow of the Institute of Food Technologists, the American Academy of Microbiology, International Association for Food Protection (IAFP) and the Society for Risk Analysis. Dr. Schaffner was the president of IAFP in 2013-2014. He co-hosts the Food Safety Talk and Risky or Not podcasts.

# **Modeling to Determine if Cooking Cured Bone-In Ham Meets Regulatory Requirements**

**JIM DICKSON**

**DEPARTMENT OF ANIMAL SCIENCE**

**IOWA STATE UNIVERSITY**

# Lethality

- ▶ **FSIS Cooking Guideline for Meat and Poultry Products (Revised Appendix A) December, 2021**
- ▶ [https://www.fsis.usda.gov/sites/default/files/media\\_file/2021-12/Appendix-A.pdf](https://www.fsis.usda.gov/sites/default/files/media_file/2021-12/Appendix-A.pdf)

# Critical Operating Parameters

- ▶ **Come Up Time (CUT)**
  - ▶ An establishment must also consider the heating CUT to be a critical operating parameter unless the establishment can provide a science-based rationale why heating CUT does not need to be addressed.
  - ▶ **Come-Up-Time Option: Total time product temperature is between 50°F and 130°F is 6 hours or less.**
- ▶ **BUT: USDA FSIS recognizes that large diameter products may have longer come up times (CUT)**

# Predictive Microbial Modeling to Support CUT

- ▶ Alternatively, **establishments may use predictive microbiology modeling to develop custom critical operating parameters.** Predictive food microbiology uses models (i.e., mathematical equations) to describe the growth, survival, or inactivation of microbes in food systems from knowledge of the intrinsic and extrinsic factors of the food over time.

# Cooking Deviations

## Long Heating Come Up Time (CUT)

- ▶ If modeling estimates the growth of vegetative pathogens to be 1-Log or less, modeling is adequate to show that the process prevented vegetative pathogen outgrowth



# What do we need to model?

- ▶ **Physical properties of the food (pH, NaCl or  $A_w$ , nitrite, etc.)**
- ▶ **Elapsed time - the time, in hours, between temperatures**
- ▶ **Temperature**

# Cooked Cured Ham

## Elapsed Time and Linear Model

- ▶ **Starting temperature and time**
  - ▶ 50°F, 7:40AM (0 hours)
- ▶ **Final temperature and time**
  - ▶ 130°F, 3:30PM (? hours)

# Elapsed Time Example - 01

▶ **Start: 7:40AM 50°F; End :  
3:30PM 130°F**

▶ **7:40 to 8:00 = 20 min**

▶ **8:00 to 9:00 = 60 min**

▶ **9:00 to 10:00 = 60 min**

▶ **10:00 to 11:00 = 60 min**

▶ **11:00 to 12:00 = 60 min**

▶ **12:00 to 1:00 = 60 min**

▶ **1:00 to 2:00 = 60 min**

▶ **2:00 to 3:00 = 60 min**

▶ **3:00 to 3:30 = 30 min**

▶  **$20 + 60 + 60 + 60 + 60 + 60 + 60 + 60 + 60 + 30 = 470$  minutes**

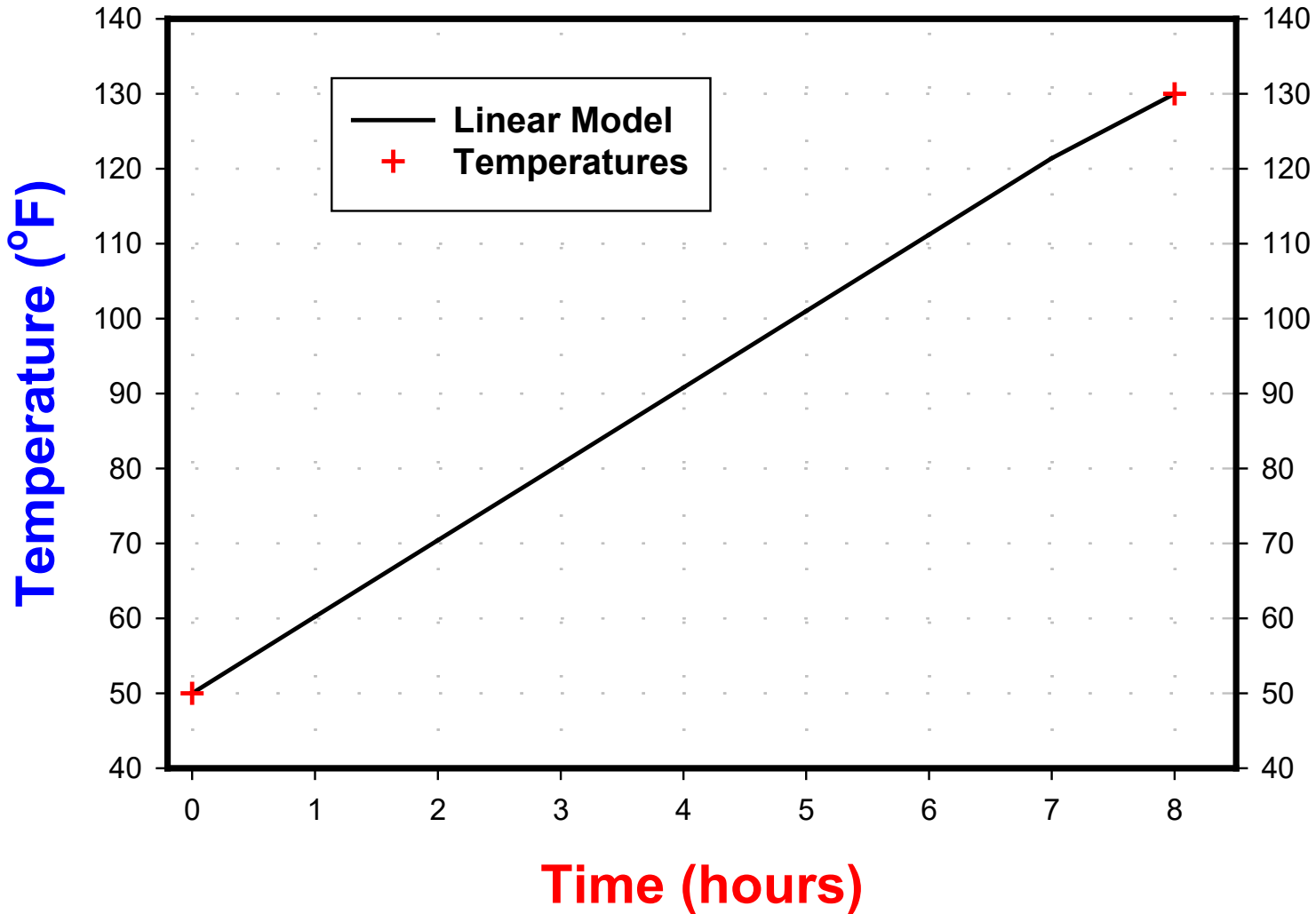
▶  **$470 \text{ minutes} / 60 \text{ minutes per hour} = 7.83$  hours**

# Linear Modeling

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- ▶ **Drawing a straight line between two points**

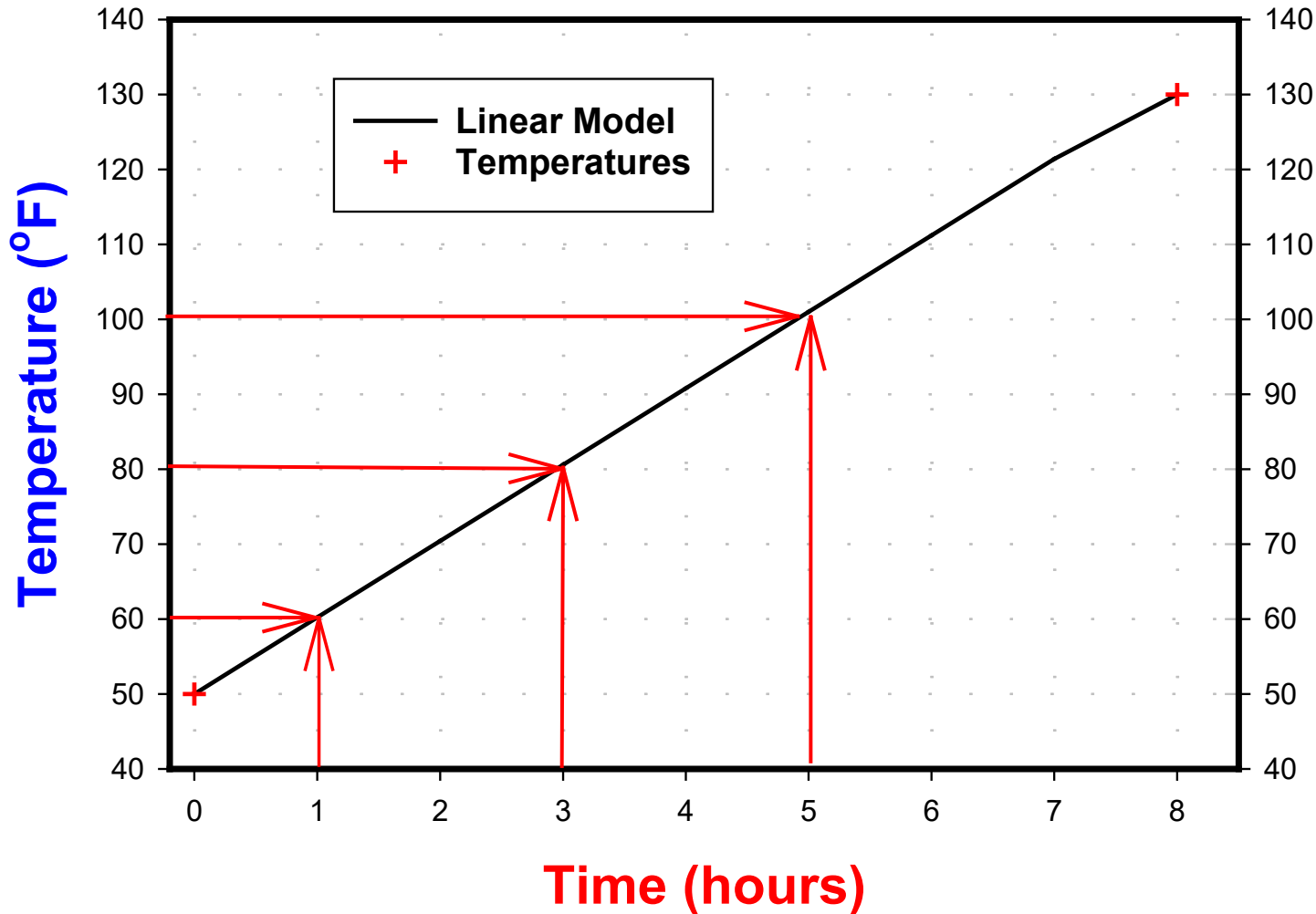
# Linear Model



Time between 50°F and 130°F = ~ 8 hours

**This process is not in compliance**

# Linear Model



**Read 3 additional temperature and time points from the graph**

**That will give 5 points: start, finish and additional points**

# Linear Model – Example 01

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- ▶ Temperature(2) – Temperature(1)
  - ▶ Time(2) – Time(1)

# Linear Model

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- ▶ 130°F – 50°F
  - ▶ 7.83 – 0
    - ▶ Or
      - ▶ 80°F
  - ▶ 7.83 hours
- ▶ 10.2°F/hour



# Linear Model - Example - 01

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Time (hours)	Temperature (°F)
0	50
1	60.2
2	70.4
3	80.6
4	90.8
5	101
6	111.2
7	121.4
8	130

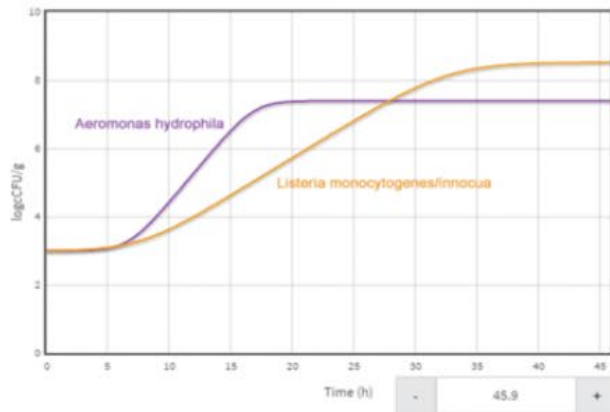
# Linear Modeling

- ▶ **Linear modeling –**
  - ▶ **overestimates pathogen growth during the cooling**
  - ▶ **overestimates pathogen growth**
- ▶ **FSIS considers this modeling result fail-safe**
  
- ▶ **Therefore – more temperature and time information provides a more accurate estimate in the model**

# ComBase

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## ComBase Predictor



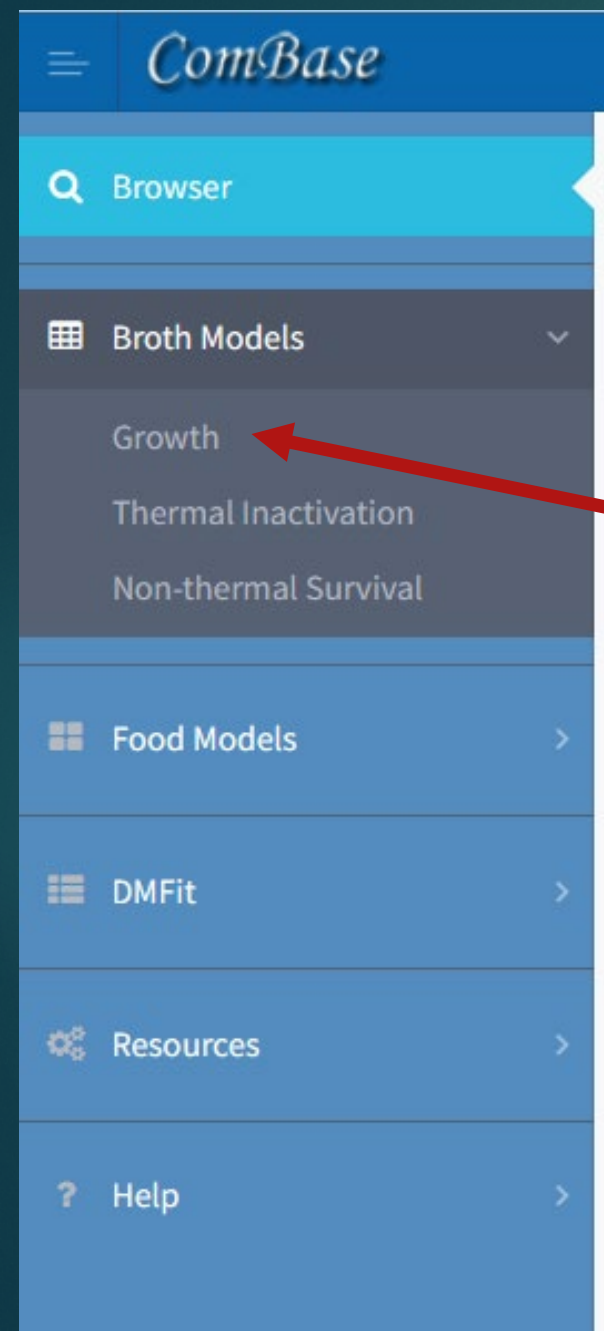
The **ComBase Browser** enables you to search thousands of microbial growth and survival curves that have been collated in research establishments and from publications

The **ComBase Predictive Models** are a collection of software tools based on ComBase data to predict the growth or inactivation of microorganisms

[Login/Register](#)

58,500 + records | 70,000 + users

<https://www.combase.cc/index.php/en/>



- ▶ After log in
- ▶ Upper left corner
- ▶ “Growth” in the “Broth Models” tab

## Growth Model (Disclaimer)

Note: the *Listeria monocytogenes/innocua* (acetic) model has been removed while under review.

[ Static | Dynamic ]

Aeromonas hydrophila

Aeromonas hydrophila

Bacillus cereus

Bacillus cereus (CO<sub>2</sub>)

Bacillus licheniformis

Bacillus subtilis

Brochothrix thermosphacta

Clostridium botulinum (non-prot.)

Clostridium botulinum (prot.)

Clostridium perfringens

Escherichia coli

Escherichia coli (CO<sub>2</sub>)

[A] Listeria monocytogenes/innocua

Listeria monocytogenes/innocua (CO<sub>2</sub>)

Listeria monocytogenes/innocua (lactic)

Listeria monocytogenes/innocua (nitrite)

Pseudomonads

Salmonella spp

Salmonella spp (CO<sub>2</sub>)

Salmonella spp (nitrite)

Shigellae (s.flexneri and relatives) (nitrite)

7

1

37

1

1

1

g.conc/h) 0.433

g CFU/g) 7.39

Min. time(Hours) 0.696

Lag time (Hours) 6.72

▶ Dynamic

▶ Select organism

## Growth Model (Disclaimer)

Note: the *Listeria monocytogenes/innocua* (ace

[ Static | Dynamic ]

Time(h)	Temp (°C)

Salmonella spp (nitrite)

Temperatures range [7,40]

Init. level

Phys.state

pH

Aw | NaCl (%)

nitrite(ppm)

- ▶ Time and Temperature (°C!!)
- ▶ **At least 5 time and temperature points**
- ▶ **Range 7°C to 40°C**
- ▶ Inputs
- ▶ Initial population
- ▶ pH
- ▶ Water activity ( $a_w$ ) or NaCl
- ▶ Nitrite

# Fahrenheit to Celsius Temperature conversion

▶  $128^{\circ}\text{F} = ??^{\circ}\text{C}$

▶  $(128^{\circ}\text{F} - 32^{\circ}\text{F}) * (5/9) = 53.3^{\circ}\text{C}$

▶  $212^{\circ}\text{F} = 100^{\circ}\text{C}$

▶  $32^{\circ}\text{F} = 0^{\circ}\text{C}$

# Fahrenheit to Celsius Temperature conversion

- ▶ In MS-Excel
- ▶ Covert temperature “from” “to”

<u>Temp F</u>	<u>Temp C</u>	<u>Formula</u>
50	10.0	=CONVERT (A3,"F","C")
60	15.6	=CONVERT (A4,"F","C")
70	21.1	=CONVERT (A5,"F","C")
80	26.7	=CONVERT (A6,"F","C")
90	32.2	=CONVERT (A7,"F","C")
100	37.8	=CONVERT (A8,"F","C")
110	43.3	=CONVERT (A9,"F","C")
120	48.9	=CONVERT (A10,"F","C")
130	54.4	=CONVERT (A11,"F","C")

=CONVERT(temp,"F","C")



# Fahrenheit to Celsius

## Temperature conversion

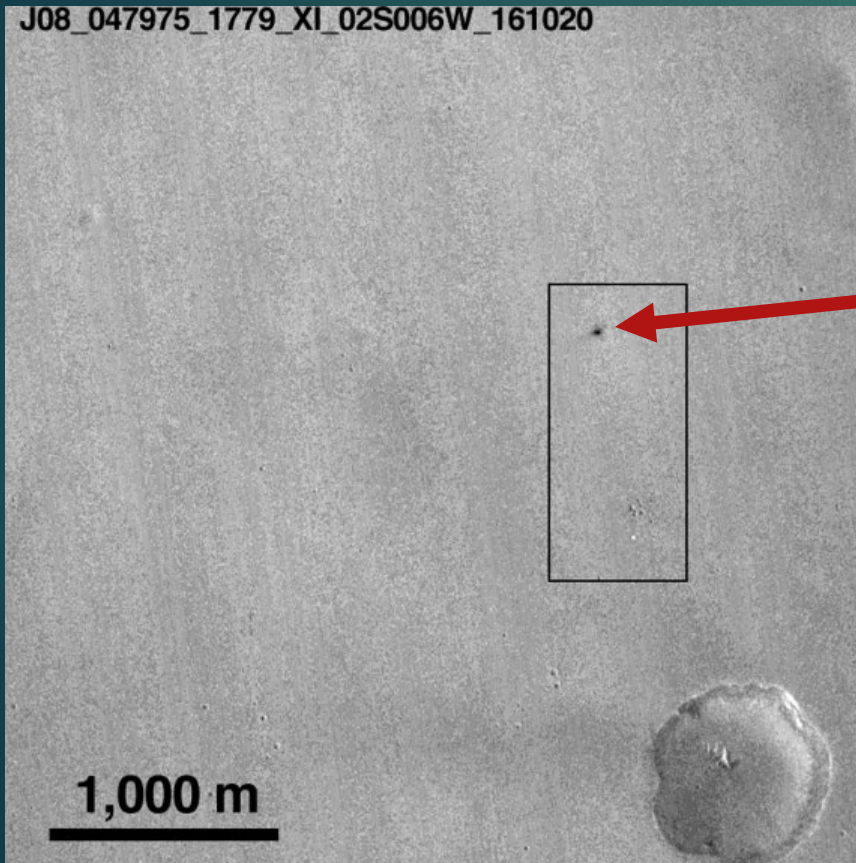
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- ▶ Mars Probe Lost Due to Simple Math Error (01 Oct 1999)

- ▶ **NASA lost its \$125-million Mars Climate Orbiter because spacecraft engineers failed to convert from Imperial to Metric measurements when exchanging vital data before the craft was launched, space agency officials said Thursday.**

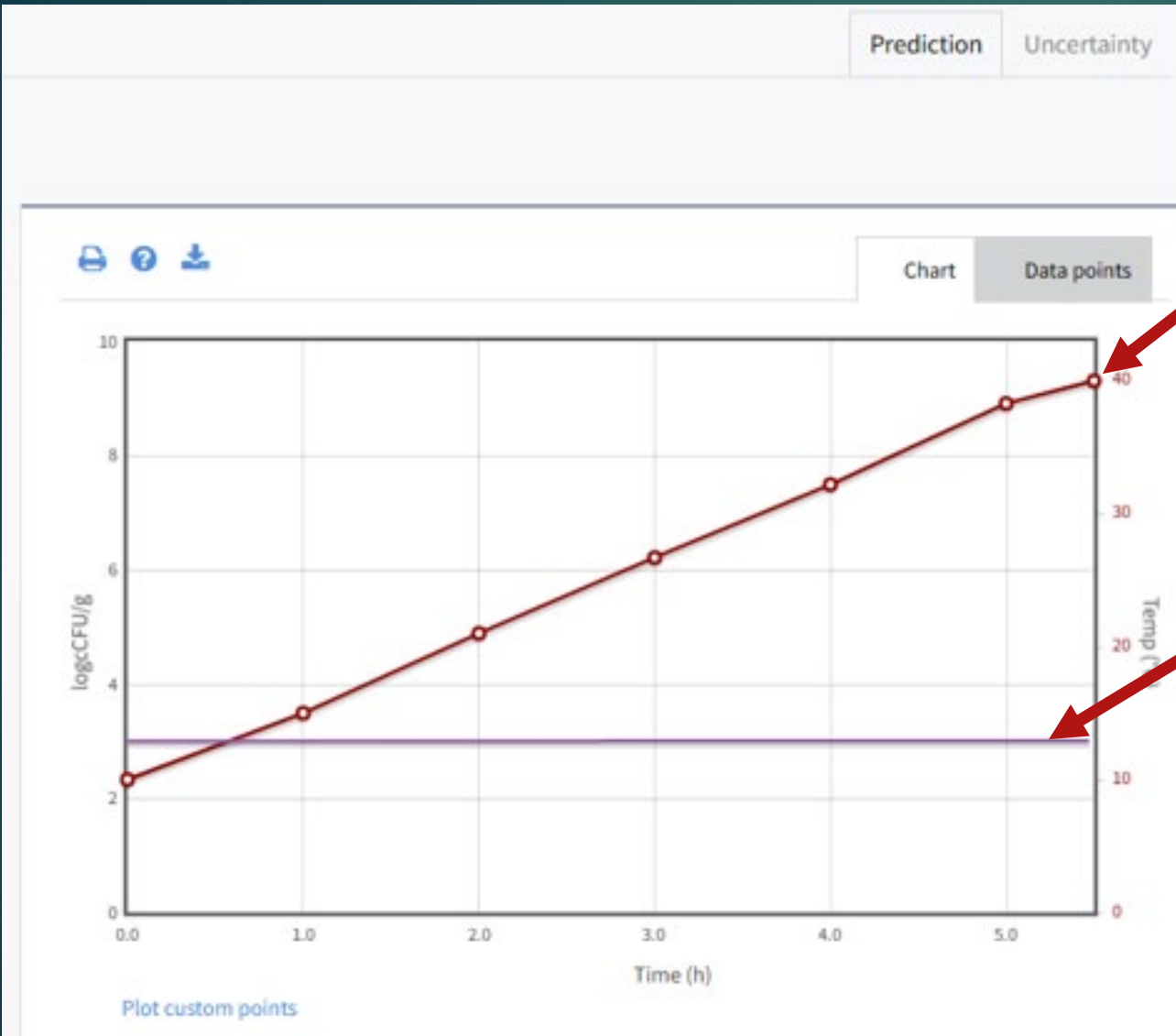
- ▶ As a result, JPL engineers ***mistook acceleration readings measured in Imperial units of pound-seconds for a metric measure of force called newton-seconds.***

- ▶ <https://www.latimes.com/archives/la-xpm-1999-oct-01-mn-17288-story.html>



# ComBase Output

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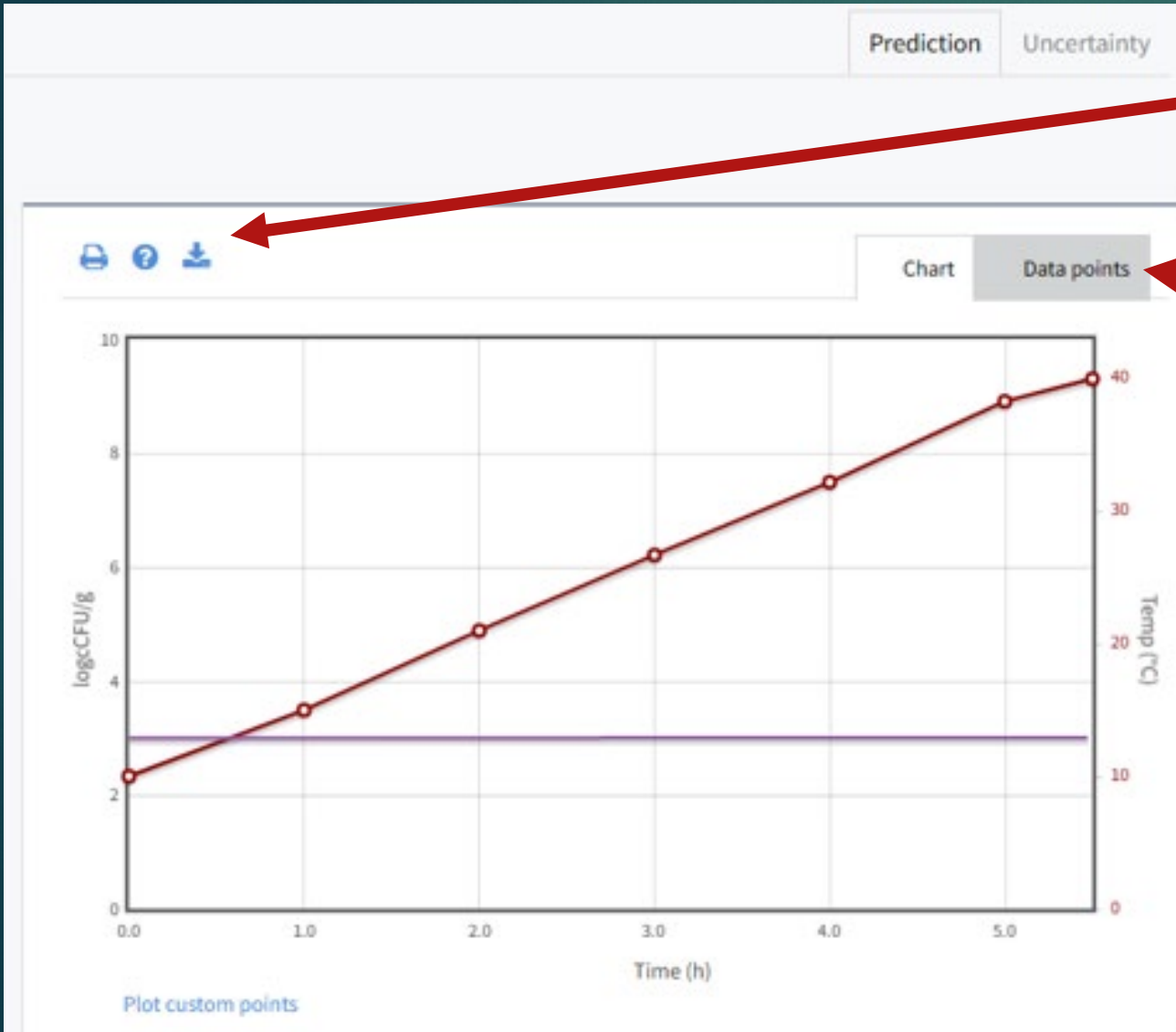


► Temperature

► Predicted Growth

# Output

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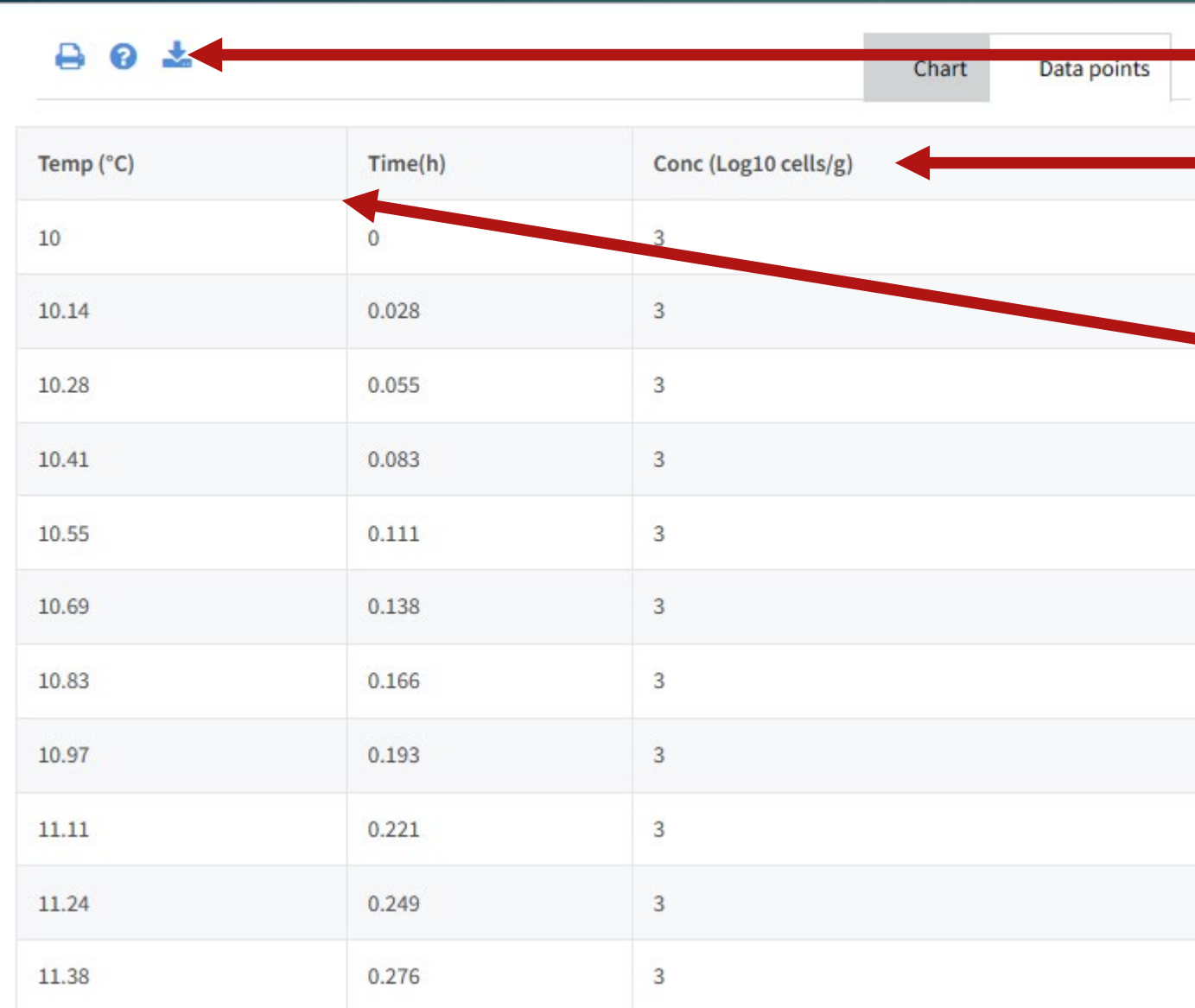


▶ Print or download

▶ Data Points

# Output

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The screenshot shows a data table with a toolbar at the top. The toolbar includes a print icon, a help icon, and a download icon, all of which are pointed to by a red arrow. To the right of the toolbar are two buttons: 'Chart' and 'Data points'. The table has three columns: 'Temp (°C)', 'Time(h)', and 'Conc (Log10 cells/g)'. A red arrow points to the 'Conc' column header, and another red arrow points to the first row of data. The table contains 12 rows of data, showing a steady increase in temperature and time, while the concentration remains constant at 3.

Temp (°C)	Time(h)	Conc (Log10 cells/g)
10	0	3
10.14	0.028	3
10.28	0.055	3
10.41	0.083	3
10.55	0.111	3
10.69	0.138	3
10.83	0.166	3
10.97	0.193	3
11.11	0.221	3
11.24	0.249	3
11.38	0.276	3

- ▶ Print or download
- ▶ Predicted Population
- ▶ Temperature and Time

What does it mean?

Long Heating Come Up Time (CUT)

- ▶ “If modeling estimates the growth of vegetative pathogens to be 1-Log or less, modeling is adequate to show that the process prevented vegetative pathogen outgrowth”
- ▶ **ComBase growth model predicts  $< 1 \log_{10}$  increase in population**

# Documentation

- ▶ **Physical parameters of the product (pH, Nitrite, NaCl or  $A_w$ )**
- ▶ **Available temperature and time information**
- ▶ **Model results**
  
- ▶ ***All parameters should represent the “worst case” scenario***



RUTGERS

New Jersey Agricultural  
Experiment Station

# Modeling below the limits of the model and cooking deviations

Don Schaffner, PhD

Distinguished Professor and Extension Specialist  
Chair, Department of Food Science

# Outline

- Modeling below the limits of the model
  - The limits of a model are often inside the limits of the growth of the organism
  - There are different solutions to the problem
- Cooking deviations with limited data
  - Modeling cooking deviations requires use of  $D$  and  $z$  values
  - My preferred tool is the NAMI spreadsheet



# Limits of the organism

- Limits to growth (NACMCF 2010)
  - ICMSF v 5 (1996)
  - FDA (2011) Fish and Fisheries Products Hazards and Controls Guidance. Appendix 4.

Pathogen		Temp (°C)		pH		a <sub>w</sub>	Water Phase NaCl (%)
		Min	Max	Min	Max	Min	Max
<i>B. cereus</i>	FDA	4	55	4.3	9.3	0.92	10
	ICMSF	4	55	5.0	8.8	0.93	
<i>C. bot</i> (growth only) (Proteolytic)	FDA	10	48	4.6	9	0.93	10
	ICMSF	10-12		4.6		0.93	10
<i>C. bot</i> (growth only) (Non-proteolytic)	FDA	3.3	45	5	9	0.97	5
	ICMSF	3.3		5.0		0.97	5
<i>C. perfringens</i>	FDA	10	52	5	9	0.93	7
	ICMSF	12	50	5.5	9.0	0.97	
Pathogenic <i>E. coli</i>	FDA	6.5	49.4	4	9	0.95	6.5
	ICMSF	7-8	44-46	4.4	9.0	0.95	
<i>E. coli</i> O157:H7	ICMSF	8	44-45	4.5			Slow at 6.5 NG at 8.5
<i>L. monocytogenes</i>	FDA	-0.4	45	4.4	9.4	0.92	10
	ICMSF	-0.4	45	4.39	9.4	0.92	
<i>S. aureus</i> (growth only)	FDA	7	50	4	10	0.83	20
	Aerobic conditions	ICMSF	7	48	4	10	0.83
	Anaerobic conditions	ICMSF			5.0		0.90
<i>Salmonella</i>	FDA	5.2	46.2	3.7	9.5	0.94	8
	ICMSF	5.2	46.2	3.8	9.5	0.94	

# Limits of the model

- Also from NACMCF (2010)

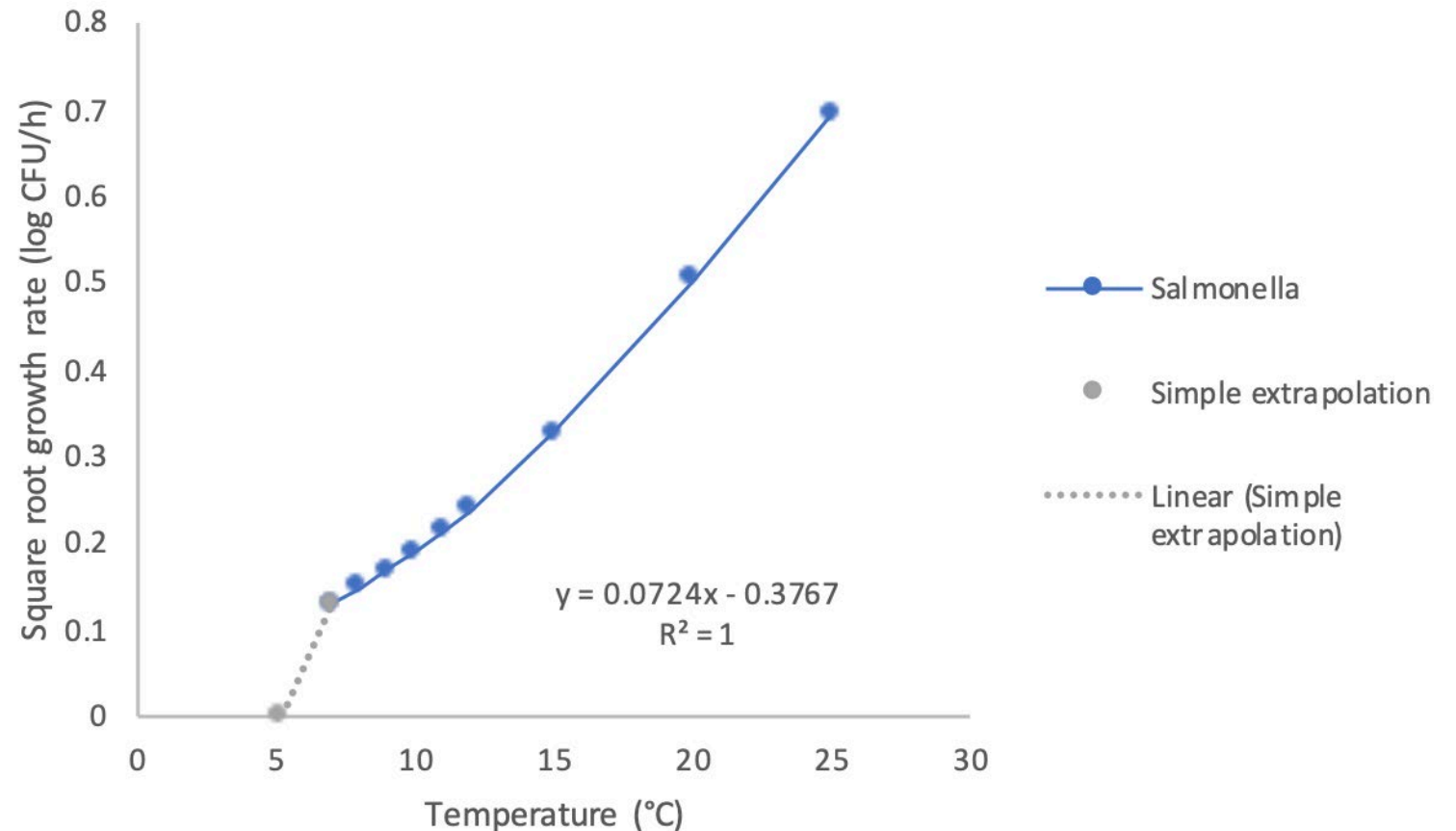
	ComBase <sup>2</sup>					PMP <sup>3</sup>				
	Temperature (°C)		pH		a <sub>w</sub>	Temperature (°C)		pH		a <sub>w</sub>
	Min	Max	Min	Max	Min	Min	Max	Min	Max	Min
<i>B. cereus</i>										
with CO <sub>2</sub>	5	34	4.9	7.5	0.974					
aerobic						5	42	4.7	7.5	0.97
anaerobic						10	42	5.0	9.0	0.97
<i>C. botulinum</i> (growth only)										
proteolytic	14	40	4.7	7.2	0.954	15	34	5.0	7.2	0.977
non-proteolytic	4	30	5.1	7.5	0.974	5	28	5.0	7.0	0.977
<i>C. perfringens</i>	15	52	5	8	0.971	19	37	6.0	6.5	0.983
<i>E. coli</i> O157:H7										
with CO <sub>2</sub>	10	30	4.5	7	0.961					
aerobic						5	42	4.5	8.5	0.97
anaerobic						5	42	4.5	8.5	0.97
<i>L. monocytogenes</i>										
with CO <sub>2</sub>	1	35	4.4	7.5	0.934					
aerobic						4	37	4.5	7.5	0.928
anaerobic						4	37	4.5	8.0	0.97
<i>S. aureus</i> (growth only)										
not specified	7.5	30	4.4	7.1	0.907					
aerobic						10	42	4.5	9.0	0.911
anaerobic						12	42	5.3	9.0	0.872
<i>Salmonella</i> spp.										
with CO <sub>2</sub>	7	30	3.9	7.4	0.973					
aerobic						10	30	5.6	6.8	0.974

# Three solutions, one wrong

- If the temperature is below the limit of the model, assume the temperature is at the limit of the model
- If the temperature is below the limit of the model, interpolate between the limit of the model and the temperature
- ~~• If the temperature is below the limit of the model, the growth rate of the organism is zero~~

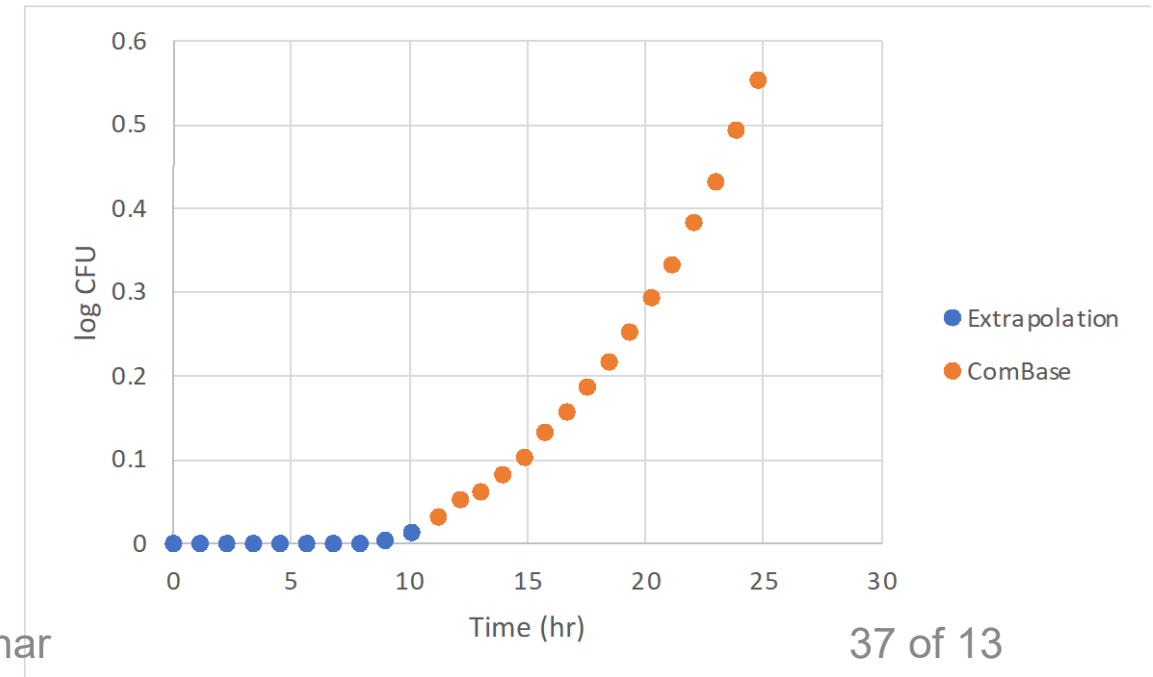
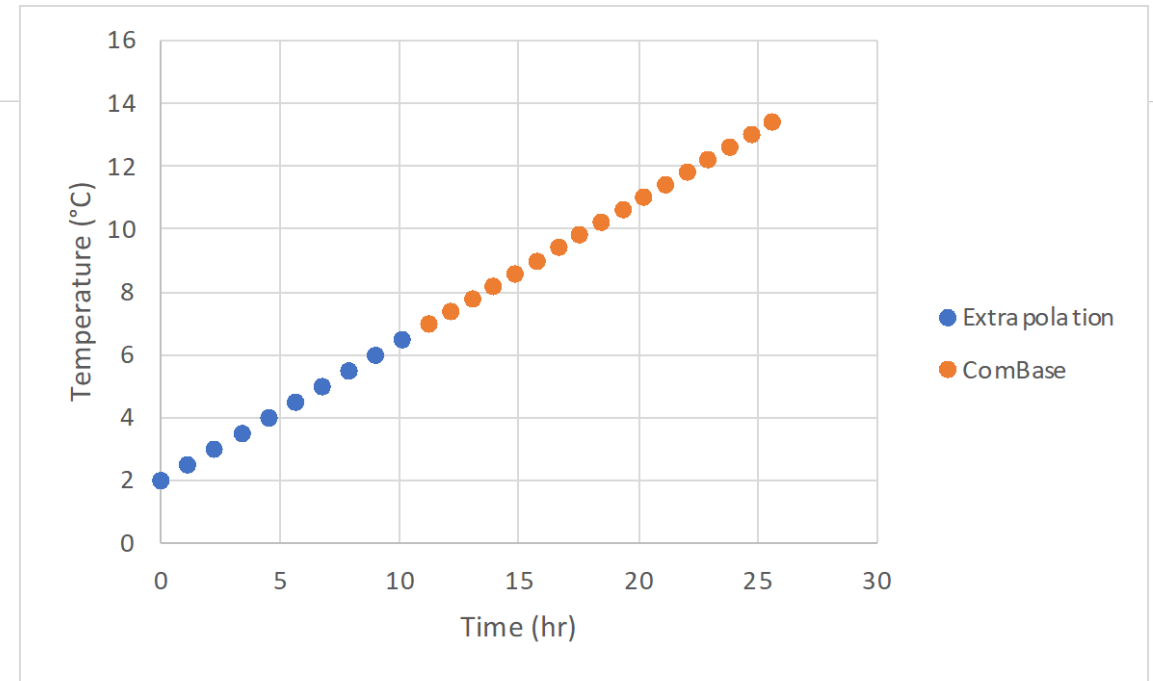
# Extrapolating below the model limit

- Vol 29, No 55 (2021), Revista Alimentos Hoy



# Example

- *Salmonella* growth under constant temperature increase
- A little growth up to 12 hours



# Example

- Company was using 1999 USDA Appendix A
- Cook to 159 °F (instantaneous)
- What to do if 159 °F is not reached?
  - Drop down to lower time and temperature
  - Integrated lethality

## Guidelines for Cooked Beef, Roast Beef, and Cooked Corned Beef

1. Cooked beef and roast beef, including sectioned and formed roasts, chunked and formed roasts, and cooked corned beef can be prepared using one of the following time and temperature combinations to meet either a 6.5- $\log_{10}$  or 7- $\log_{10}$  reduction of Salmonella. The stated temperature is the minimum that must be achieved and maintained in all parts of each piece of meat for a least the stated time:

Minimum Internal Temperature		Minimum processing time in minutes or seconds after minimum temperature is reached	
Degrees Fahrenheit	Degrees Centigrade	6.5- $\log_{10}$ Lethality	7- $\log_{10}$ Lethality
130	54.4	112 min.	121 min.
131	55.0	89 min.	97 min.
132	55.6	71 min.	77 min.
133	56.1	56 min.	62 min.
134	56.7	45 min.	47 min.
135	57.2	36 min.	37 min.
136	57.8	28 min.	32 min.
137	58.4	23 min.	24 min.
138	58.9	18 min.	19 min.
139	59.5	15 min.	15 min.
140	60.0	12 min.	12 min.
141	60.6	9 min.	10 min.
142	61.1	8 min.	8 min.
143	61.7	6 min.	6 min.
144	62.2	5 min.	5 min.
145	62.8	4 min.*	4 min.*
146	63.3	169 sec.	182 sec.
147	63.9	134 sec.	144 sec.
148	64.4	107 sec.	115 sec.
149	65.0	85 sec.	91 sec.
150	65.6	67 sec.	72 sec.
151	66.1	54 sec.	58 sec.
152	66.7	43 sec.	46 sec.
153	67.2	34 sec.	37 sec.
154	67.8	27 sec.	29 sec.
155	68.3	22 sec.	23 sec.
156	68.9	17 sec.	19 sec.
157	69.4	14 sec.	15 sec.
158	70.0	0 sec.**	0 sec.**
159	70.6	0 sec.**	0 sec.**
160	71.1	0 sec.**	0 sec.**

# Cooking deviations with limited data

- NAMI spreadsheet
- <http://www.amif.org/process-lethality/>
- Choice of D and z are important

<b>Date:</b> _____ <b>Organism:</b> _____ <b>Product name:</b> _____		<b>User Must:</b> 1. Identify organism and product of concern 2. Provide at least 20 time/temp data points																																																														
<b>Instructions:</b> 1 Select the organism and product of concern and identify corresponding T ref (Temperature reference), z, and D values and create a similar table to Table 1. These values should be obtained from your own companies challenge study data, from scientific literature, or other reliable sources. These values need to be relevant and appropriate for the type of product and the organism of concern.																																																																
<i>Note: The data in Table 1 has been selected as examples of the type of data that should be collected and extracted from your own companies challenge study data, scientific literature, safe harbor documents, or other reliable sources. The data in Table 1 SHOULD NOT BE USED EXCLUSIVELY AS SUPPORT DATA.</i>																																																																
2 Enter the T ref, z, and D values into the appropriate labeled cells below the table that contains the lethality data from literature. 3 Clear and enter at least 20 time/temperature data points into the data table.																																																																
<i>Note: This model MUST contain a MINIMUM of 20 time/temperature data points. The model is not accurate if less than 20 data time/temperature data points are used.</i>																																																																
4 Once the table is completed, a cumulative F value will be given as the very last number in the right hand column of the data table. This number adds up the lethality values for each time interval and calculates an approximation of the area under the lethal rate curve.																																																																
5 After the data is entered, a core temperature and lethality curve are produced.																																																																
6 The total log reduction of the process is automatically determined by dividing the cumulative F value by the D value that was entered into the appropriate labeled cell. The resulting value equals the total log reduction of the process.																																																																
7 By using these estimates, you or a process authority should determine if the process meets regulatory requirements as safe.																																																																
<a href="#">The USDA's FSIS Appendix A Compliance Guidelines For Meeting Lethality Performance Standards For Certain Meat And Poultry Products that addresses desired log reductions should also be considered when evaluating a lethality process.</a>																																																																
<b>Definitions:</b> <b>D-value:</b> The time (in minutes) at an associated T ref required to kill 90% of the selected microorganism, a one log reduction. <b>z-value:</b> The number of degrees F to change the D value by a factor of ten. <b>F-value:</b> The process lethality. The equivalent time of heating at a reference temperature. Total lethality will be the final computed cumulative F value.																																																																
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<i>Note: This model is a tool for calculating F-values. To ensure correct results, the proper z, T-ref, and D-values for each product and organism must be used.</i>																																																																

# Useful reference



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**Food Microbiology**

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## A systematic approach to determine global thermal inactivation parameters for various food pathogens

Esther D. van Asselt<sup>1</sup>, Marcel H. Zwietering\*

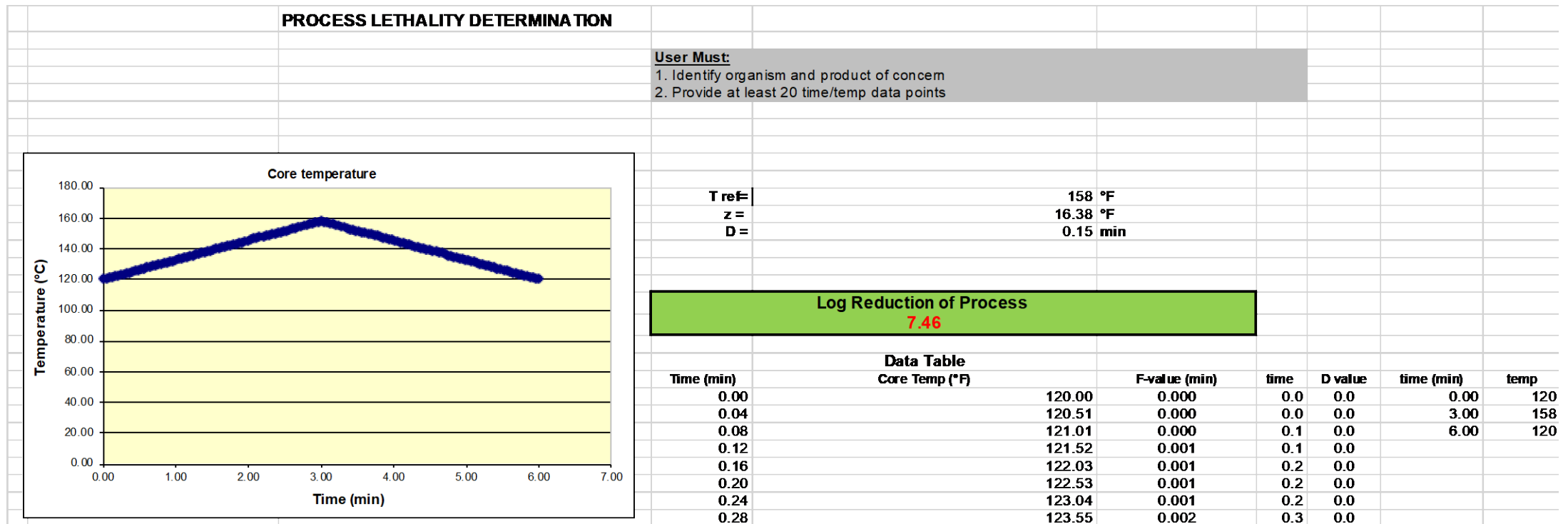
*Wageningen University, Laboratory of Food Microbiology, P.O. Box 8129, 6700 EV Wageningen, The Netherlands*

Received 19 February 2005; received in revised form 3 August 2005; accepted 7 August 2005



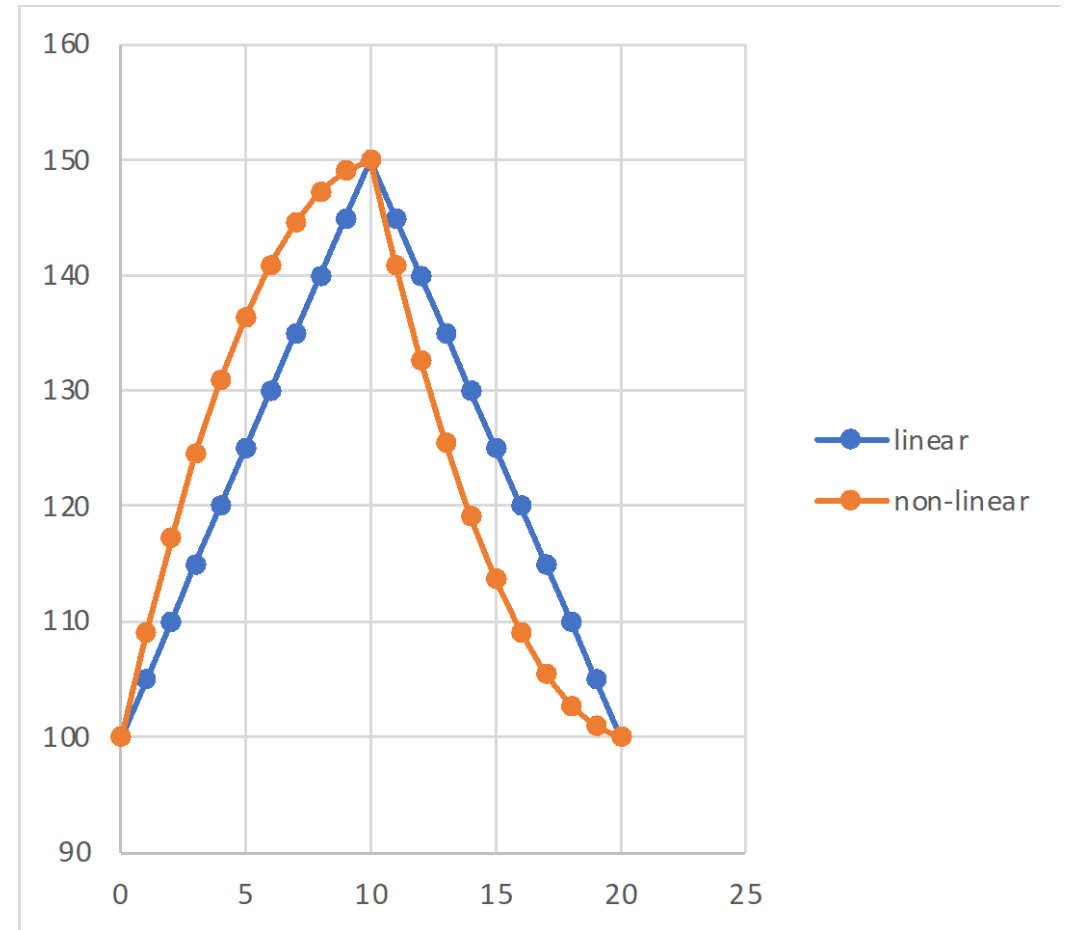
# Using NAMI spreadsheet

- 20 data points are needed, so interpolate



# Straight line interpolation

- May be close to observation
- Under-estimates lethality on rise
- Over-estimates lethality on decline
- Overall underestimation (fail safe) with linear



# Summary

- Two examples today
  - Modeling below the limits of the model
  - Cooking deviations with limited data
- Many problems can be solved with off-the-shelf modeling tools and Excel
- The key is to make fail safe assumptions, and justify them

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September 18, 2023 FDA's Food Traceability Final Rule

October 24, 2023 Managing Meat Shelf Life and Spoilage to Ensure Food Security

<https://www.foodprotection.org/events-meetings/webinars/>



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