

# Investigating the Temperature Dependency and Efficiency of *S. cerevisiae* Using MatLab

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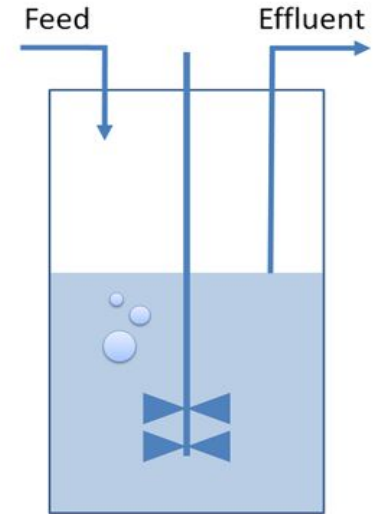
Biology 388  
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# Outline

- Introduction of the chemostat problem of Tai et al (2007)
- Data extracted from the Tai paper (2005) & (2007)
- Methods used to investigate the model
- Temperature investigation
- Efficiency/waste investigation
- Conclusions

# Chemostats are Useful for Controlling Variables

- “The chemostat is an experimental apparatus where the chemical environment can be maintained static and nutrient availability can be controlled by the experimenter.” -ScienceDirect
- Chemostats can be used to investigate the differential utilization of fermentation at different metabolic rates.
- Tai paper experimented with a limiting nutrient- either Glucose (Carbon) or Ammonium (Nitrogen).
- Tai experiment done under anaerobic conditions.
- It is not possible to only change a single parameter without having an effect on any others, even using the chemostat culture approach rather than the batches.



**Inflow = outflow**

# Parameters Included in Tai et al

**Table 1.** Physiological characteristics of *S. cerevisiae* grown in ammonium- and glucose-limited anaerobic chemostat cultures

Limiting nutrient	Growth temperature (°C)	$Y_{Glu/X}$ ( $g_{DW} \cdot g_{glucose}^{-1}$ )	$q_{Glu}^a$	$q_{EtOH}^a$	$q_{CO_2}^a$	Carbon recovery (%)	Residual glucose (mM)	Residual ammonia (mM)
Glucose	12	$0.07 \pm 0.01$	$-2.5 \pm 0.2$	$3.8 \pm 0.3$	$4.4 \pm 0.3$	$100 \pm 3$	$2.8 \pm 1.1$	$65.2 \pm 2.2$
Glucose	30	$0.07 \pm 0.00$	$-2.3 \pm 0.0$	$3.5 \pm 0.0$	$3.8 \pm 0.2$	$95 \pm 1$	$0.3 \pm 0.1$	$61.3 \pm 4.5$
Ammonium	12	$0.05 \pm 0.00$	$-3.6 \pm 0.2$	$6.1 \pm 0.3$	$6.0 \pm 0.6$	$97 \pm 4$	$90.0 \pm 9.8$	$1.5 \pm 0.2$
Ammonium	30	$0.04 \pm 0.00$	$-4.0 \pm 0.1$	$6.8 \pm 0.2$	$7.4 \pm 0.2$	$97 \pm 2$	$85.1 \pm 8.2$	$0.2 \pm 0.1$

Cultures were grown at 30 and 12°C ( $D = 0.03 \text{ h}^{-1}$ ). Values represent the mean  $\pm$  SD of data from three independent steady-state chemostat cultivations.  $Y_{Glu/X}$ , biomass yield on glucose; DW, dry weight.

<sup>a</sup> Values expressed as  $\text{mmol} \cdot g_{DW}^{-1} \cdot \text{h}^{-1}$ .

\*Did not know: r,k,L,F

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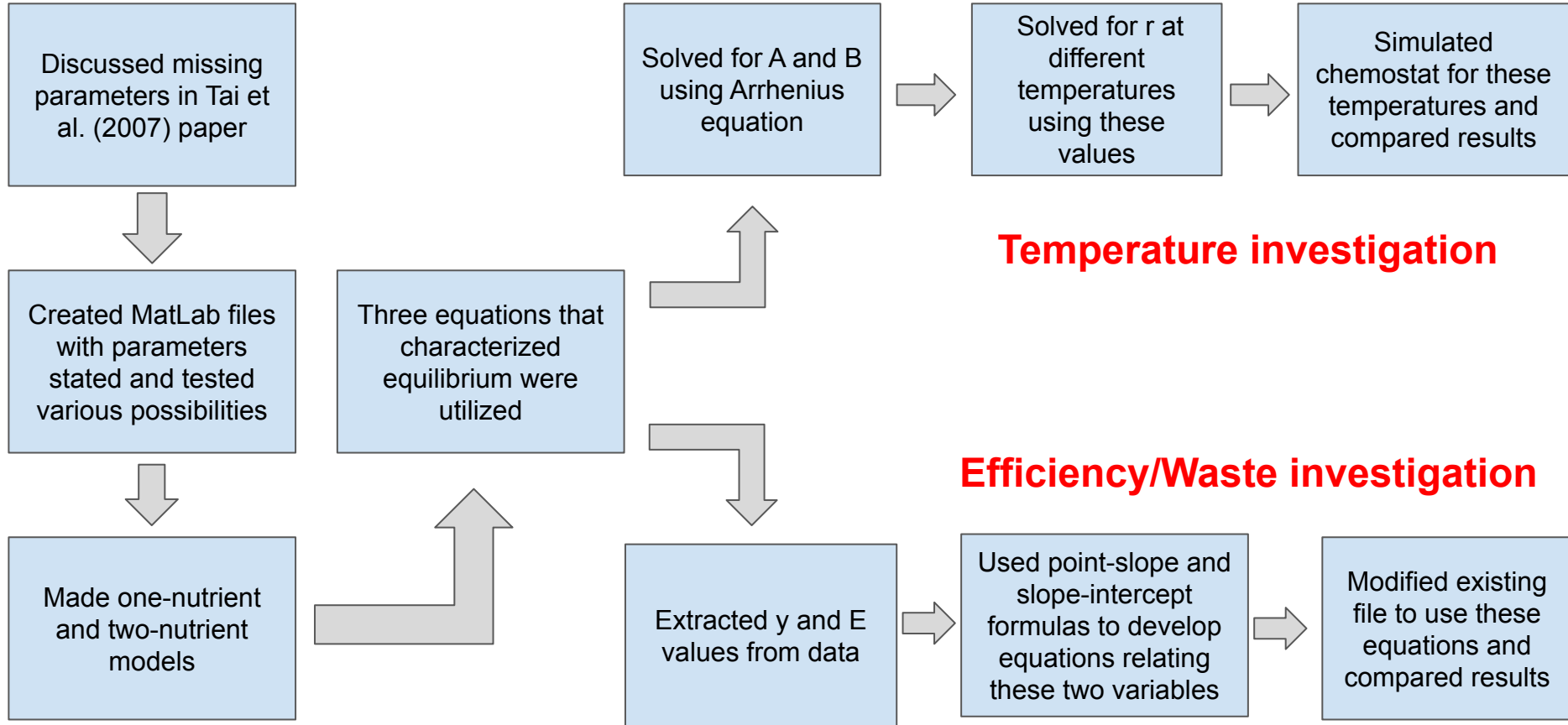
# Parameters from Data Used to Define Unknowns

	<b>12C Glucose Limiting</b>	<b>30C Glucose Limiting</b>	<b>12C Ammonium Limiting</b>	<b>30C Ammonium Limiting</b>
<b>T</b>	285K	303K	285K	303K
<b>r</b>	0.08	0.46	0.08	0.46
<b>A</b>	$4.9 \cdot 10^{11}$	$4.9 \cdot 10^{11}$	$4.9 \cdot 10^{11}$	$4.9 \cdot 10^{11}$
<b>B</b>	$6.98 \cdot 10^4$	$6.98 \cdot 10^4$	$6.98 \cdot 10^4$	$6.98 \cdot 10^4$
<b>E</b>	14.2857	14.2857	20.0000	25.000
<b>y</b>	0.504	0.054	16.216	15.333

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# Investigation of Chemostat Model





# Remaining Parameters Calculated from the following

**Equilibrium equations used in MATLAB:**

$$y(t) = \frac{dy}{dt} = \frac{-Exryz}{(y+k)(z+L)} - qy + qu$$

$$z(t) = \frac{dz}{dt} = \frac{-Fxryz}{(y+k)(z+L)} - qz + qv$$

$$x(t) = \frac{dx}{dt} = \frac{xryz}{(y+k)(z+L)} - qx$$

**Arrhenius Equation:**

$$K = Ae^{-Ea/RT}$$

**Point-slope Equation:**

$$y - y_1 = m(x - x_1)$$

**Slope-intercept Formula:**

$$y = mx + b$$

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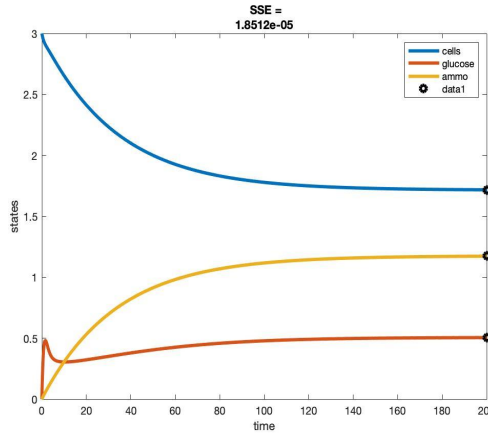
# Chemostat Simulated at Different Temperatures

- Used Glucose-limited condition
- Calculated rates using Arrhenius equation

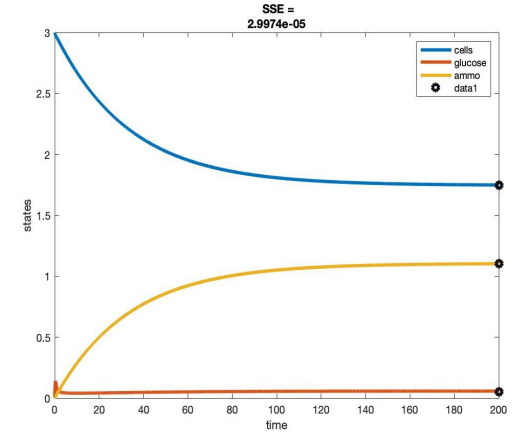
Temperature	Rate
15°	0.1087
20°	0.1787
25°	0.289

# Glucose Steady States Remained the Same

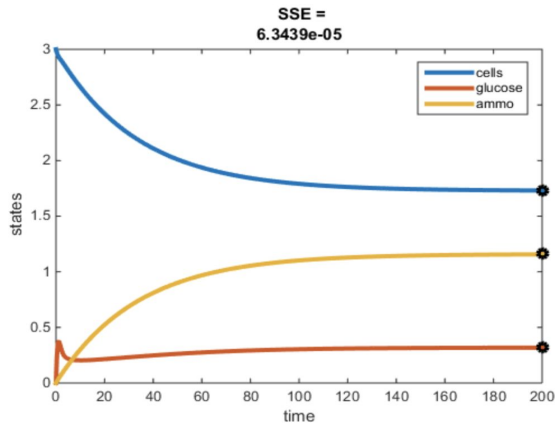
12 degrees Glucose limited



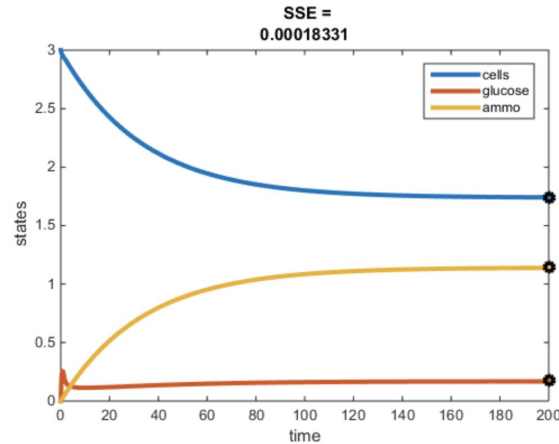
30 degrees Glucose limited



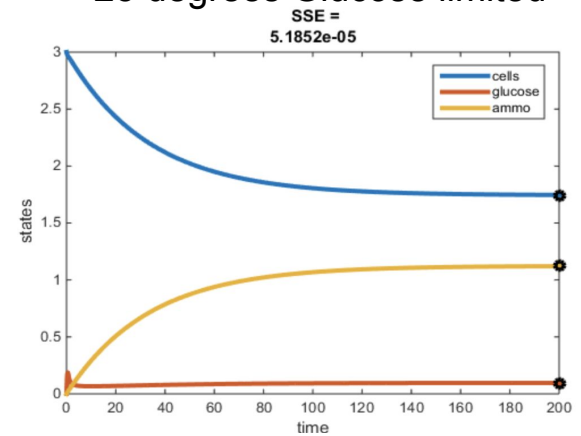
15 degrees Glucose limited



20 degrees Glucose limited



25 degrees Glucose limited



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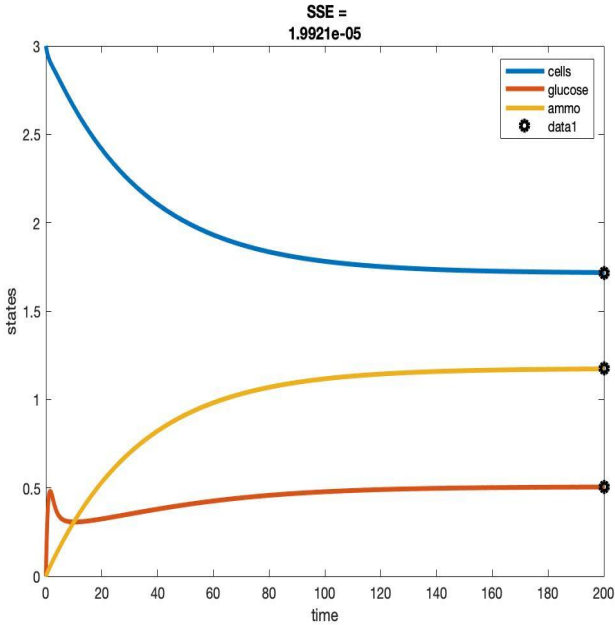
# Investigating E as a Variable Versus a Constant

- Calculated y values by multiplying residual glucose (rG) by the conversion factor to grams
- Used point-slope and slope-intercept formulas to develop equations that match both (y,E) points for each temperature

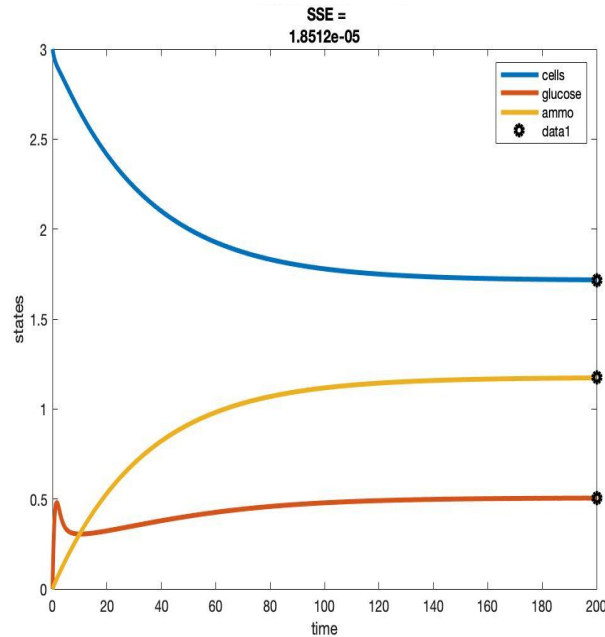
Temperature	Function
12°C	$E = 0.3637y + 14.1022$
30°C	$E = 0.7012y + 14.2483$

# Using the 12°C Equation Steady States Varied

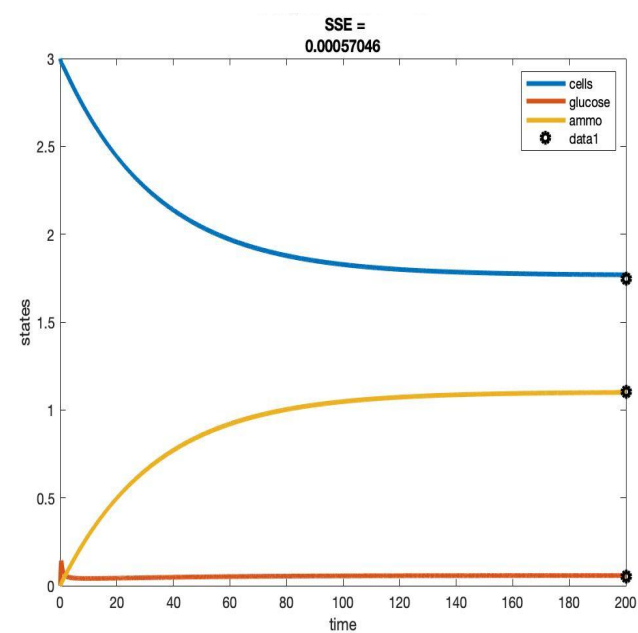
12 degrees Glucose limited adjusted



12 degrees Glucose limited



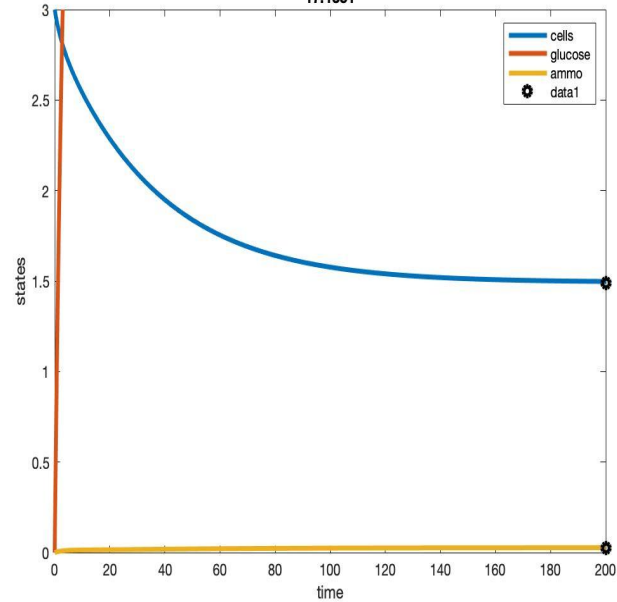
30 degrees Glucose limited adjusted



# The 30°C Equation Produced Similar Results

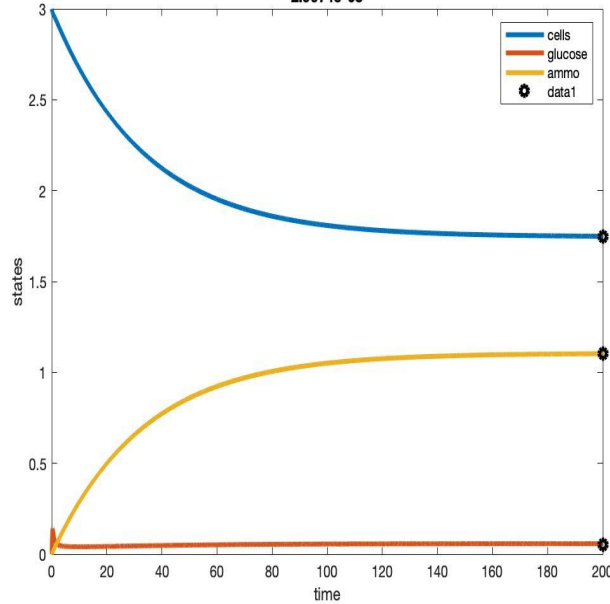
12 degrees Glucose  
limited adjusted

SSE =  
17.1351



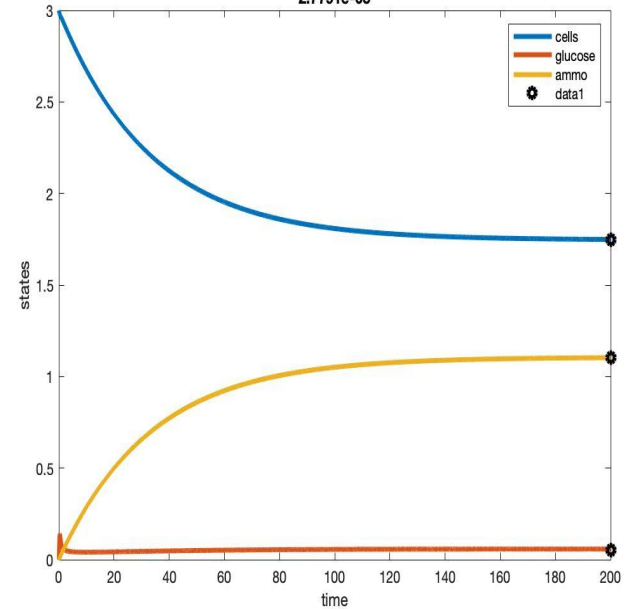
30 degrees Glucose  
limited

SSE =  
2.9974e-05



30 degrees Glucose  
limited adjusted

SSE =  
2.7791e-05





# Conclusions

- Changing the temperature did not affect the steady states
- Glucose levels decreased as temperature levels increased
- The Efficiency (E) did not change with temperature

# Future Directions

- Include temperatures outside of the 12 and 30 degree range and see if that makes a bigger difference than using the intermediary 15, 20, 25 degrees.
- Investigate a different nutrient's steady state over time, such as Trehalose.

# Acknowledgments

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**Kam D. Dahlquist, Ph.D.**



**Ben G. Fitzpatrick, Ph.D.**

**Dahlquist Lab**

# References

1. Dahlquist, K., & Fitzpatrick, B. (n.d.). BIOL388/S19: Week 14/15. Retrieved from [https://openwetware.org/wiki/BIOL388/S19:Week\\_14/15](https://openwetware.org/wiki/BIOL388/S19:Week_14/15)
2. Tai, S. L., Daran-Lapujade, P., Walsh, M. C., Pronk, J. T., & Daran, J. M. (2007). Acclimation of *Saccharomyces cerevisiae* to low temperature: a chemostat-based transcriptome analysis. *Molecular Biology of the Cell*, 18(12), 5100-5112. DOI: 10.1091/mbc.e07-02-0131
3. Tai, S. L., Boer, V. M., Daran-Lapujade, P., Walsh, M. C., de Winde, J. H., Daran, J. M., and Pronk, J. T. (2005). Two-dimensional transcriptome analysis in chemostat cultures: combinatorial effects of oxygen availability and macro- nutrient limitation in *Saccharomyces cerevisiae*. *J. Biol. Chem.* 280, 437–447.