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BS, Massachusetts Institute of Technology, June 2021 (Chemical-biological Engineering, Computer Science & Molecular Biology)

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Carolyn Bertozzi and K. Barr Sharpless chat about sharin the 2022 Nobel Prize in Chemistry



Bioorthogonal, click chemistry clinch the Nobel Prize October 5, 2022



Lithium mining's water use sparks bitter conflicts and novel chemistry



For John Goodenough's 100th birthday, Stereo Chemistry revisits a fan-favorite interview with the renowned scientist



TINY
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Wastinana

The sticky science of why we





The helium shortage wasn't supposed to March 24, 2022

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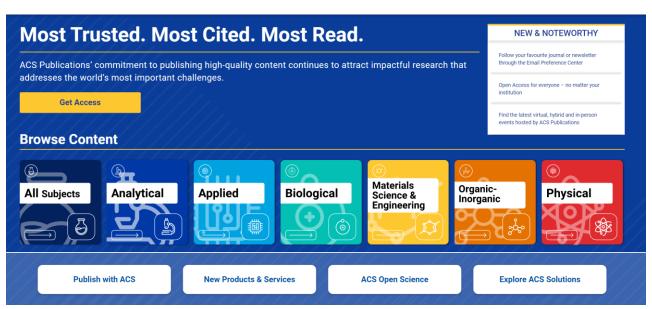


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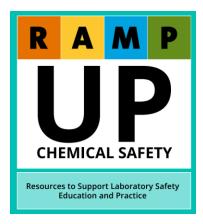
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A complete listing of ACS Safety Programs and Resources



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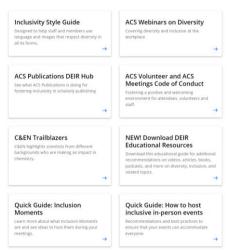


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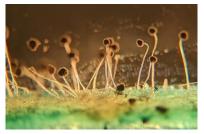
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Thursday, March 7, 2024 2-3pm ET
The Art of Self-Reinvention

Co-produced with the ACS Women Chemists Committee



Wednesday, March 13, 2024 | 11am-12:30pm ET Fungal Foes: Understanding the Challenges and Exploring New Treatment Options

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Powering the Future: The Latest

Battery Technologies

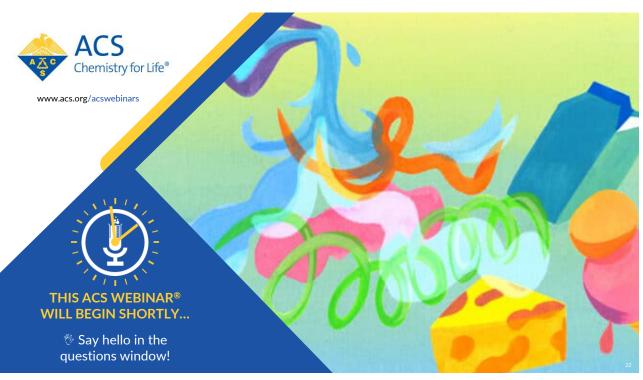
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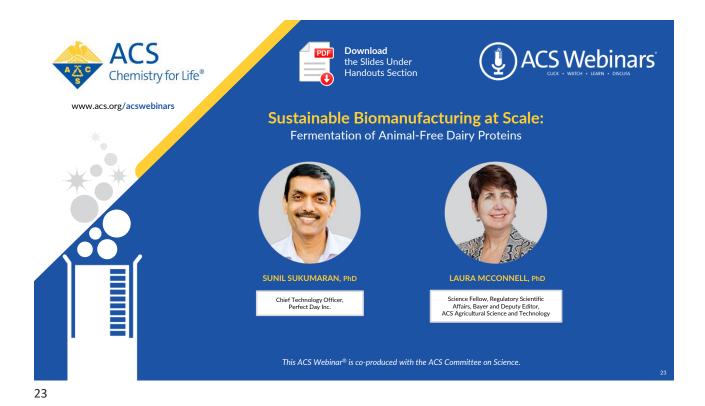
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- Identify and promote new frontiers of chemistry
- Examine scientific basis & formulate public policies related to chemical sciences

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ACS Committee on Science (COMSCI)



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Policy Statements

- ✓ Energy
- ✓ Sustainability
- ✓ Hydraulic Fracturing
- √ Forensic Science

Symposia at Fall ACS Meeting

Scaling New Heights of Chemistry Education with Artificial Intelligence Tools

Organized by: Robert Pribush, Judith Benham, Tom Holme, Mary Carroll

Elevating Atmospheric Chemistry Measurements and Modeling with Artificial Intelligence

Organized by: Carl Picconatto, Jeff Arnold, and Mary Carroll

<u>Awards</u>

- > National Medal of Science
- > National Medal of Technology and Innovation
- > Dreyfus Award in the Chemical Sciences

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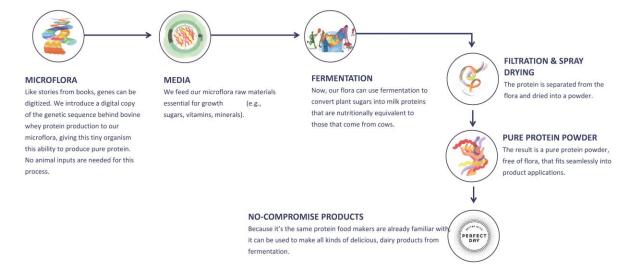
Sustainable Biomanufacturing at Scale: Fermentation of Animal-Free Dairy Proteins



All information and images are confidential and proprietary









*Compared to whey protein found in traditional milk. Data from ISO-certified, third-party-validated report.

All information and images are confidential and proprietary.

We're bringing higher quality, sustainable protein to the category.

Protein Source	BCAA (g/100g powder)	Function	Animal Free	Lactose Free	Soy Free	GHG Emission: (per 100g protein)
Dairy (WPI)	20	***	×	8	Ø	9.50
Pea	11	★☆☆				0.44
Oat	7	★★☆				2.70
Almond	N/A	★益益				0.26
Soy	9	***			×	1.98
Perfect Day	23	***	Ø	Ø	Ø	0.29

Acronyms: WPI (whey protein isolate), BCAA (branched-chain amino acids), GHG (greenhouse gas)

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The below brands, and many more, are now in over 5K grocery stores around the globe



BEL GROUP Cream Cheese



Ice Cream



NICK'S Ice Cream



RENEWAL MILL **Bakery Mix**



BORED COW Milk



Milk



MYPROTEIN **Sports Nutrition**



Natreve Sports Nutrition

Partners in **Testing**







Barista Milk + Ice Cream



MARS Chocolate

Sca (g/100g powder): Gorissen et al., 2008, "Protein content and amino acid composition of commercially available plant-based protein isolates", https://doi.org/10.100782F500726-018-2640-5
2. GHG Emissions per 100g Protein (Poore & Nemecek, 2018), "Plant Proteins: Assessing Their Nutritional Quality and Effects on Health and Physical Function," Hertzler, Lieblein-Boff, Weller, Allgeler, Gras Notice for Partilly Defatted Almond Protein Flour, Blue Glimmod.

Partilly Defatted Almond Protein Flour, Blue Glimmod.

Global Footprint



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The State of Our Industry **Opportunities** Challenges High consumer demand for Scale required to meet demand sustainable products outpacing available infrastructure Global market for products and Geographic expansion requires varied cross-border appeal regulatory approval processes Requires proactive policy engagement by Government interest is high and potential funding available existing leaders Path to price parity is proven Largely price intolerant during and achievable period of scale-up Common sustainability goal Requires pushback against narrative drives deep collaboration of disruption as success

Growing Consumer Demand

Research shows 90 million consumers will be interested in precision fermentation food and beverages when they understand its benefits

	F	Ready	Easily			nced with enefits	Unco	nvinced	Curren	tly out of reach	Total across groups in eacl age cohort
Gen Z (18-25)	14%	4,440,553	19%	6,379,001	14%	4,439,579	39%	12,625,547	15%	4,896,658	100%
Millennials (26-41)	27%	18,911,267	15%	10,050,526	13%	8,887,815	27%	18,672,987	18%	12,331,595	100%
Gen X (42-57)	16%	10,074,889	8%	4,830,936	14%	9,015,227	37%	23,099,536	25%	15,697,681	100%
Boomers (58-73)	3%	1,899,418	6%	3,698,010	14%	7,919,066	45%	25,960,005	32%	18,299,525	100%
Total in each segment	35.326.127		24,958,473		30,261,687		80,358,074		51,225,459		
Total addressable market today			willi	546,287 ng to buy ODAY							
•				~							

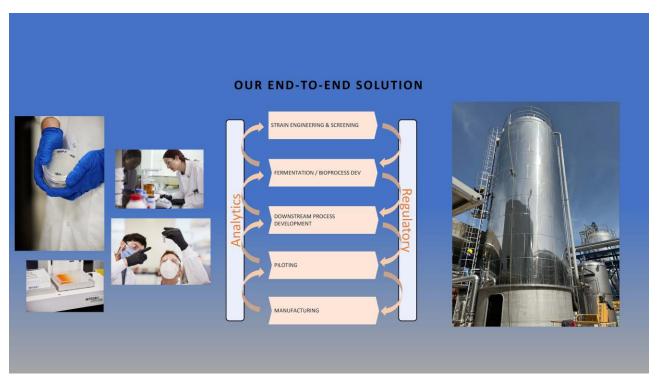
Data from The Hartman Group for Perfect Day & Cargill

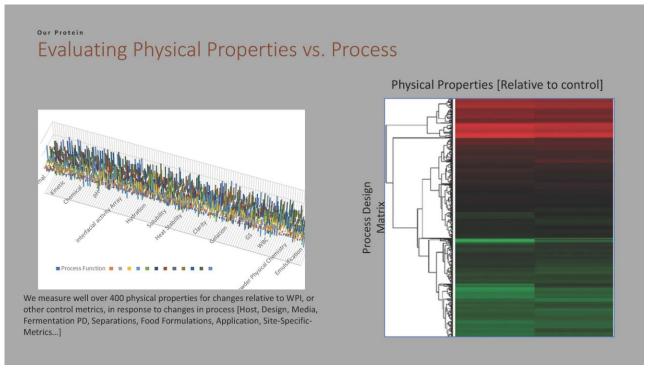
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We've built our business to meet demand and scale our

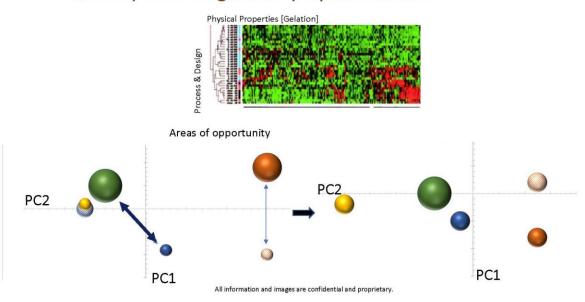




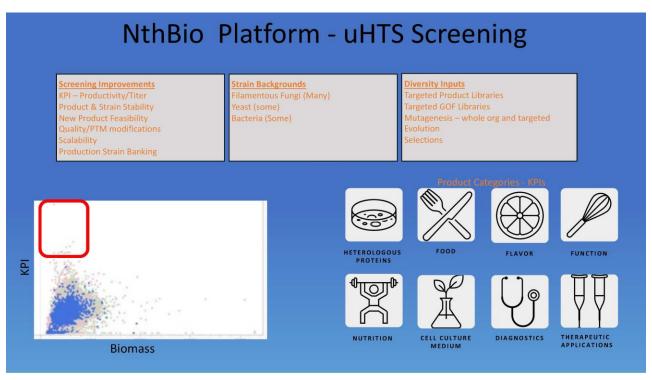




Identifying the Process Functions that matter for Quality and Regulatory Specification

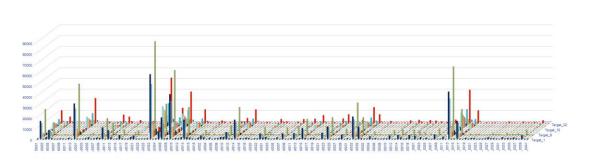


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Analysis of Hits

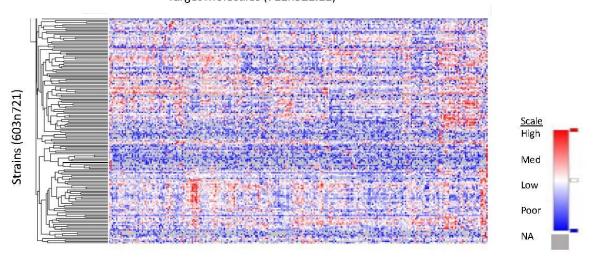




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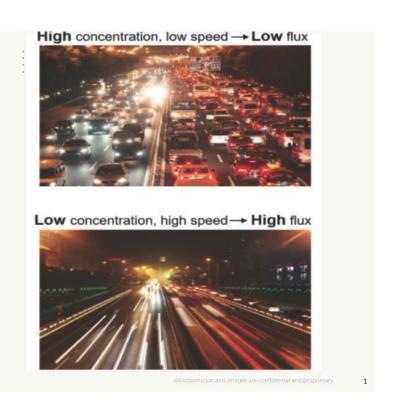
Large Scale Analysis - Strains vs. Product Expression

Target Molecules (711n321.11)







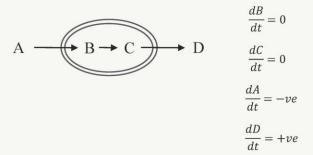


Why should we carry out metabolic flux analysis

- · To quantify the carbon flux distribution
- · To understand the complex interplay between energy metabolism, carbon fixation, and assimilation pathways
- · Connects genomics with metabolism

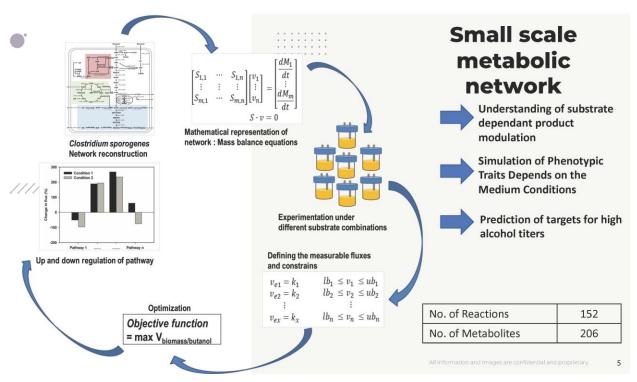
Input Carbon source Nitrogen source Biomass H2 CO2

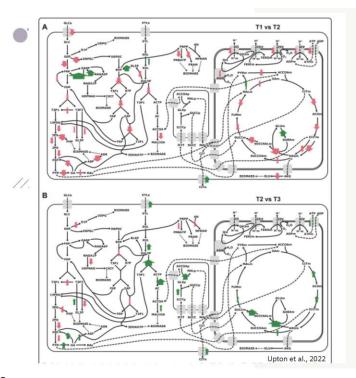
Theory: Pseudo steady state Approximation



 $v = flux (rate \ of \ reaction) = \frac{mmole \ of \ substrate \ consumed \ or \ Product \ formed}{g \ biomass \ * hour}$

All information and images are confidential and proprietary.





Integration of A. niger transcriptomic profile with metabolic model identifies potential targets to optimise citric acid production from lignocellulosic hydrolysate



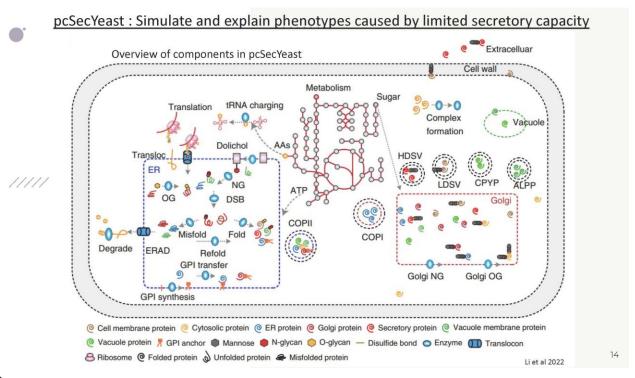
Prediction of targets for higher citric acid titers

T1: Glucose consumption phase before the onset of citric acid production and phosphate-limited growth.

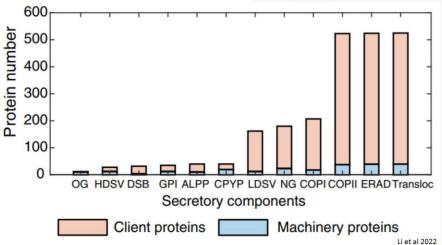
T2: Glucose consumption phase, citric acid producing, phosphate-limited growth.

T3: Xylose consumption phase after glucose was fully consumed. Citric acid producing and phosphate-limited growth.

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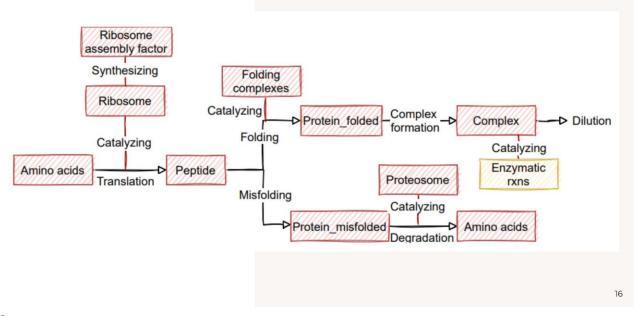
Subsystems in the secretory pathway and the protein number in each subsystem



- 1. OG: O-glycosylation
- 2. HDSV : high-density secretory vesicles
- 3. DSB: disulfide bond formation
- 4. GPI: glycosylphosphatidylinositol
- 5. ALPP: alkaline phosphatase pathway
- 6. CPYP: carboxypeptidase Y pathway
- 7. LDSV: low-density secretory vesicles
- 8. NG: N-glycosylation
- 9. COPI: Coat Protein Complex I
- 10. COPII: Coat Protein Complex II
- 11. ERAD: ER-associated degradation
- 12. Transloc: translocation

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Flowchart of the protein related process in the pcSecYeast.







- \checkmark Reactions for production of post-translational modification precursors
- ✓ Reactions catalyzed by isozymes were also split into multiple identical reactions with various isozymes. This step was performed to facilitate later kcat match and enzyme constraining step
- ✓ Translation initiation, elongation, and termination reactions added for each protein (total 1639)
- Protein translocation pathways added: co-translational translocation, post-translational translocation, and post-translational translocation-tail-targeting

Co-translational translocation

- 1. signal peptide recognition
- 2. ER receptor biding to peptide-SRPC
- 3. binding of peptide -SRPC-SRC to the translocator (Sec61C) $\,$
- 4. binding of peptide -SRPC-SRC to the translocator (Ssh1C)
- 5. signal peptidase
- 6. export the signal peptide out of ER for degradation

Post-translational translocation

- 1. exit the ribosome
- 2. bind to the cytosolic chaperone
- 3. Translocation
- 4. pulling of nascent protein

*The coefficient of ATP in step 4 was set as length/40, since the ATP molecule bound to the chaperone Kar2, is assumed to be hydrolyzed to ADP for every 40 amino acids that pass through the translocon pore

Post-translational translocationtail targeting

- 1. load the TA proteins
- 2. bind to Get3
- 3. bind to ER receptor

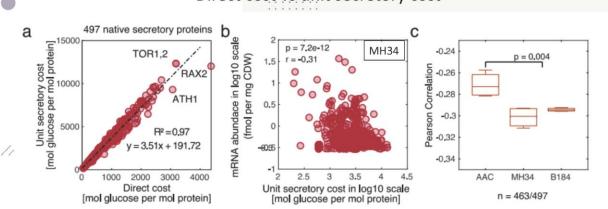
Reactions for complex formation for enzymes used in either of these pathways are added to the model.

Li et al 2022

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Direct cost vs unit secretory cost



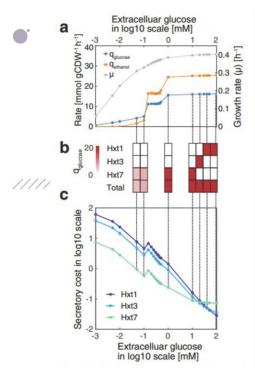
Direct cost: Includes the energetic cost for synthesis, modification and secretion of this protein **Unit secretory cost:** Direct cost + cost for the corresponding increased fraction of the catalytic machineries in these processes caused by the increase of this protein

Total no of proteins: 1639 Metabolic proteins: 1156 Secretory proteins: 483 Secretory $cost_i = unit secretory <math>cost_i \cdot [E_i] = unit secretory cost_i \cdot \frac{V_{glc.total}}{k_{cat.i}} \frac{|S|}{|S|+K_{u.t.}}$

AAC: low yield $\alpha\text{-amylase}$ strain MH34 and B184: high yield $\alpha\text{-amylase}$ strain

Li et al 2022

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Simulated physiological response of *S. cerevisiae* as a function of the extracellular glucose concentration

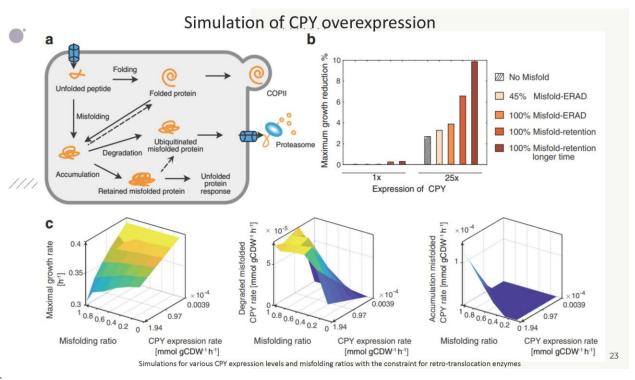
Hxt 1 and Hxt 3: Low affinity transporters : Low unit secretory cost

Hxt 7: High affinity transporters : High unit secretory cost



Li et al 2022 21

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-	_							Overview of protein
	a Protein	abbr.	DSB	NG	OG	GPI	Length	features for eight
	Insulin precursor	IP	3	0	0	0	53	recombinant proteins
	Human granulocyte colony stimulating factor	hGCSF	2	0	1	0	174	produced by S. cerevisiae
	Hemoglobin	Hemoglobin	0	0	0	0	299	
	β-glucosidase	BGL	0	0	0	0	421	
	α-amylase	a-amylase	4	1	0	0	478	
	Acid phosphatase	PHO	8	9	0	0	435	
	Human serum albumin	HSA	17	1	0	0	585	
111,	Human transferin	HTF	19	0	1	0	679	
* * * *	b 4×10 ⁻³	с	×1e-5					
	Protein production rate [mmol gCDW** h*] Output Specific growth rate [mmol gCDW** h*]	0.3 0.4 te [h·1]	8 · 6 · 4 · 2 · 50 · 50 · 6 · 6 · 6 · 6 · 6 · 6 · 6 · 6 · 6 ·	n≅x≷n	-04¤		ative impact	

Need of the hour?

- o Scale down models ?
- ////o In-depth appreciation of stage specific Physiology
 - o A relevant mathematical model with relevant constraints put in place
 - Alternate feed-stock
 - Recycling resources

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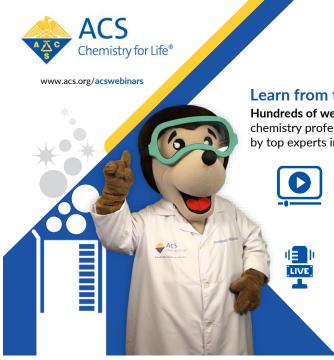
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