

Rumen-Derived Next Generation Probiotics for Sustainable Bioplastic Production and Microplastic Toxicity Mitigation

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Curriculum Vitae

- ▶ 2019~현재 서울대학교 농생명공학부 동물생명공학전공 교수
- ▶ 2025~2026 University of Florida Visiting Professor
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- ▶ 2022~현재 한국축산학회 영문지 JAST 편집위원장
- ▶ 2021~2024 한국연구재단 생명과학단 기반생명 동물자원학분야 전문위원



Rumen-Derived Next Generation Probiotics for Sustainable Bioplastic Production and Microplastic Toxicity Mitigation

Seoul National University
Department of Agricultural Biotechnology
Younghoon Kim

Source: Tanaka et al. (2022) Polymers, 14(15):3063

1 | **Background**

2 | **Results (1)**
- PHA producing bacteria

3 | **Results (2)**
- Microplastic toxicity

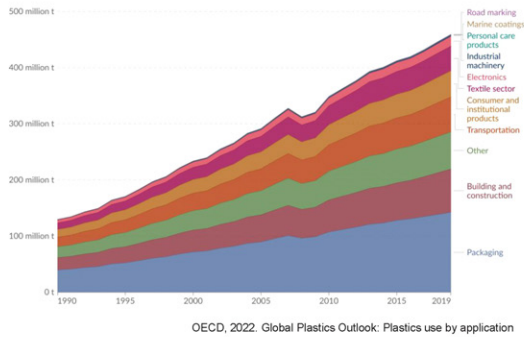
4 | **Further study**

01

Background

Current problems of petroleum-based plastics

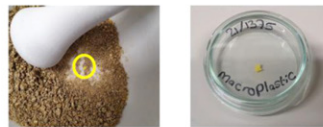
Global primary plastic production
By industrial sector, 1990 to 2019



Occurrence and exposure of
microplastics



Walker, Tony R. and Lexi Fequet (2023)



Van-der Veen et al., 2022

Problems of petroleum-based plastics



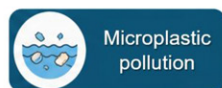
Fossil fuel dependency



Low recyclability

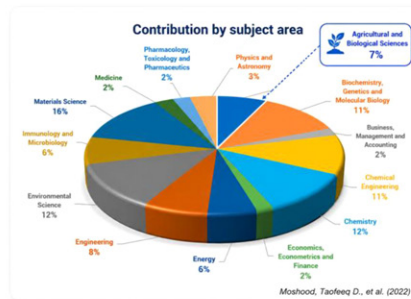
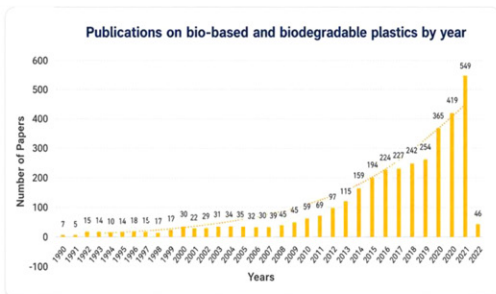


Long-term persistence

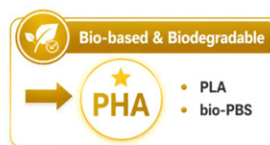
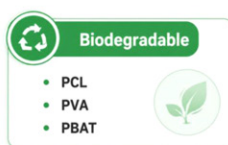
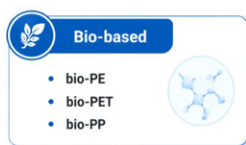


Microplastic pollution

Bioplastics: Plastics that are bio-based, biodegradable, or both

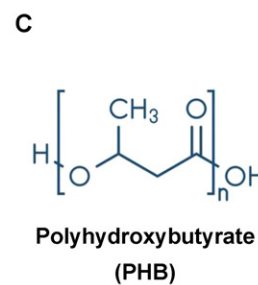
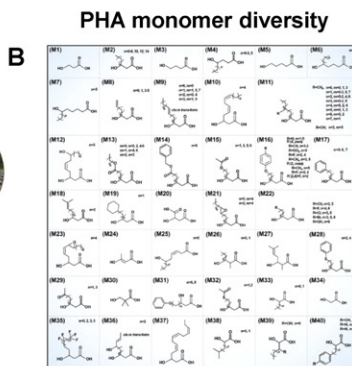
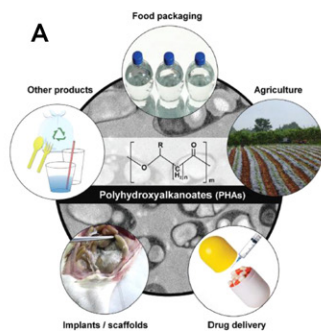


Three types of bioplastics



5

PHAs for sustainable bioplastics



- Microbial polyesters accumulated as intracellular storage granules
- Bio-based and biodegradable bioplastics
- Monomer composition determines polymer properties
- Representative polyhydroxyalkanoates (PHAs) type: PHB, PHBV

6

Discovery of PHA producing bacteria

RESEARCH

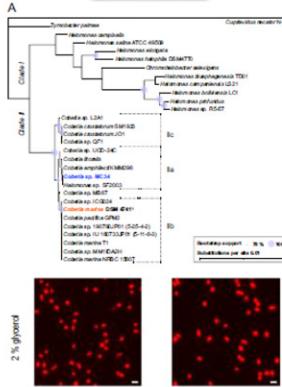
Open Access

High natural PHA production from acetate in *Cobetia* sp. MC34 and *Cobetia marina* DSM 4741^T and in silico analyses of the genus specific PhaC₂ polymerase variant

Mikkel Christensen¹, Piotr Jablonski², Bjørn Altemark³, Knut Tjørum⁴ and Hilde Hansen⁵*

Microbial Cell Factories (2021)

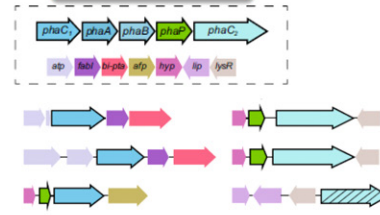
Isolation



Carbon source screening

Strain	Carbon substrate	CO ₂ at harvest	CDW (g/L)	PHB (% of CDW)	PHV (% of CDW)	PHA (g/L)
<i>Cobetia</i> sp. MC34	Acetate	15.0 (± 1.3)	3.4 (± 0.5)	72 (± 11)	–	2.5
	Acetate + valerate	14.2 (± 2.1)	3.4 (± 0.8)	48 (± 3.4)	14 (± 2.3)	2.1
	Glycerol	6.2 (± 1.4)	2.5 (± 0.5)	26 (± 1.2)	–	0.7
	Glucose	6.5 (± 0.3)	1.8 (± 0.1)	35 (± 4.8)	–	0.6
	Fructose	6.0 (± 0.8)	3.9 (± 0.8)	9.9 (± 1.3)	–	0.4
<i>Cobetia marina</i> DSM 4741 ^T	Acetate	15.0 (± 2.9)	3.9 (± 0.6)	61 (± 8.3)	–	2.4
	Acetate + valerate	17.0 (± 1.4)	4.4 (± 0.3)	59 (± 11)	26 (± 2.6)	3.7
	Glycerol	14.0 (± 2.1)	4.0 (± 0.1)	61 (± 8.1)	–	2.5
	Glucose	8.0 (± 0.2)	2.4 (± 0.3)	46 (± 2.9)	–	1.1
	Fructose	8.1 (± 0.2)	3.1 (± 0.2)	28 (± 1.1)	–	0.9

Genome analysis



- ❖ *Cobetia* sp. MC34: 2.5 g/L PHB from acetate
- ❖ *Cobetia marina* DSM 4741^T: 2.5g/L PHB from glycerol
- ❖ Wild type bacteria can show strong PHB accumulation

7

Large-scale production

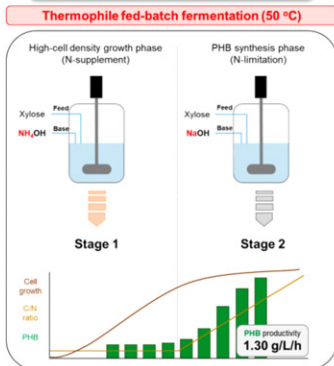
Production of polyhydroxybutyrate with high cell density cultivation using thermophile *Caldimonas thermodepolymerans*

Jun Won Jang^{1,2}, In Yeub Hwang^{1,2}, Ok Kyung Lee³, Eun Yeol Lee^{1,2}*

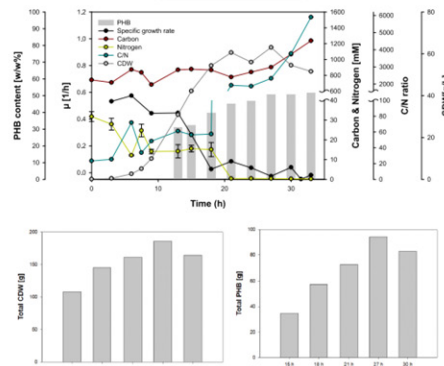
¹ Department of Chemical Engineering (BK21 FOUR Integrated Engineering Program) College of Engineering, Kyung Hee University, Republic of Korea

Bioresource Technology (2024)

Large-scale fermentation



Optimization



- ❖ Nitrogen limitation promoted PHB accumulation (31.9 g/L with 1.30 g PHB/L/h).
- ❖ Thermophilic PHB production may reduce contamination risk.

8

Metabolic engineering

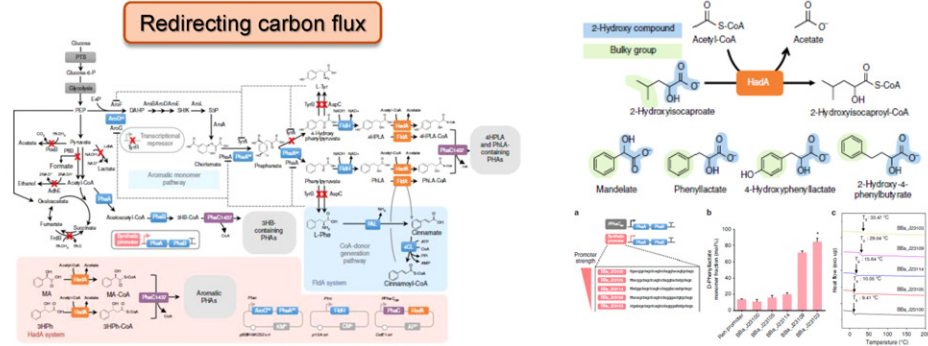
ARTICLE

DOI: 10.1038/s41467-017-03498-w OPEN

One-step fermentative production of aromatic polyesters from glucose by metabolically engineered *Escherichia coli* strains

Jung Eun Yang¹, Si Jae Park², Won Jun Kim¹, Hyeon Jun Kim¹, Bumjoon J. Kim¹, Hyuk Lee⁴, Jihoon Shin⁵ & Sang Yup Lee^{1,6,7}

Nature communications (2018)



- ❖ Engineered *E. coli* produced poly(3HB-co-D-phenyllactate) from glucose.
- ❖ Gene deletions redirected carbon flux toward aromatic monomer production.
- ❖ Expression of HadA evolved PHA synthase, and PhaAB enabled 3HB-containing aromatic polyester synthesis.

9

Industrial by-product upcycling for PHB production

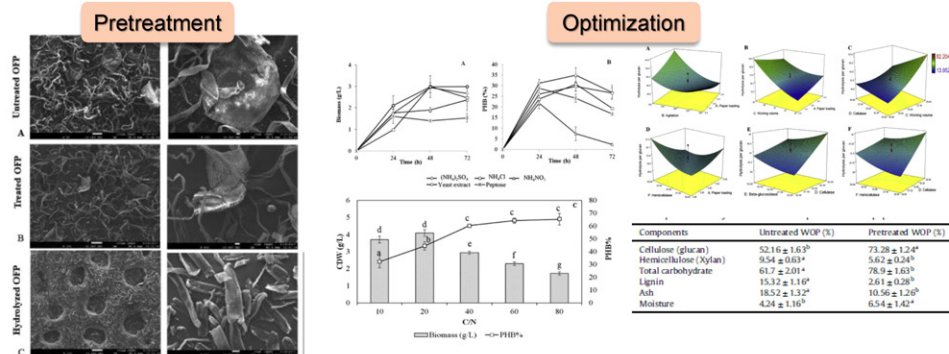
Production of bioplastic (poly-3-hydroxybutyrate) using waste paper as a feedstock: Optimization of enzymatic hydrolysis and fermentation employing *Burkholderia sacchari*



Journal of Cleaner Production (2018)

Huda Al-Battashi¹, Neelamegam Annamalai², Shatha Al-Kindi², Anu Sadasivan Nair², Saif Al-Bahry², Jay Prakash Verma², Nallusamy Sivakumar^{2*}

¹ Department of Biology, College of Science, Sultan Qaboos University, PO Box 36, PC 123, Oman
² Hawkebury Institute for Environment, Western Sydney University, Hawkebury Campus, Penrith, Sydney, New South Wales, Australia



- ❖ Converting waste office paper into a fermentable carbon source of PHB production.
- ❖ 92.1% sugar utilization with 1.60 g/L PHB.
- ❖ Waste upcycling can reduce feedstock cost.

10

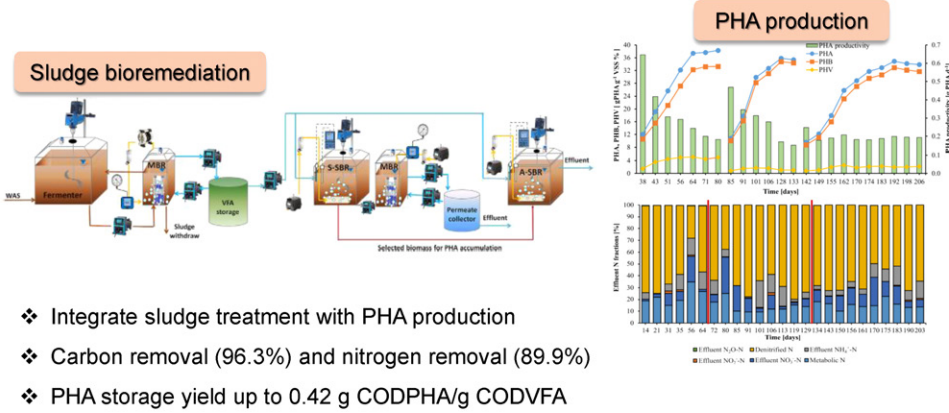
Bioremediation of activated sludge

From waste activated sludge to polyhydroxyalkanoate: Insights from a membrane-based enrichment process

Antonio Mineo^a, Mark M.C. van Loosdrecht^b, Giorgio Mannina^{a,*}

^a Engineering Department, Palermo University, Viale delle Scienze ed. 9 90129 Palermo, Italy

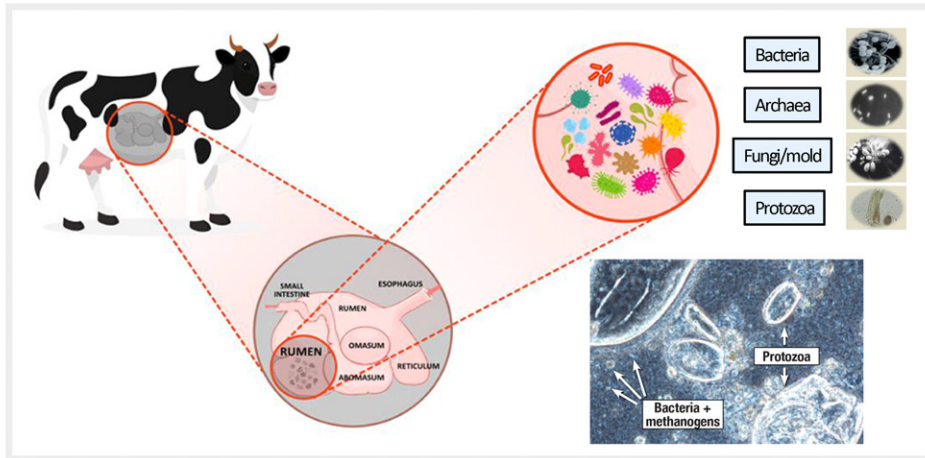
^b Department of Biotechnology, Delft University of Technology, Van der Maasweg 9 2629 HZ Delft, The Netherlands *Chemical Engineering Journal (2025)*



11

Rumen microbiome

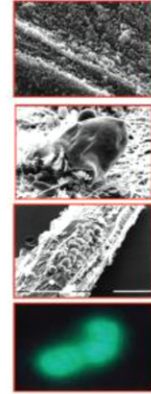
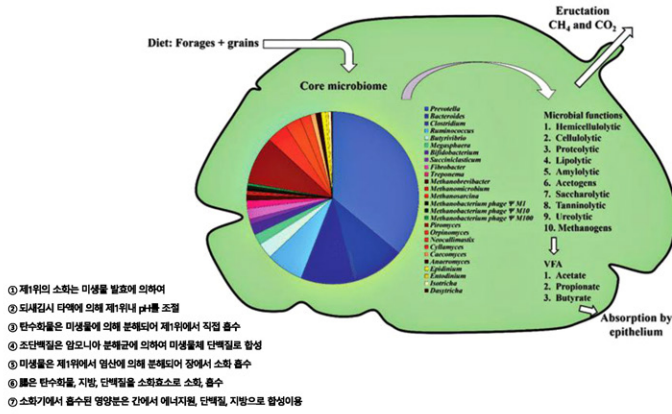
Biodiversity of Rumen Microbiome



Rumen Microbiome: Reservoir of uncultivable microorganisms and new functional biological enzyme resources

Rumen microbiome

Biodiversity of Rumen Microbiome

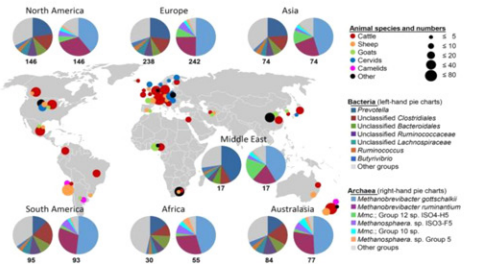


Diet composition directly influences the rumen microbiome structure, which affects microbial functions and consequently biomass degradation, resulting in the release of methane (CH₄) and carbon dioxide (CO₂) via eructation and volatile fatty acids that are absorbed by the epithelium abomasum.

- ① 제1위 소의는 미생물 발효에 의하여
- ② 되새김시 타액에 의해 제1위내 pH를 조절
- ③ 탄수화물은 미생물에 의해 분해되어 제1위에서 직접 흡수
- ④ 조단백질은 알모니아 분해에 의해 미생물에 단백질로 합성
- ⑤ 미생물은 제1위에서 암산에 의해 분해되어 장에서 소화 흡수
- ⑥ 높은 탄수화물, 지방, 단백질을 소화효소로 소화 흡수
- ⑦ 소화기에서 흡수된 영양분은 간에서 에너지원, 단백질, 지방으로 합성 이용

Rumen microbiome

Biodiversity of Rumen Microbiome



Henderson et al. 2015

nature biotechnology

Cultivation and sequencing of rumen microbiome members from the Hungate1000 Collection

2015, Volume 33, Number 10, October 1, 2015

RMG NETWORK

Global Rumen Census

Discover the diversity of rumen microbiomes across different regions and animal species.

Carbon neutrality and rumen microbiome

Science Advances

A heritable subset of the core rumen microbiome dictates dairy cow productivity and emissions

2017, Volume 12, Number 10, October 13, 2017

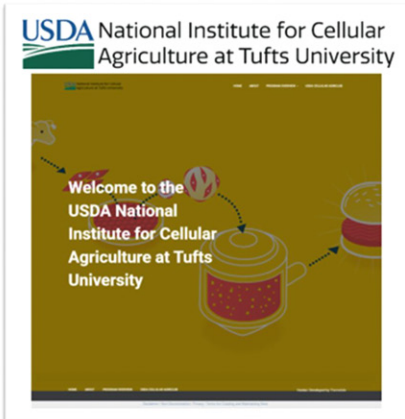
방귀 적게 쐬는 젖소, 장내미생물로 글라 메탄가스 줄인다

MYTHERISA

Discover the power of rumen microbiomes in improving dairy cow productivity and reducing methane emissions.

Future strategy for the animal science and industry that improves productivity and achieves carbon neutrality through the discovery of highly functional microbiome derived from rumen fluids

Rapid growth of cell technology for cultured meat



USDA \$10 million investment (2021)

Lab-Grown Meat's Carbon Footprint Potentially Worse Than Retail Beef

Study Finds Scaling Up Production Using Existing Processes Highly Energy-Intensive

By Amy Quinlan | May 22, 2023

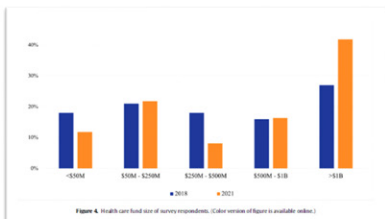


State of California \$5 million investment (2022)

Raszap Skorbiansky, S., McFadden, J., & Saavoss, M. (2024). The Economics of Cellular Agriculture.

Cell culture spent medium from cultured meat production

Cell culture industry



1 ton of cultivated-meat
= 10,000 L of culture medium

Generation of the spent media



Identified compound	Quantity of Formula	Spent medium (%) From formula
2-Ethoxyethanol acid	C ₈ H ₁₆ O ₃	1
2-Propylene	C ₃ H ₆ O	1
Dehydrogen	C ₁₀ H ₁₆ O	1
Water	C ₁ H ₂ O	1
Carbon	C ₁ H ₀ O	1
Hydrogen sulfide	C ₁ H ₂ S ₂ O	1
Hydroxylamine	C ₁ H ₃ O ₂ N	1
Glutamine	C ₅ H ₉ O ₂ N	1
2-Pyridoxaldehyde	C ₆ H ₇ O ₂ N	1
1-Aminoethanol	C ₂ H ₅ O ₂ N	1
2-Hydroxyethanol acid	C ₂ H ₄ O ₃	1
4-Hydroxy-2-pyridone (2-Pyridone)	C ₅ H ₅ O ₂ N	1
Acetyl-L-cysteine	C ₄ H ₇ O ₂ S	1
Cysteine	C ₃ H ₇ O ₂ S	1
Cysteine	C ₃ H ₇ O ₂ S	1
Sodium 1-carboxylate	C ₁ H ₃ O ₂ N	1
2-oxoethanol	C ₂ H ₃ O ₂	1
Water (hydroxylamine)	C ₁ H ₃ O ₂ N	1
L-homoserine	C ₄ H ₇ O ₂ N	1
L-homoserine	C ₄ H ₇ O ₂ N	1
L-homoserine	C ₄ H ₇ O ₂ N	1

Lactate, proteins, metabolites, ammonium

PHB production

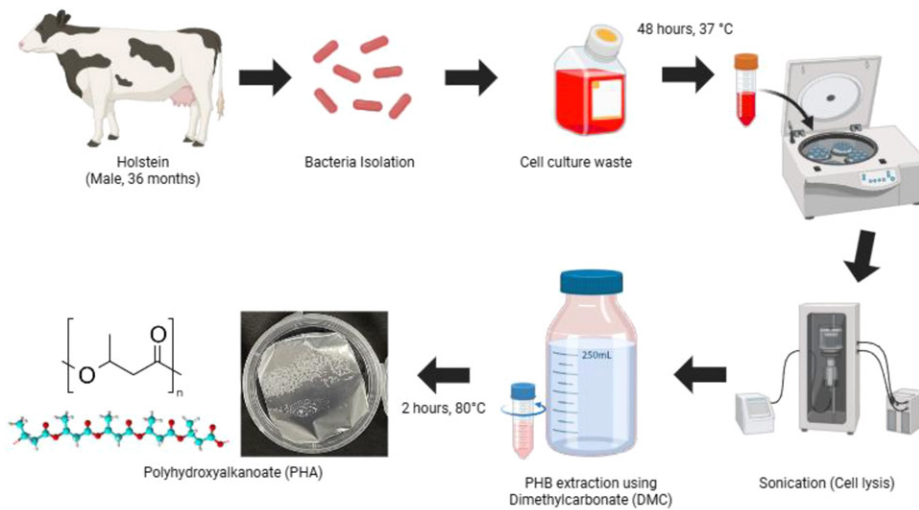
Remediation

Carbon valorization

Bioplastic

Circular economy

Bioconversion of cell culture spent medium



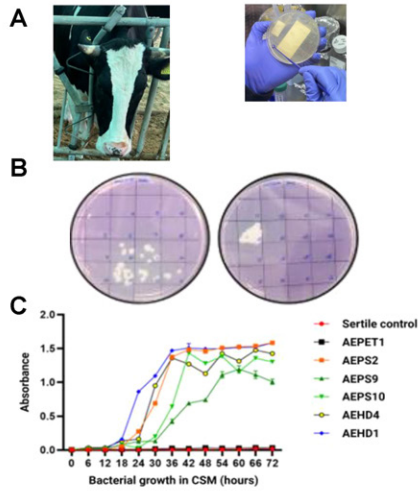
02

Results (1)

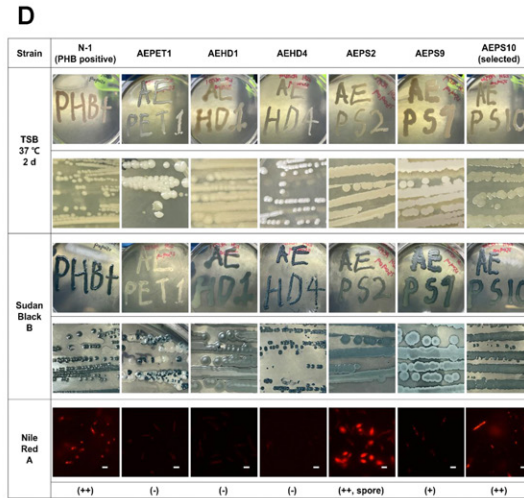
- Isolation and Characterization of PHA producing bacteria from rumen fluids

Bacterial isolation for PHA producing bacteria

Isolation of CSM-growing bacteria



Screening of PHA-producing bacteria

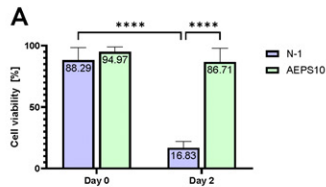


- ❖ Among 60 rumen fluid-derived strains, 6 isolates formed visible colonies.
- ❖ AEPS10 was selected for further analysis

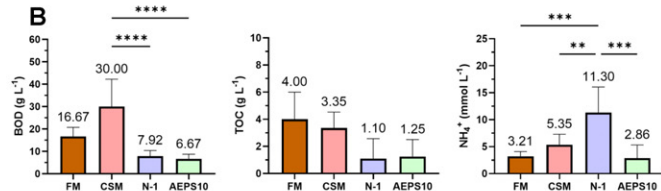
19

Bioremediation and polymer production

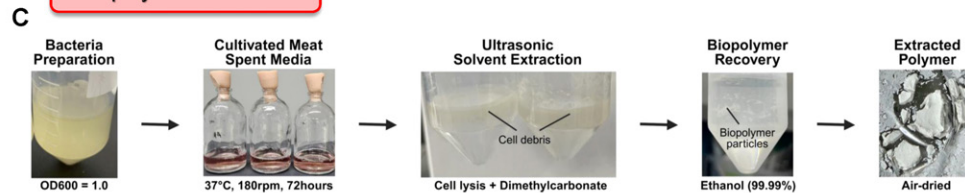
Cell viability



Bioremediation



Biopolymer extraction

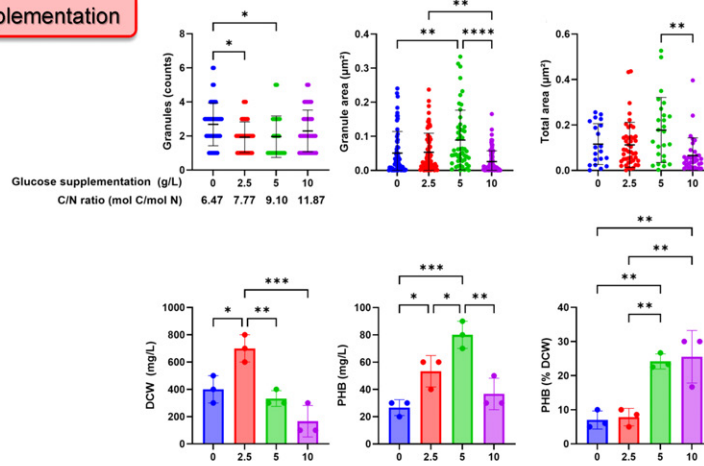


- ❖ Biochemical oxygen demand (- 77.8%)
- ❖ Total organic carbon (- 62.7%)
- ❖ Ammonium (- 46.5%)

20

Dynamics of PHB production

Carbon supplementation

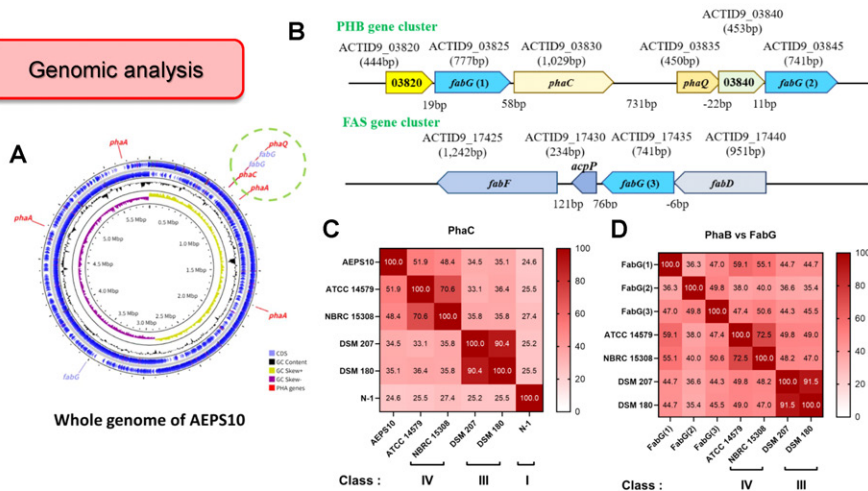


- ❖ Increasing the C/N ratio promoted PHB accumulation in AEPS10.
- ❖ The 5 g/L glucose condition showed the highest PHB production.
- ❖ PHB granules became more distinct under carbon-supplemented CSM.
- ❖ Carbon availability strongly influenced PHB storage capacity.

23

Molecular characterization of PHB producing bacteria using WGS

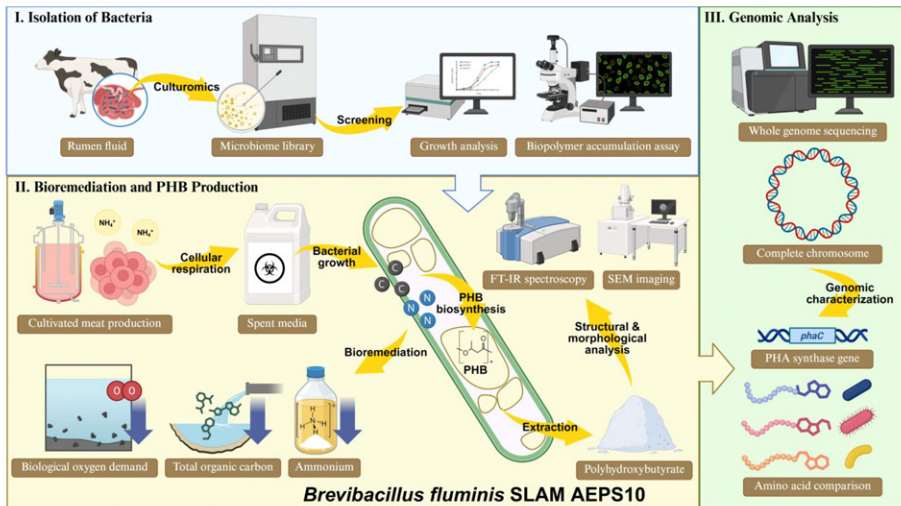
Genomic analysis



- ❖ AEPS10 carried a Class IV-like *phaC* region.
- ❖ PHB-related enzymes showed similarity to Bacillus-type PHA systems.
- ❖ Conserved enzyme identity supported PHB biosynthetic potential.
- ❖ Gene organization suggested strain-specific PHB regulation.

24

Molecular characterization of PHB producing bacteria



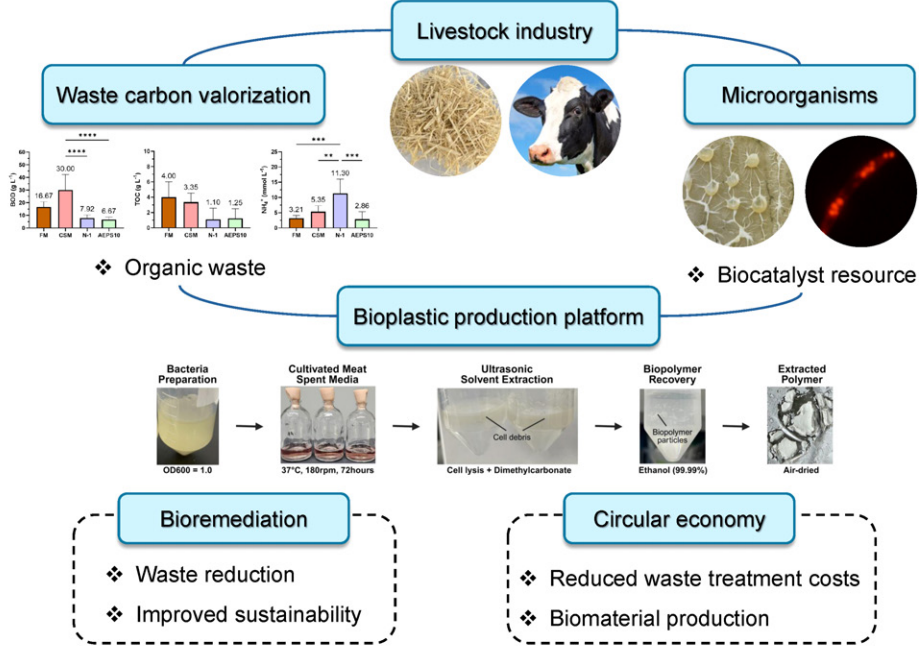
- ❖ This study connects waste treatment and bioplastic production in one bacterial process.
- ❖ AEPS10 reduced BOD, TOC, and ammonium in CSM and accumulated recoverable PHB.
- ❖ This strain provides a starting point for applying animal-associated microbial resources to waste-carbon conversion.

25

04

Further study

“Enbiotics” for microbe-based sustainable animal production



43

“Culturomics” for Unknown but Functional microbes in Rumen fluid

Gut microbiome is important role on the production and robustness of livestock as well as health function of companion animal

- It is necessary to secure live and active gut microbiome for applying to animal industry using **culturomics**, not just observation of microbial change with metagenomics

16S rRNA gene phylogeny

Phylogenetic tree showing various bacterial species and their relative abundance.

Species list:

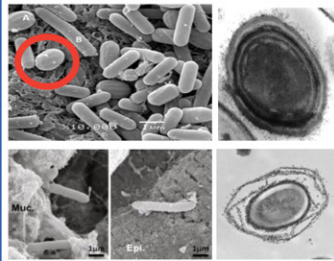
- OTU1**: *Streptococcus* sp. (100%)
- OTU2**: *Streptococcus* sp. (100%)
- OTU3**: *Streptococcus* sp. (100%)
- OTU4**: *Streptococcus* sp. (100%)
- OTU5**: *Streptococcus* sp. (100%)
- OTU6**: *Streptococcus* sp. (100%)
- OTU7**: *Streptococcus* sp. (100%)
- OTU8**: *Streptococcus* sp. (100%)
- OTU9**: *Streptococcus* sp. (100%)
- OTU10**: *Streptococcus* sp. (100%)
- OTU11**: *Streptococcus* sp. (100%)
- OTU12**: *Streptococcus* sp. (100%)
- OTU13**: *Streptococcus* sp. (100%)
- OTU14**: *Streptococcus* sp. (100%)
- OTU15**: *Streptococcus* sp. (100%)
- OTU16**: *Streptococcus* sp. (100%)
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- OTU69**: *Streptococcus* sp. (100%)
- OTU70**: *Streptococcus* sp. (100%)
- OTU71**: *Streptococcus* sp. (100%)
- OTU72**: *Streptococcus* sp. (100%)
- OTU73**: *Streptococcus* sp. (100%)
- OTU74**: *Streptococcus* sp. (100%)
- OTU75**: *Streptococcus* sp. (100%)
- OTU76**: *Streptococcus* sp. (100%)
- OTU77**: *Streptococcus* sp. (100%)
- OTU78**: *Streptococcus* sp. (100%)
- OTU79**: *Streptococcus* sp. (100%)
- OTU80**: *Streptococcus* sp. (100%)
- OTU81**: *Streptococcus* sp. (100%)
- OTU82**: *Streptococcus* sp. (100%)
- OTU83**: *Streptococcus* sp. (100%)
- OTU84**: *Streptococcus* sp. (100%)
- OTU85**: *Streptococcus* sp. (100%)
- OTU86**: *Streptococcus* sp. (100%)
- OTU87**: *Streptococcus* sp. (100%)
- OTU88**: *Streptococcus* sp. (100%)
- OTU89**: *Streptococcus* sp. (100%)
- OTU90**: *Streptococcus* sp. (100%)
- OTU91**: *Streptococcus* sp. (100%)
- OTU92**: *Streptococcus* sp. (100%)
- OTU93**: *Streptococcus* sp. (100%)
- OTU94**: *Streptococcus* sp. (100%)
- OTU95**: *Streptococcus* sp. (100%)
- OTU96**: *Streptococcus* sp. (100%)
- OTU97**: *Streptococcus* sp. (100%)
- OTU98**: *Streptococcus* sp. (100%)
- OTU99**: *Streptococcus* sp. (100%)
- OTU100**: *Streptococcus* sp. (100%)

“Next Generation Probiotics” in Animal Science

Multimiomics analysis

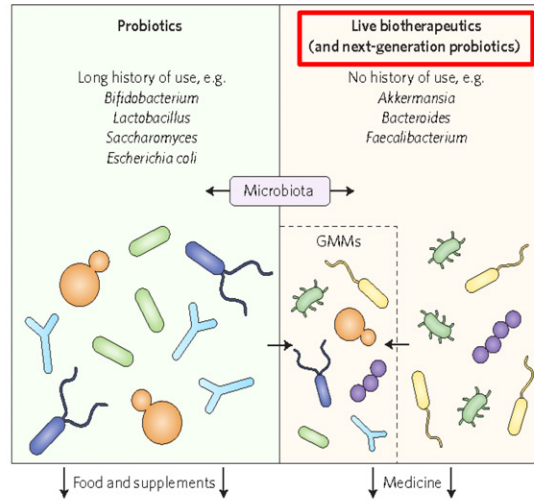


DFM application with spore-forming probiotics



Spore-forming bacteria like *Bacillus coagulans* and *Clostridium butyricum* have various health benefits on the livestock as well as companion animal

O'Toole et al. NM 2017



Combination of multiomics analysis and DFM application with spore-forming probiotics will be open new era for **next generation probiotics** with **concept of LBP** in the animal industry

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SNU Lab. of Animal Microbiology (SLAM)



Thank you
For your attention

