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Regional Workshop on
Waste Agricultural Biomass

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Technology and Innovation for Production of Cellulosic Biofuels

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Research Institute of Innovative Technology for the Earth (RITE)



Today's outline

- RITE: Who we are
- Biofuel: current status
- RITE Bioprocess
 - Features
 - Biofuels
 - Biochemicals

Research Institute of Innovative Technology for the Earth (RITE)

Established: 1990

**Status: Non-profit organization under
the Ministry of Economy, Trade and Industry**

Annual budget: JPY 5.1 billion (\$ 51 million)

Mission:

- **Development of environmental technologies
against global warming problems**

Main research fields:

- **Biorefinery**
- **CO₂ geological sequestration**



Biorefinery group

Production of biofuels /
chemicals from biomass



Research fields

Biorefinery: Production of energy and chemicals from Biomass

Research projects

Chemicals

- Succinic acid
- L-Lactate, D-Lactate
- Propanol (Raw material for propylene)
- Ethanol (Raw material for ethylene)
- Amino acid

Biofuels

- Ethanol
- Propanol
- Butanol

Energy

- H₂

Biofuels: Current situation

■ Sustainability?

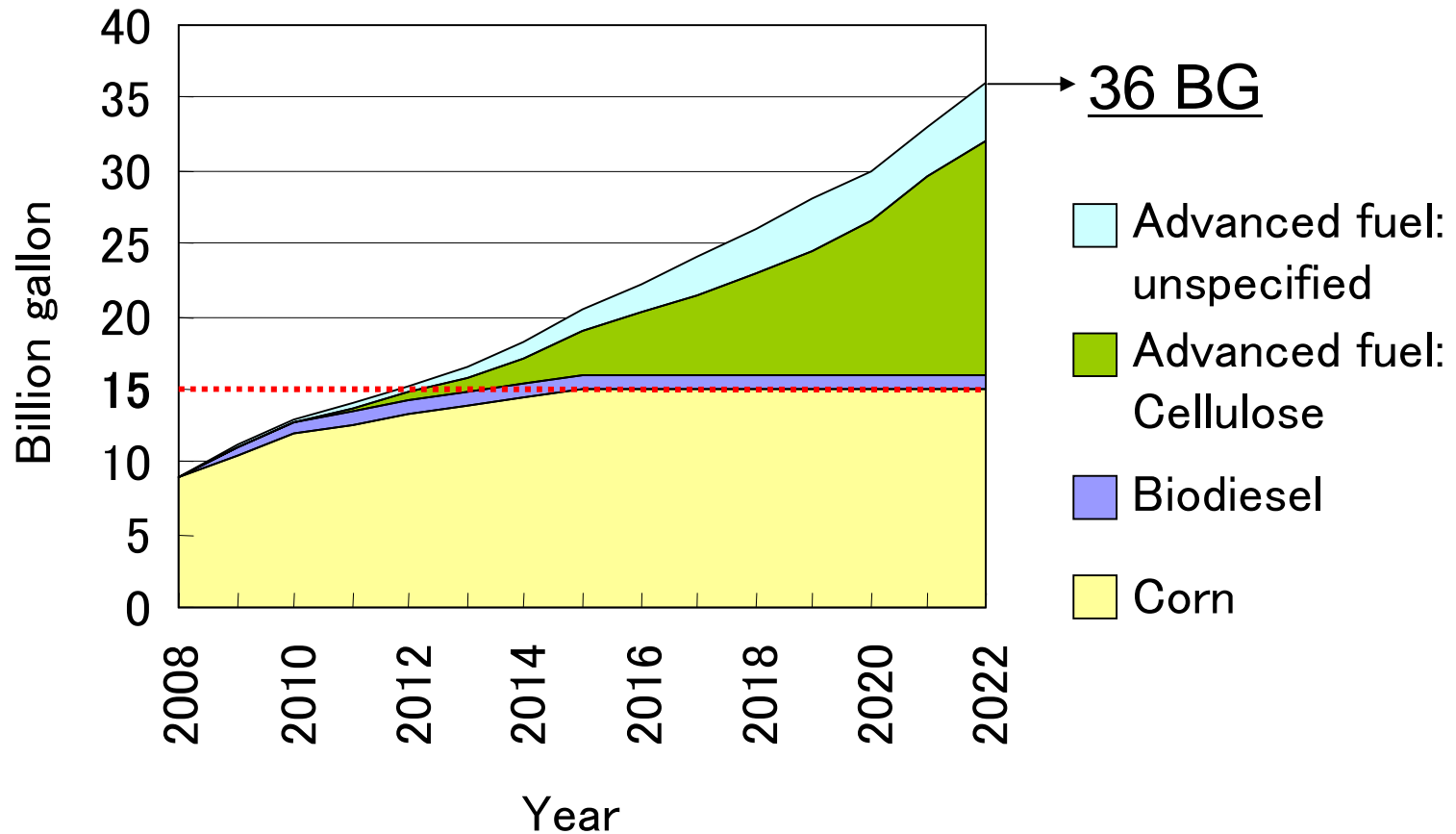
Food-Fuel issue, environmental effect...

■ Solution: Cellulosic biofuels

- Non-food resources
- Countermeasures against
global warming problems

Challenge: Development of a cost-effective
production process

Cellulosic biofuel in USA



Source: EPA Renewable Fuel Standard2

A novel and highly efficient bioprocess

**Growth-Arrested Bioprocess
(RITE bioprocess)**

RITE strain

Corynebacterium glutamicum

Under oxygen deprivation

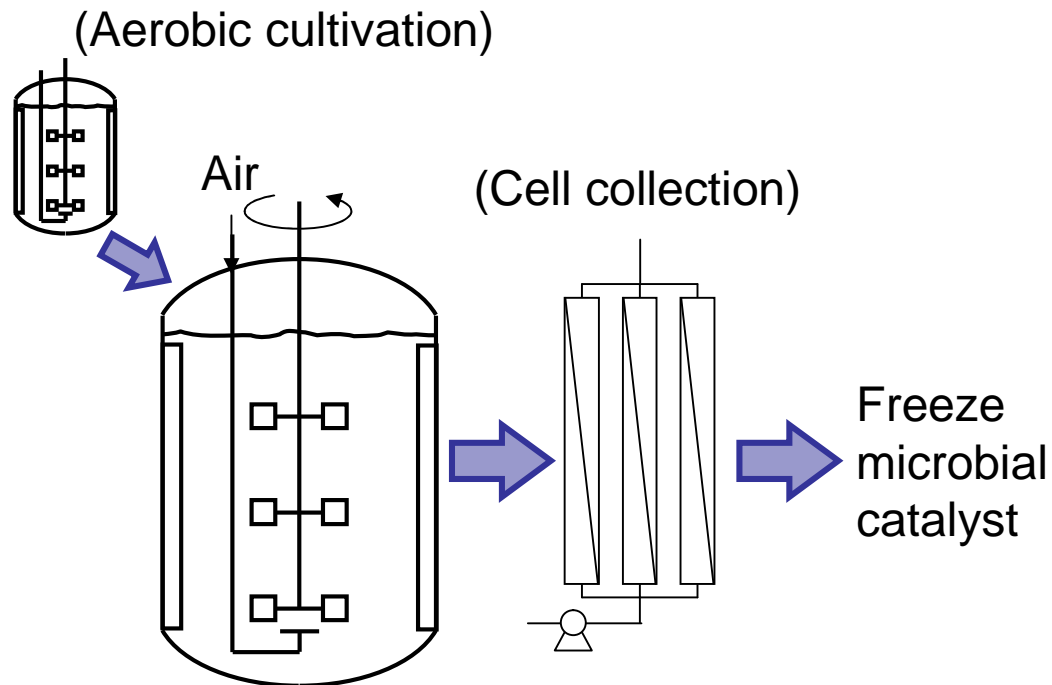
- **Growth-arrested**
- **Maintains main metabolic capabilities**

RITE bioprocess

JP-Patent 3869788

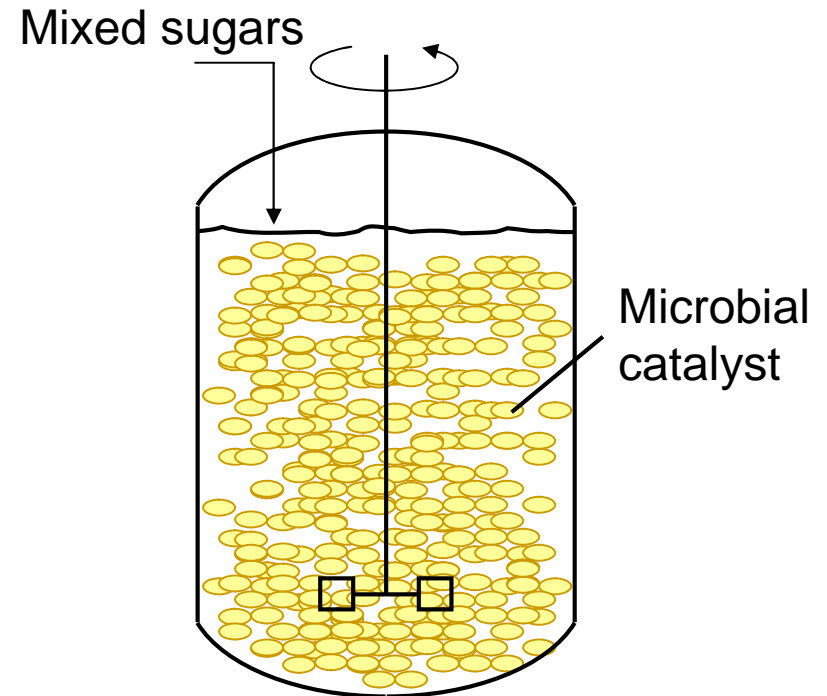
INDIA 209524

Microbial catalyst preparation



Growth by cell division

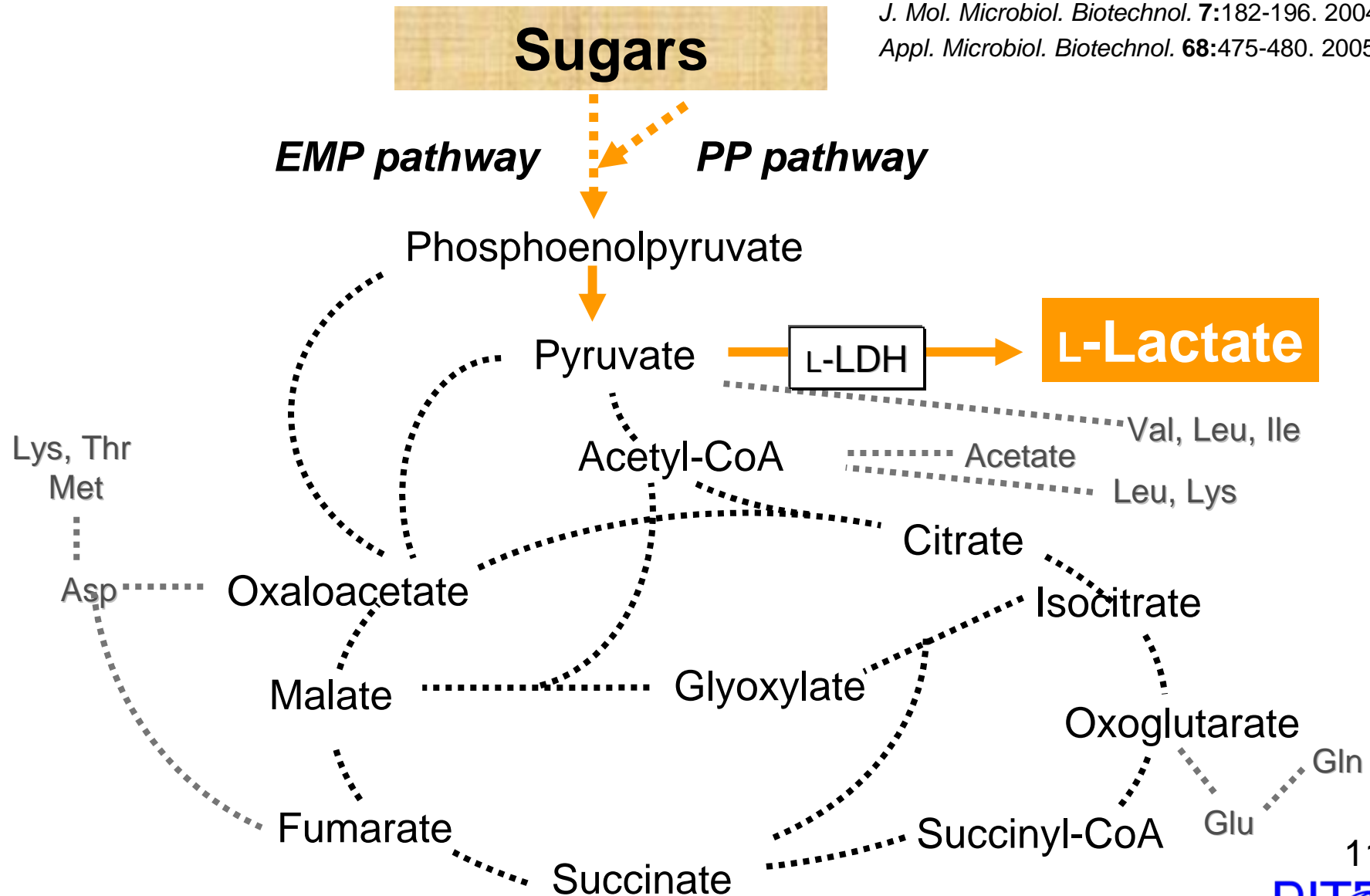
Bioconversion



Growth-arrested cells 10

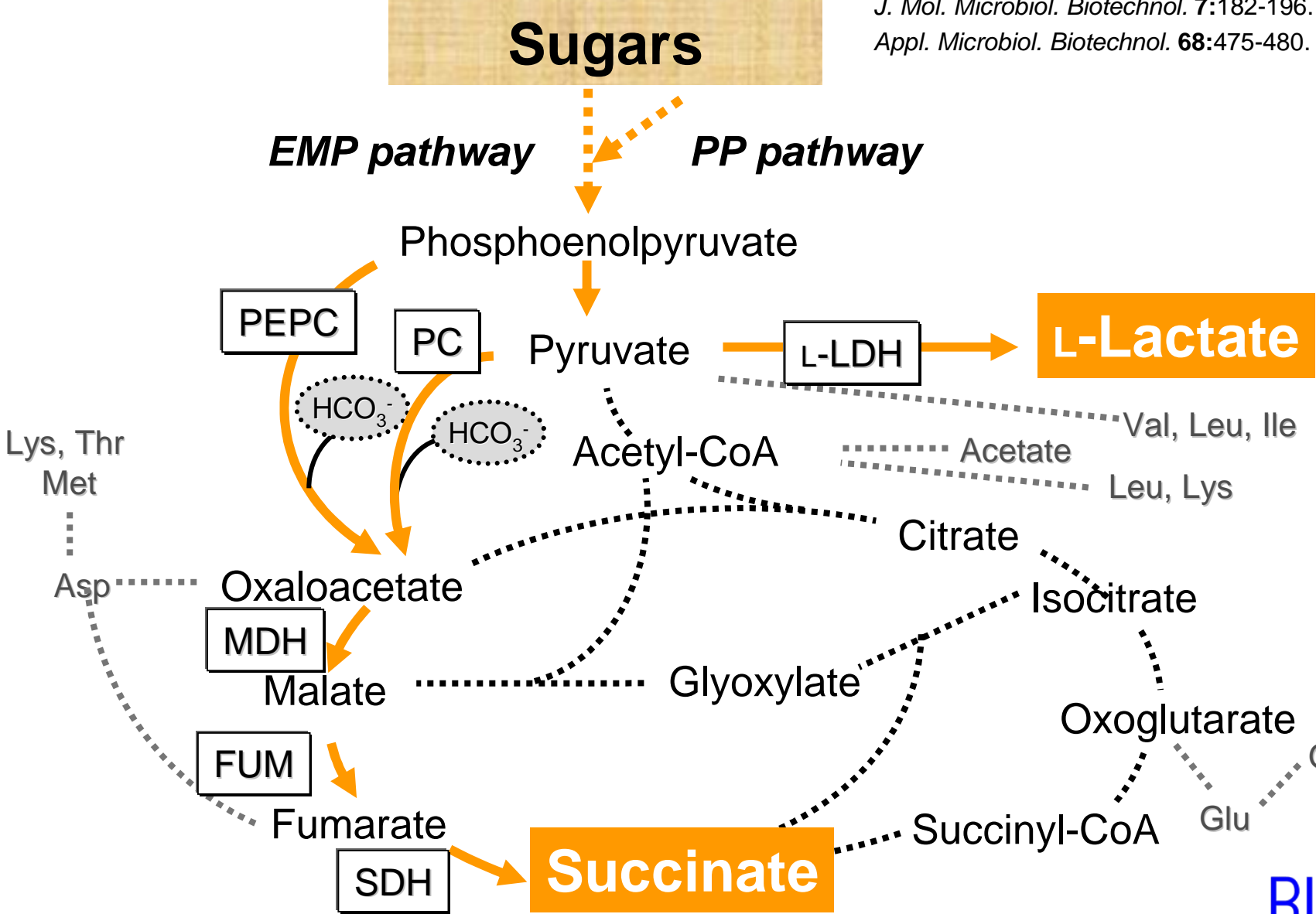
Metabolic pathways of *C. glutamicum* under oxygen deprivation (without CO₂)

J. Mol. Microbiol. Biotechnol. 7:182-196. 2004.
Appl. Microbiol. Biotechnol. 68:475-480. 2005.

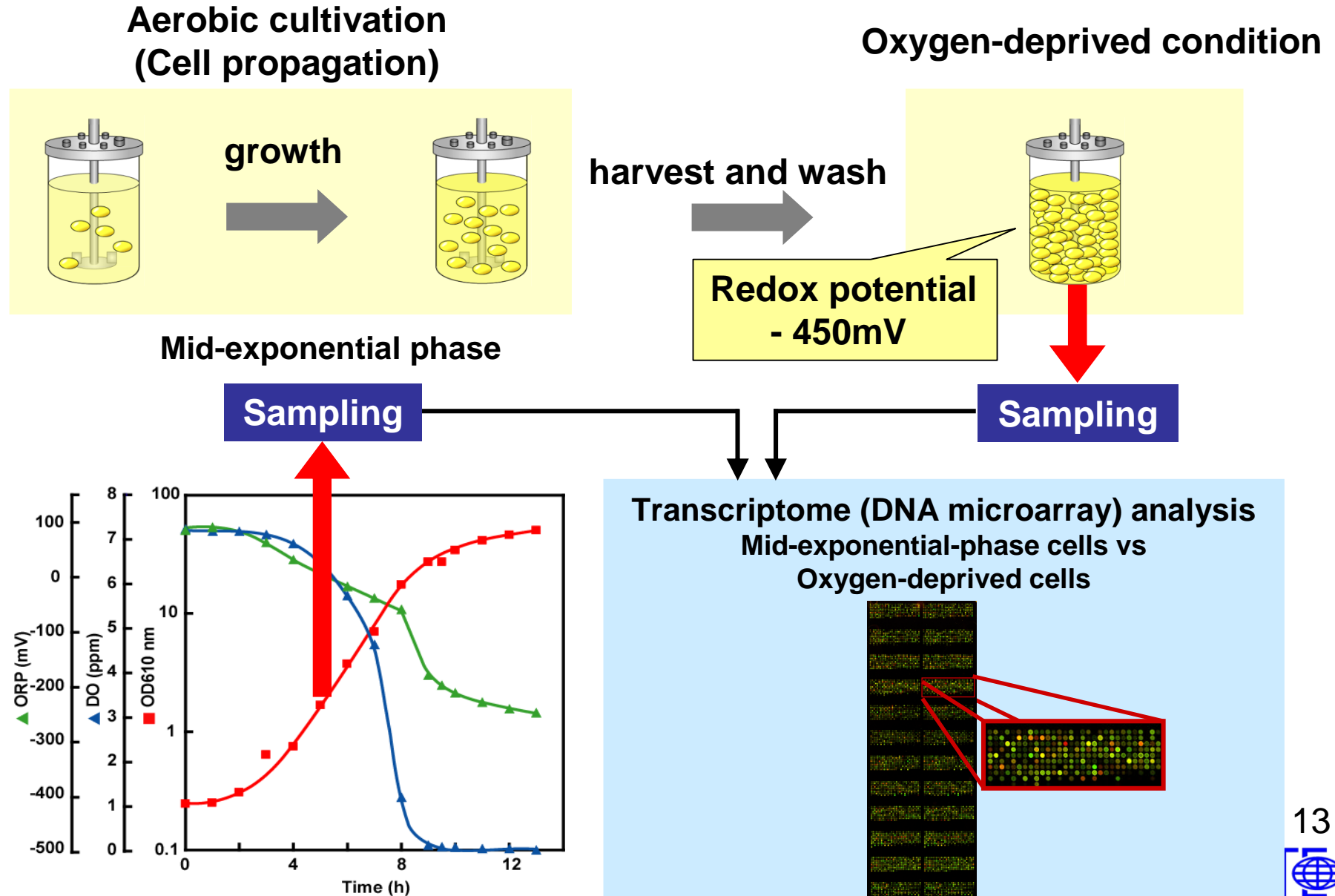


Metabolic pathways of *C. glutamicum* under oxygen deprivation (with CO₂)

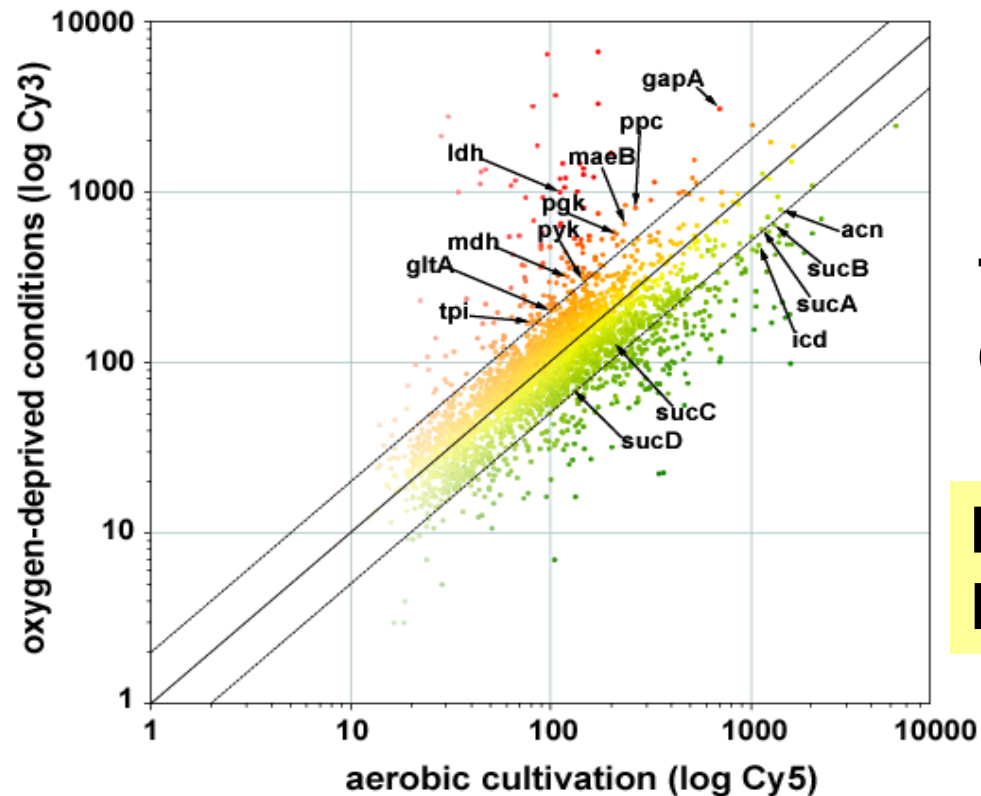
J. Mol. Microbiol. Biotechnol. 7:182-196. 2004.
Appl. Microbiol. Biotechnol. 68:475-480. 2005.



Analysis of metabolic shift under oxygen deprivation



Gene expression analysis



• Entire gene (3080 genes)

The ratios of mRNA levels
(oxygen-deprived conditions
/aerobic cultivation)

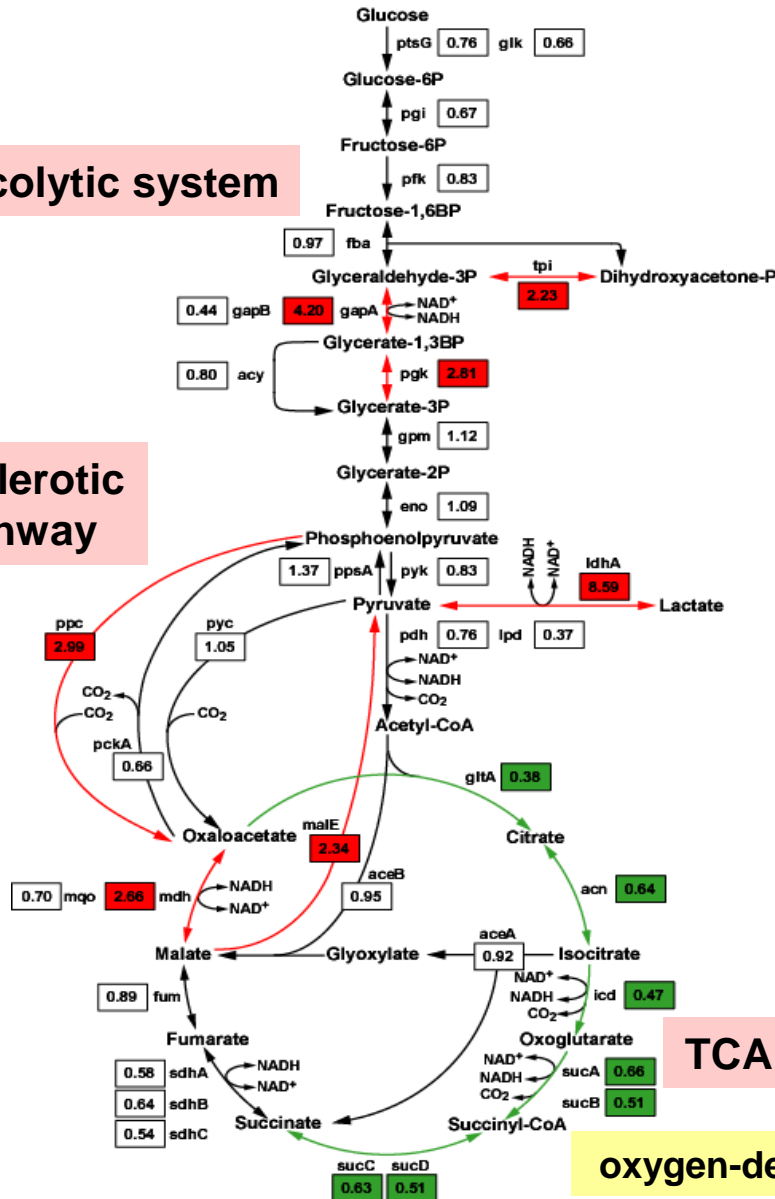
More than 2-fold 161 genes
Less than 1/2-fold 221 genes

A gene expression profile is different greatly between aerobic and oxygen-deprived conditions

Expression analysis of glucose metabolism

glycolytic system

anaplerotic pathway



Enzyme Relative enzyme activities
oxygen-deprived conditions / aerobic

GAPDH	5.3
PGK	10.5
TPI	19.1
PEPC	4.5
LDH	14
MDH	25.8

Genes encoding several key enzymes involved in the glycolytic and organic acid production pathways were significantly up-regulated under growth-arrested bioprocess.

Microbiology. 153:2491-2504. 2007.

TCA cycle

oxygen-deprived conditions / aerobic cultivation

Application of “RITE bioprocess” For the production of biofuels

- Ethanol
- Propanol
- Butanol

Biofuels: Current situation

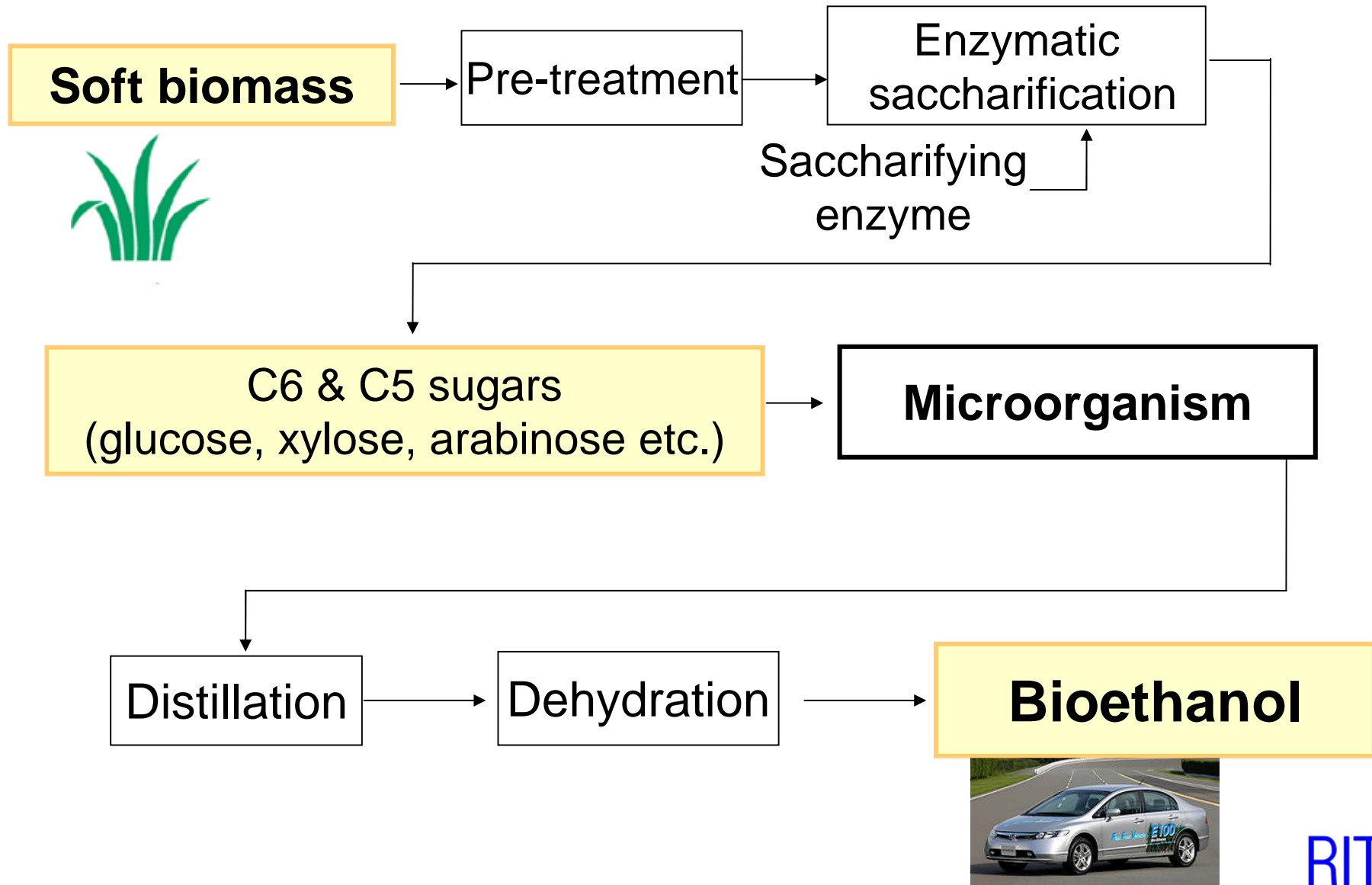
■ Sustainability?

Biofuels “pros & cons”

■ Cellulosic biofuels:

- Non-food resources
- Countermeasures against
global warming problems

Ethanol production from soft biomass

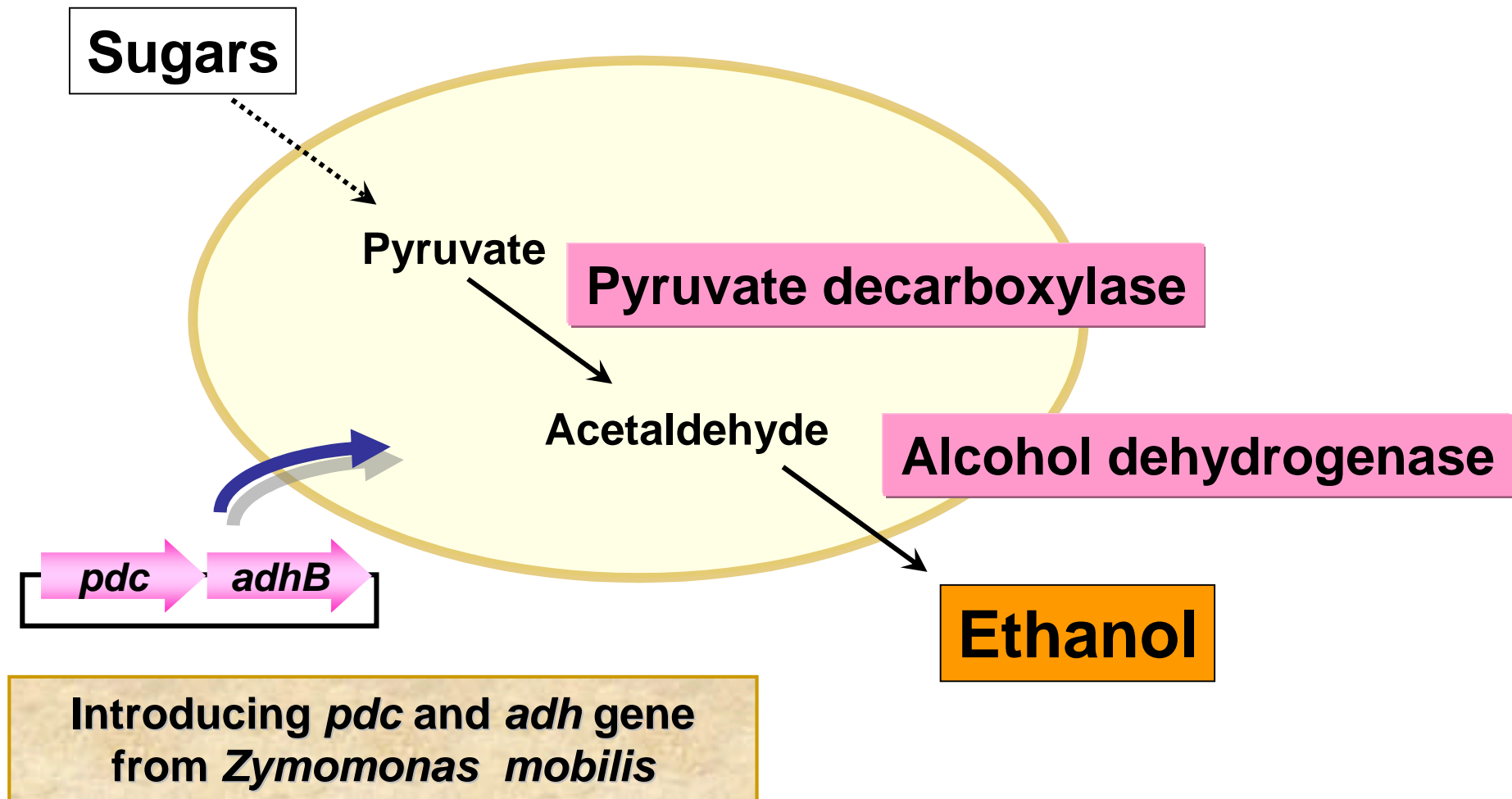


Ethanol production

Important traits for industrialization

- **High productivity**
- Simultaneous utilization of C₆ & C₅ sugars
- Tolerance to “fermentation inhibitors”

Development of the ethanol producing strain



Ethanol production

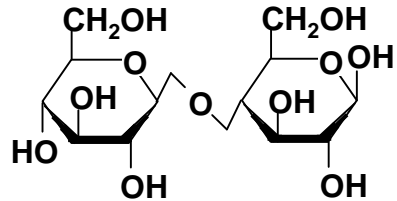
Important traits for industrialization

- High productivity
- **Simultaneous utilization of C₆ & C₅ sugars**
- Tolerance to “fermentation inhibitors”

Introducing ability to utilize sugars derived from biomass

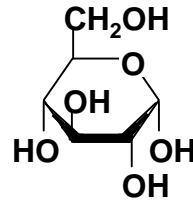
Cellulose

Cellulose (C₆-C₆)



Adaptive mutant for cellobiose uptake ability¹⁾

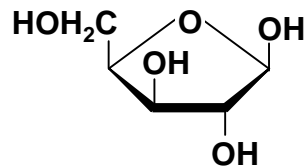
Glucose (C₆)



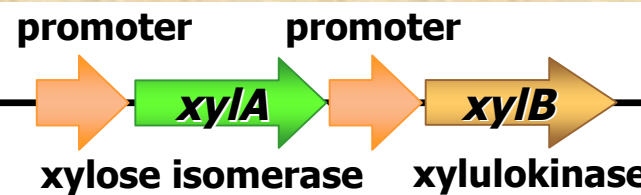
RITE strain

Hemicellulose

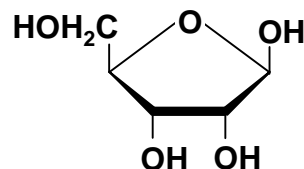
Xylose (C₅)



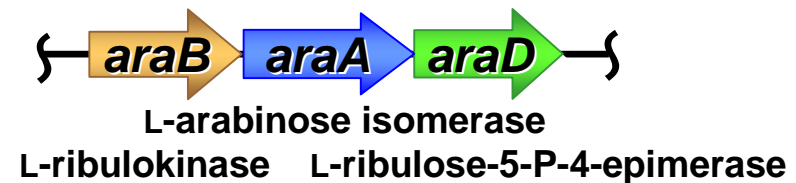
Chromosomal integration for xylose metabolic ability²⁾



Arabinose (C₅)

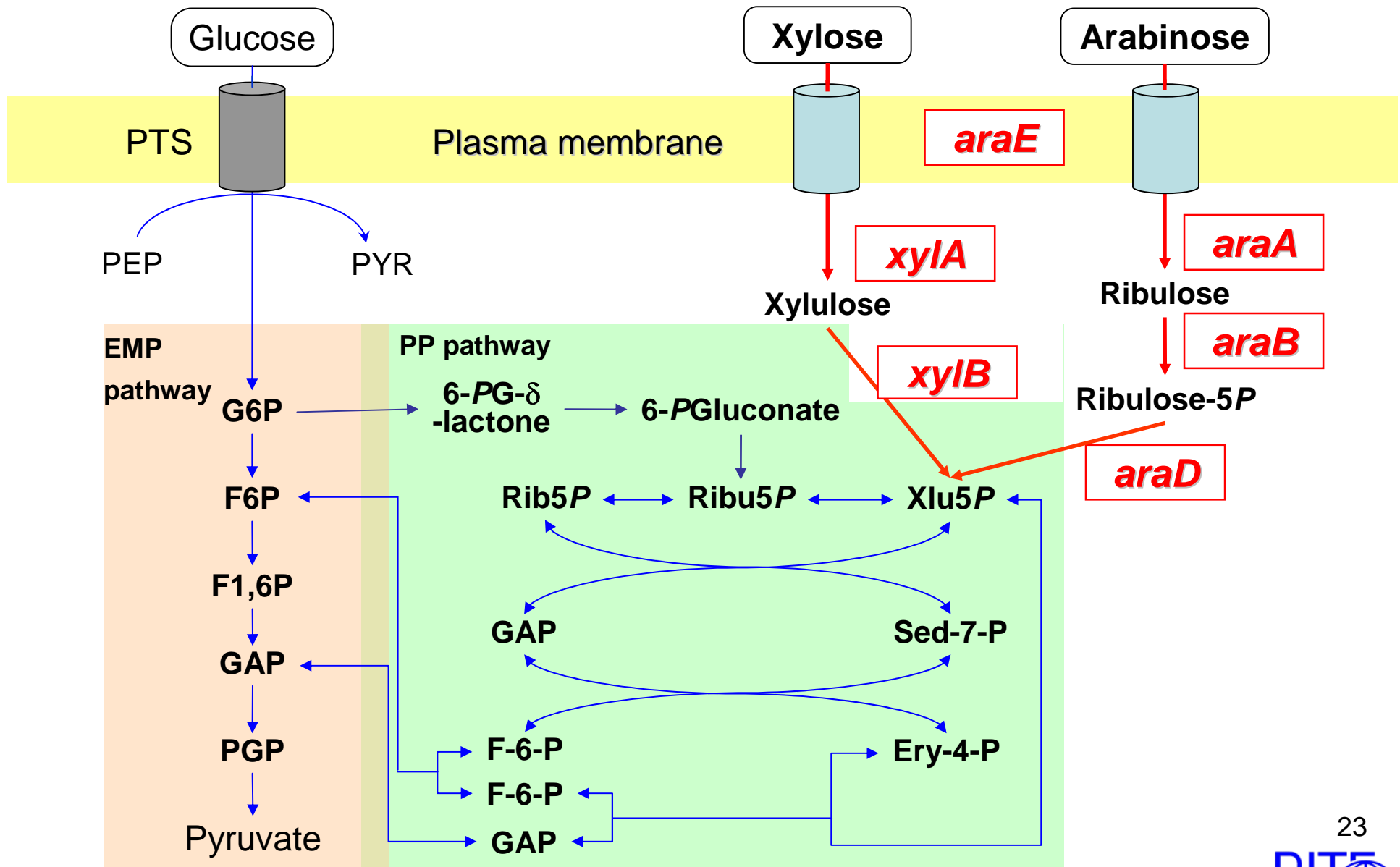


Chromosomal integration for arabinose metabolic ability³⁾



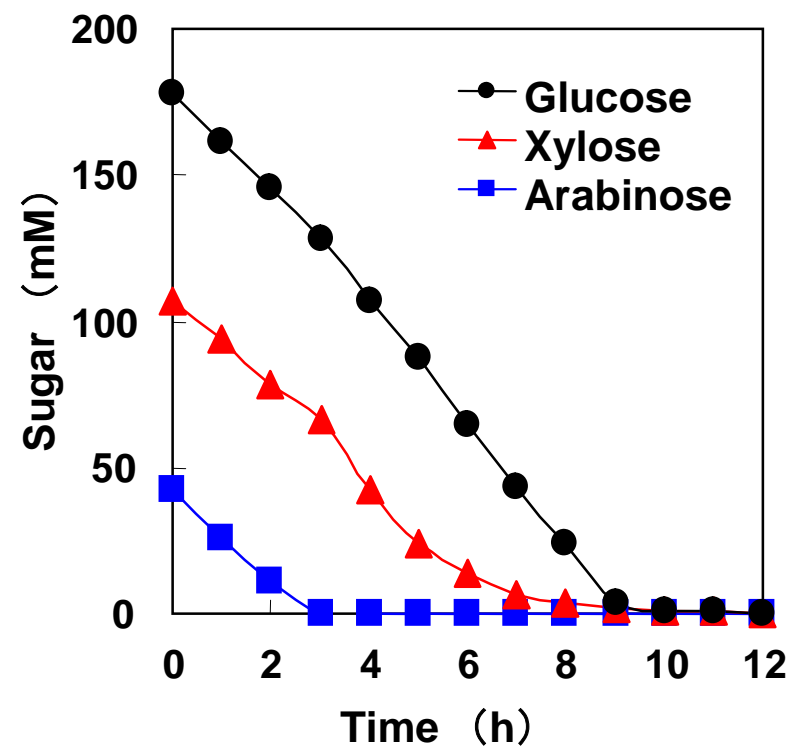
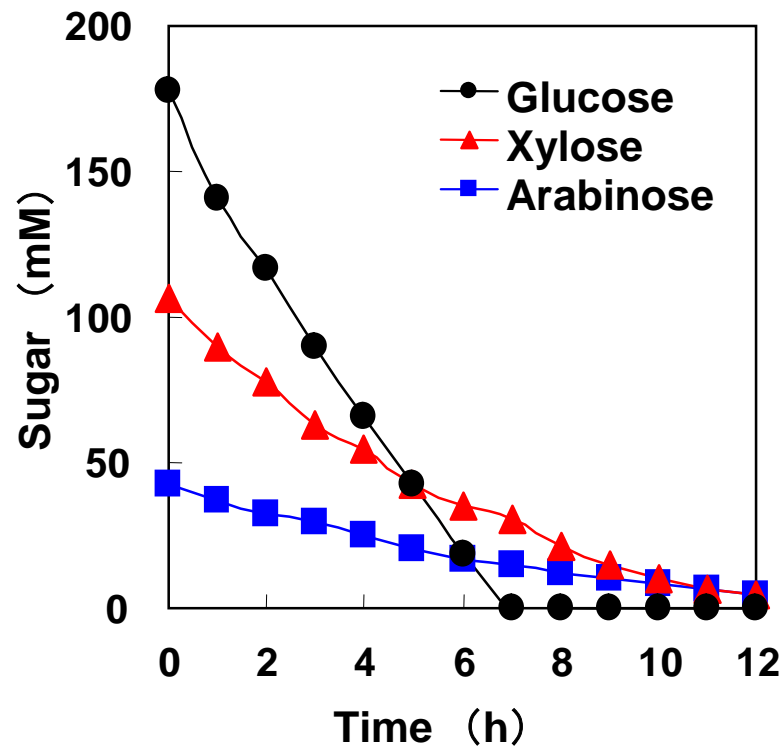
1) *Microbiology* **149**:1569-80. 2003. 2) *Appl. Environ. Microbiol.* **72**:3418-28. 2006. 3) *Appl. Microbiol. Biotechnol.* **77**:1053-62. 2008.

Engineering of xylose and arabinose metabolic pathways



Simultaneous Utilization of Mixed Sugar

Introduction of a pentose transporter

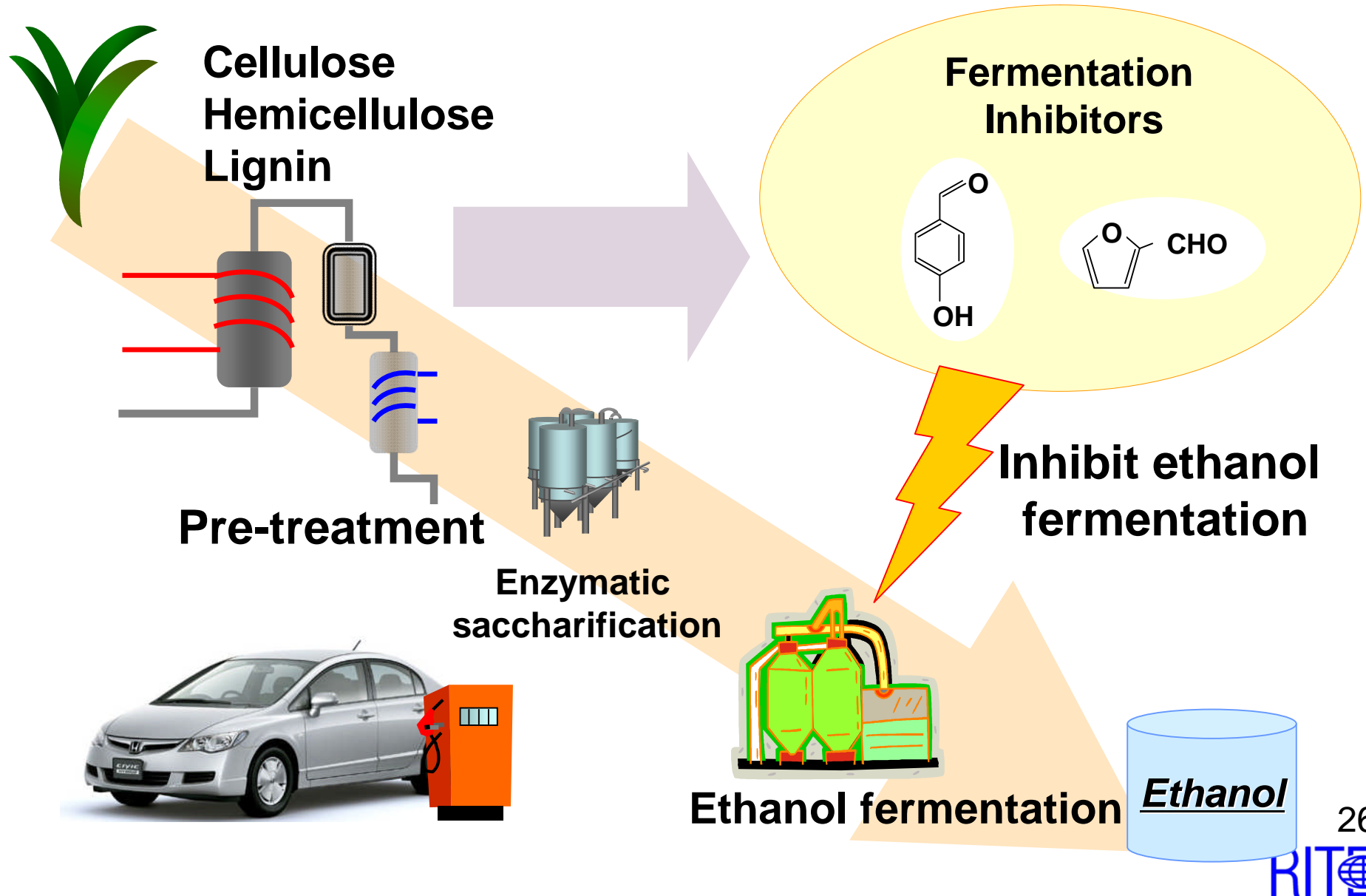


Ethanol production

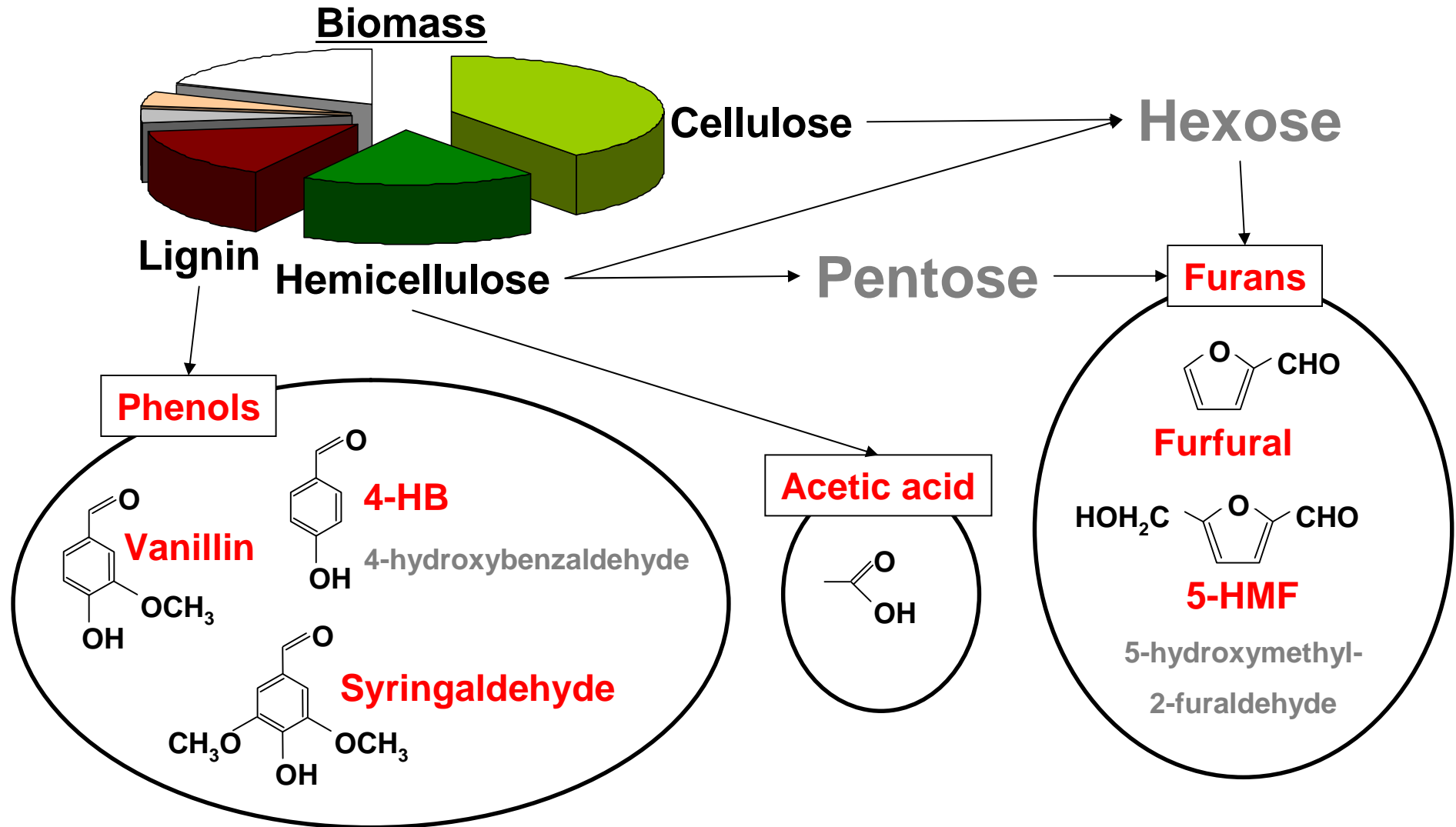
Important traits for industrialization

- High productivity
- Simultaneous utilization of C₆ & C₅ sugars
- **Tolerance to “fermentation inhibitors”**

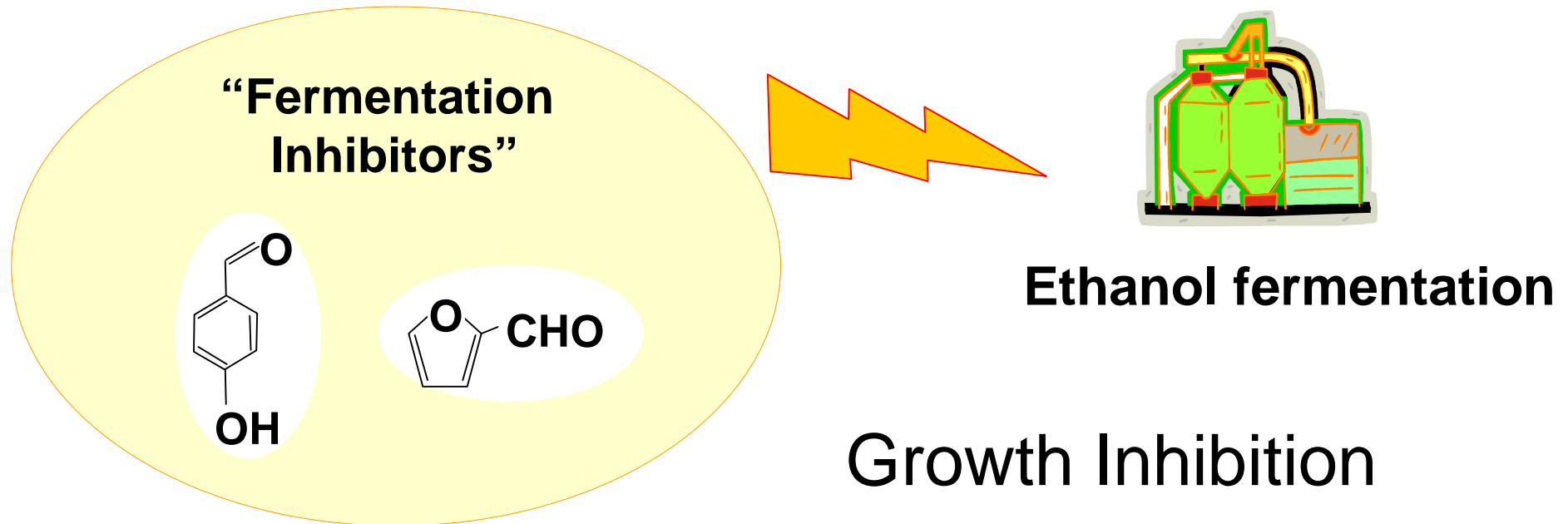
What is “fermentation inhibitors”?



Major “fermentation inhibitors”



Inhibition mechanism



No inhibition to the ethanol producing metabolic pathway!

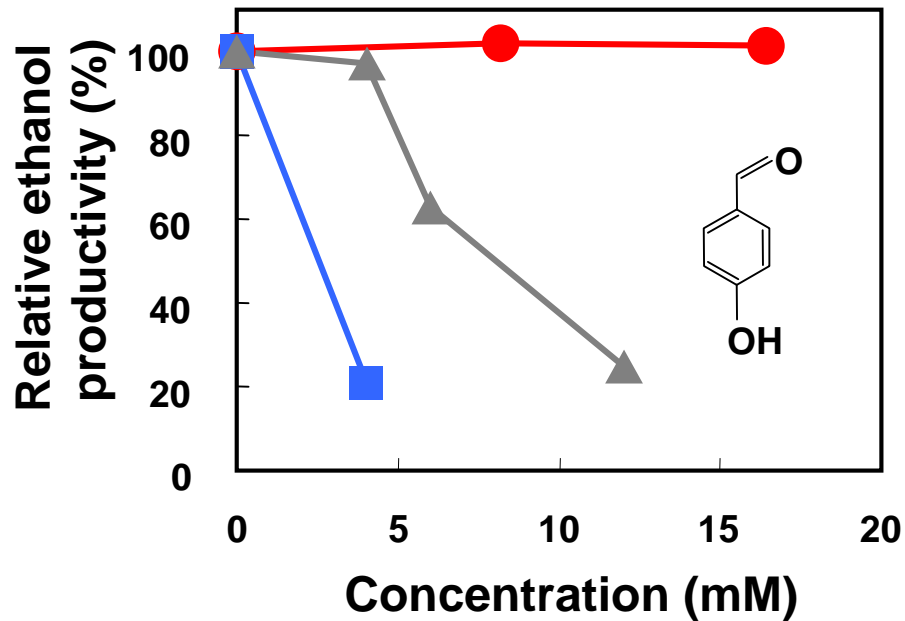
Effect of lignocellulose-derived inhibitors on growth and ethanol production by growth-arrested *Corynebacterium glutamicum* R. *Appl. Environ. Microbiol.* **74**:754-760. 2007.

28

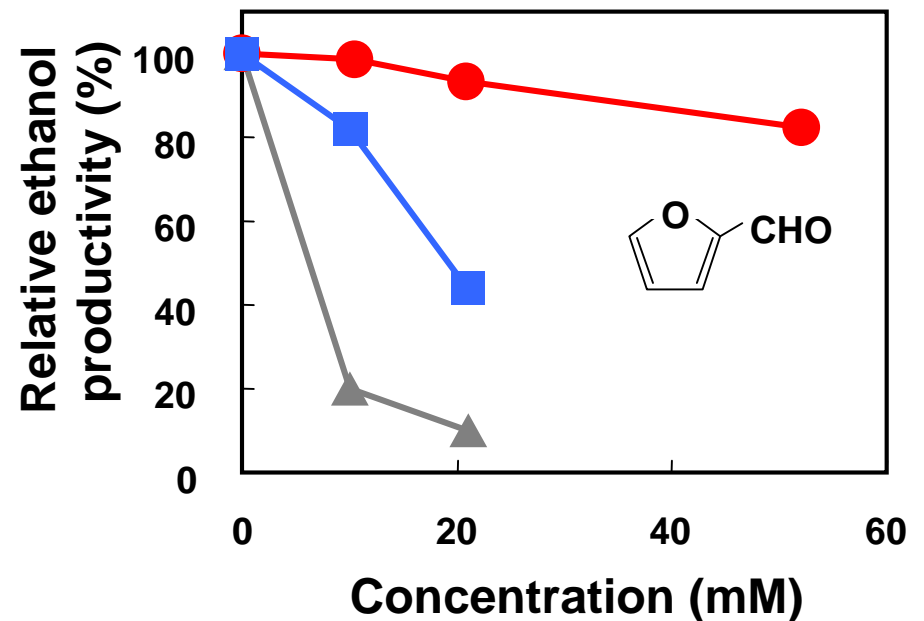
Tolerance to the inhibitors

● RITE Bio-Process ■ *Z. mobilis* ▲ *S. cerevisiae*

4-HB



Furfural



Biobutanol as “fuel”

- Diesel additive
- High energy density
- Low water solubility

Expected use as;

- Fuel for diesel engines
- Aviation fuel
- Ethanol – Butanol: Mixed-use complementally
(Synergistic effects)

Trends in Biobutanol R&D

- Will improve ABE fermentation
- Create novel producing microorganisms

Prediction of practical application

Fundamental research (3-5 years)

+

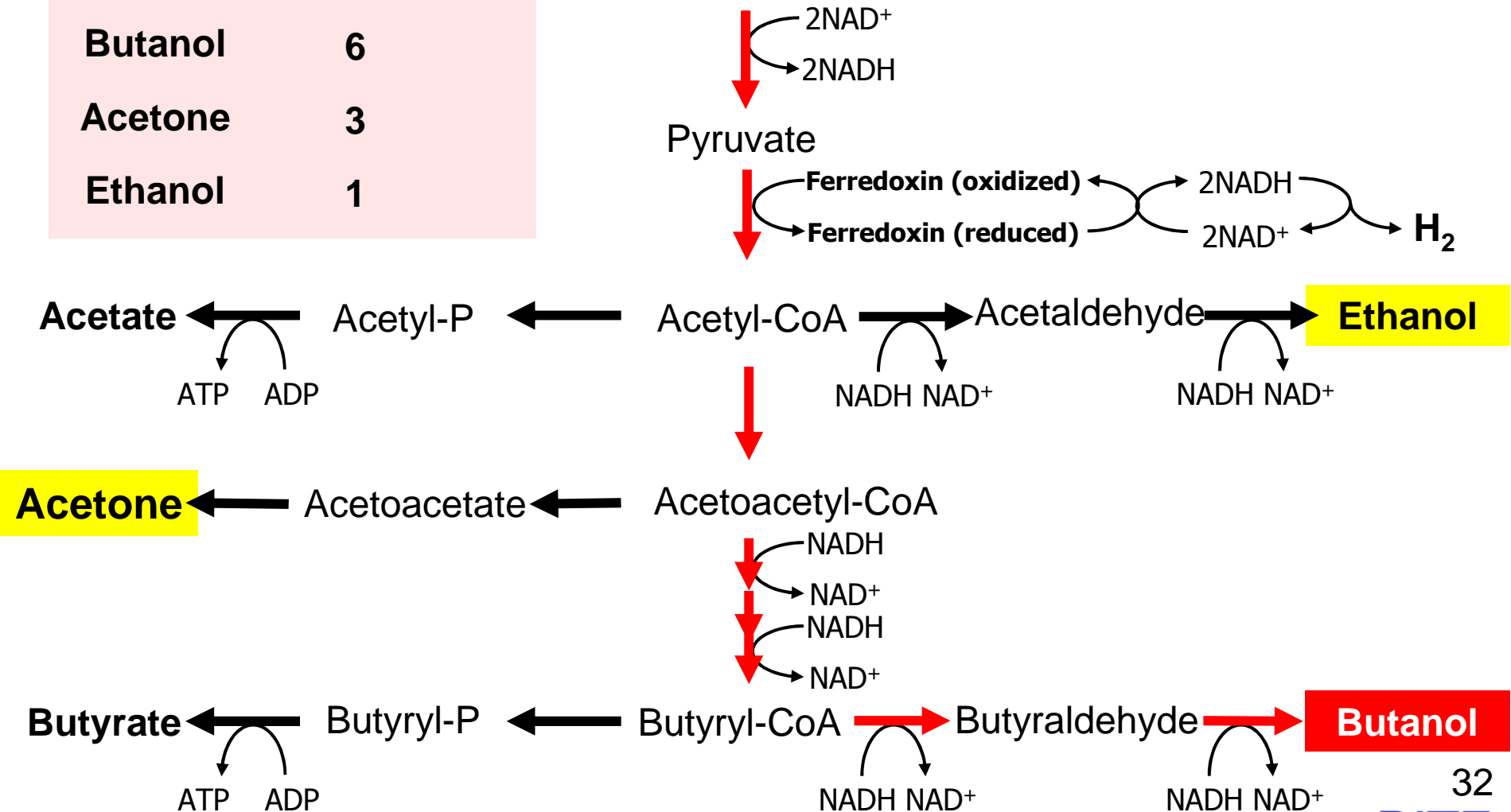
Industrialization research (2 years)

Clostridial ABE fermentation pathway

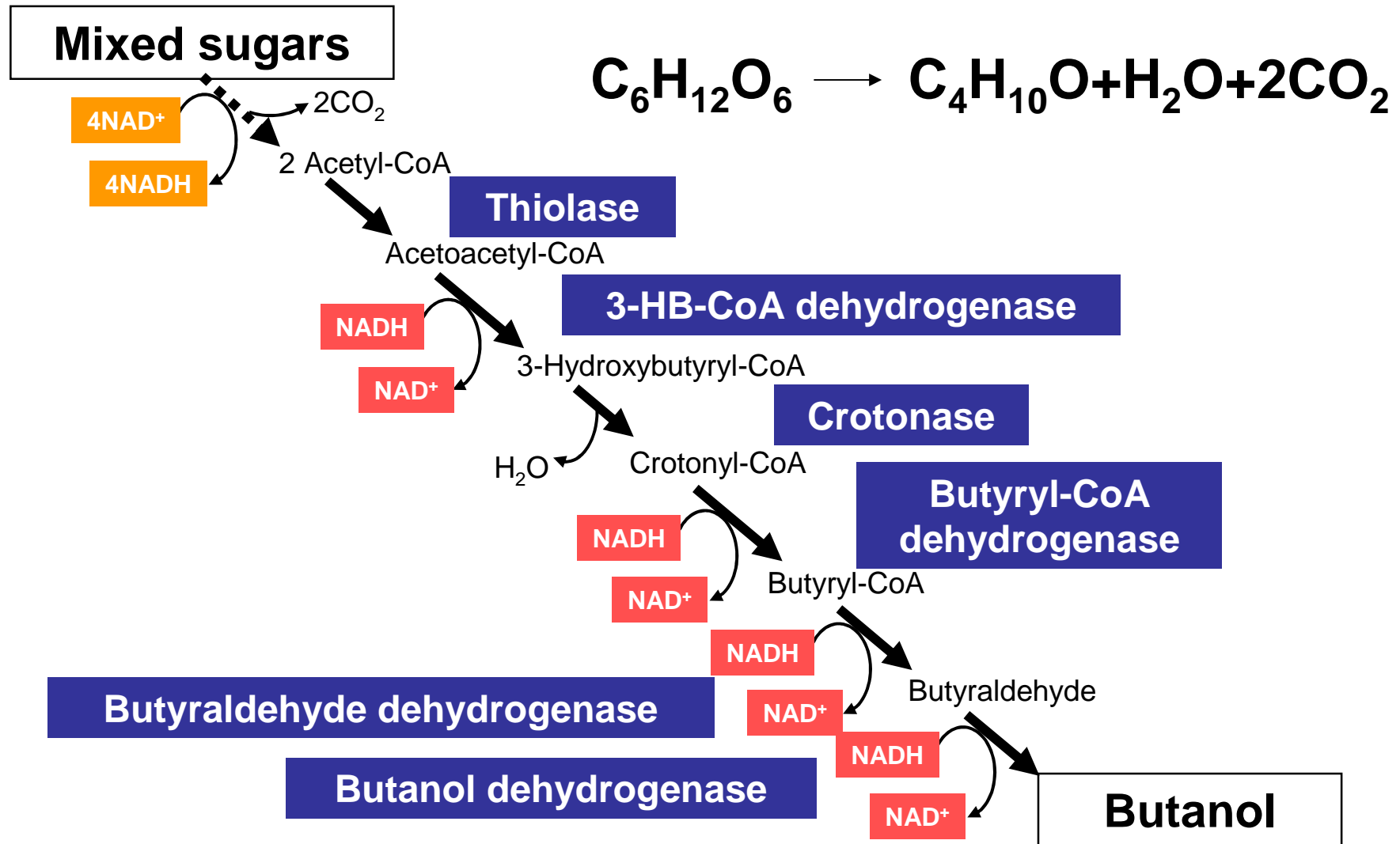
Production ratio (mol)

Butanol	6
Acetone	3
Ethanol	1

Mixed sugars



Butanol production by recombinant *E.coli*.



Application of “RITE bioprocess” for the production of biochemicals

Examples;

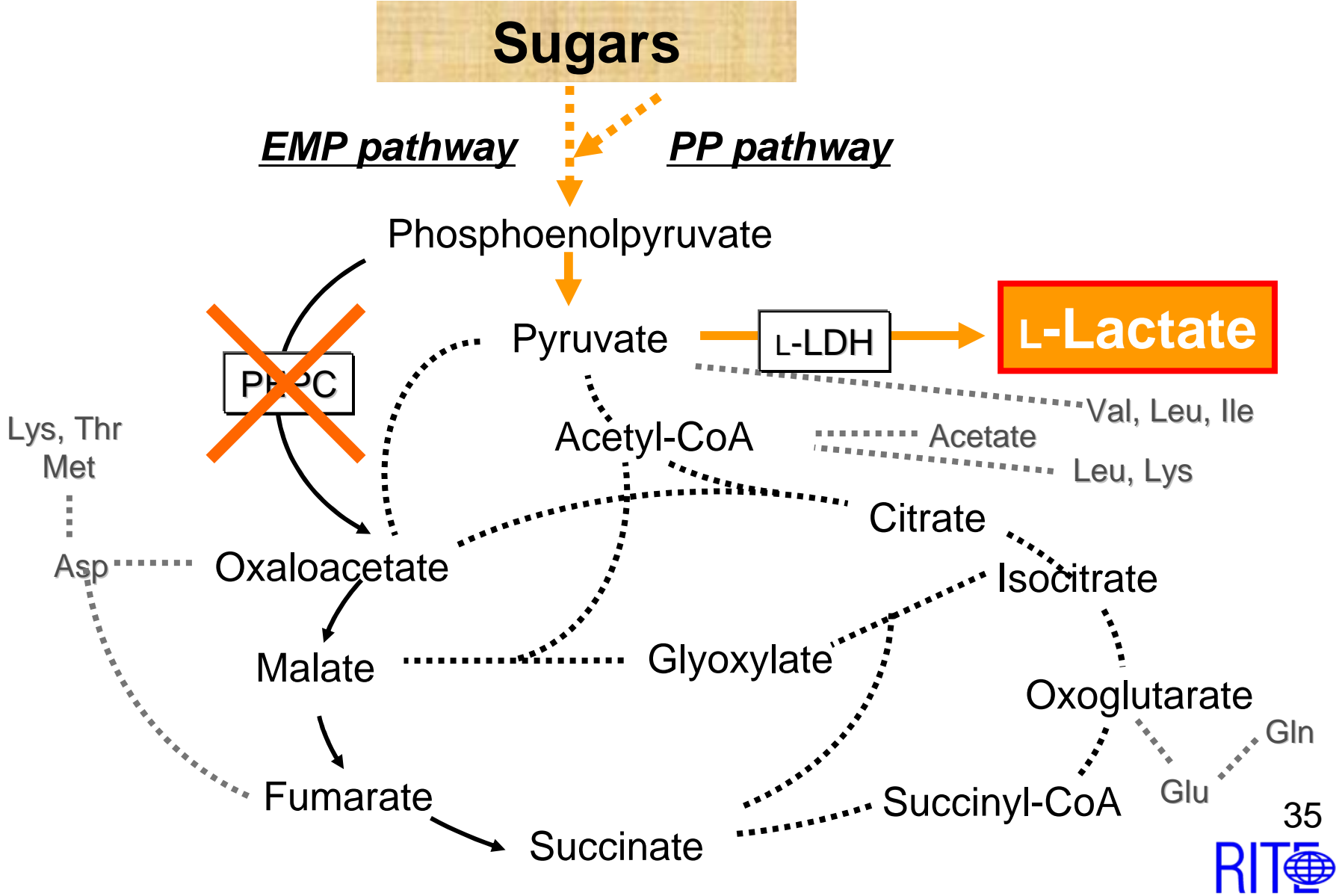
L-Lactate

D-Lactate

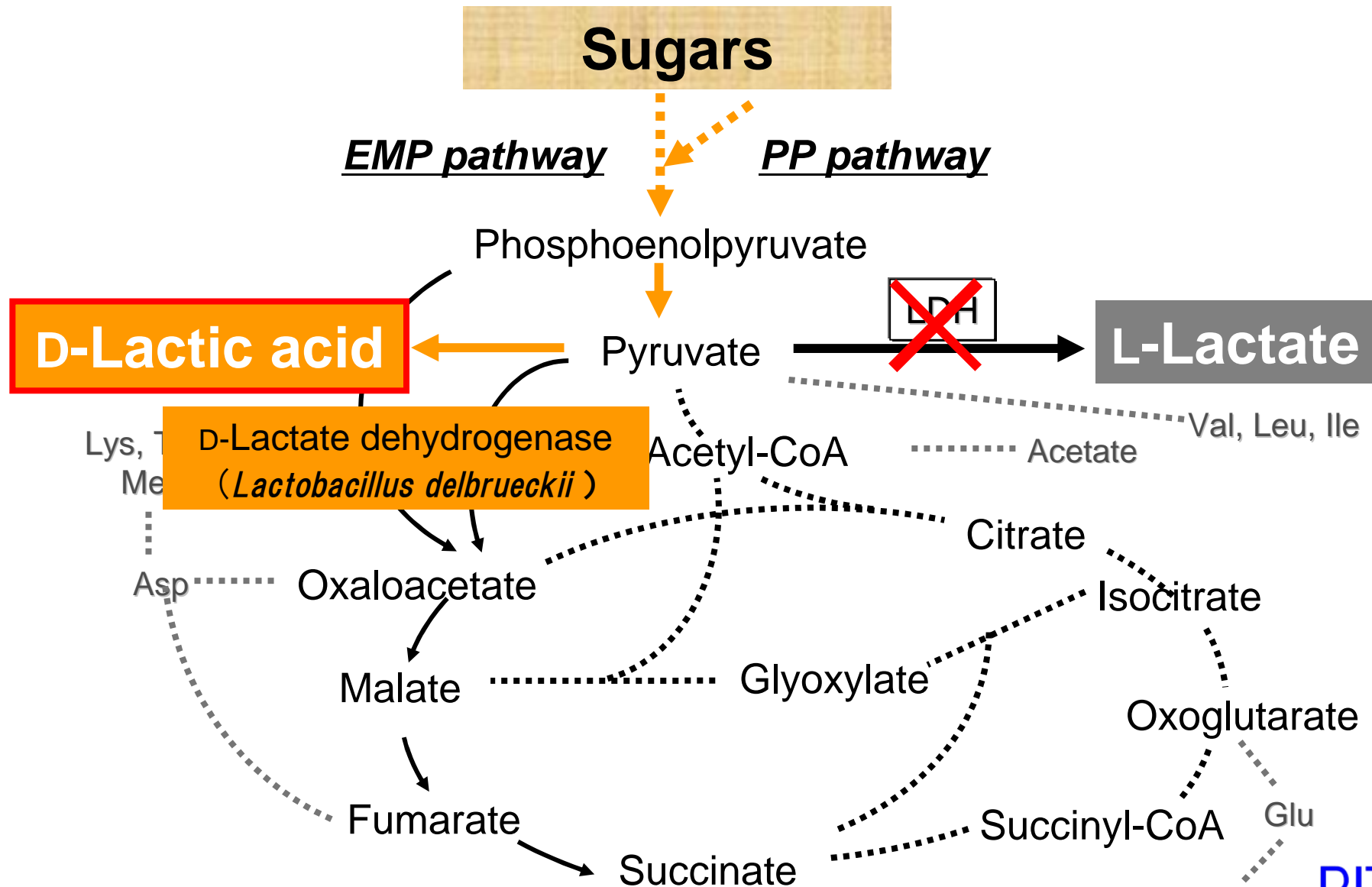
Succinate

Amino acid (L-Alanine)

Metabolic engineering for L-Lactate production



Metabolic engineering for D-Lactate production



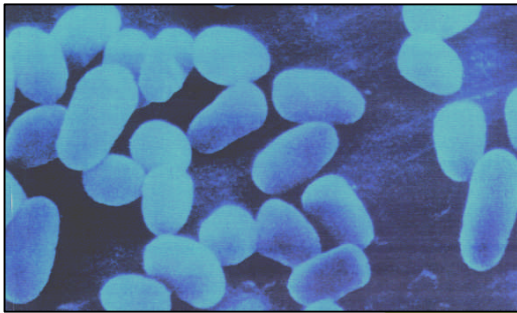
Comparison of D-lactate production

Microorganism	Titer (g/l)	Productivity (g/l/h)	Reference
<i>Lactobacillus delbrueckii</i>	63	1.3	<i>Macromolecul Biosci.</i> 4: 1021-1027. (2004)
<i>E.coli</i> RR1	62	2.1	<i>Appl Environ Microbiol.</i> 65: 1384-1389. (1999)
<i>E. coli</i> W3110 SZ63	49	0.5	<i>Appl Environ Microbiol.</i> 69: 399-407. (2003)
<i>E. coli</i> MT-10934/pGlyldhA	65	1.3	JP 2005-102625 (2005)
<i>Saccharomyces cerevisiae</i> OC2	62	1.4	<i>J Biosci Bioeng.</i> 101: 172-177. (2006)
<i>E. coli</i> SZ194	92	2.1	<i>Biotechnol lett.</i> 28: 663-670. (2006)
RITE bioprocess	110	40.0	<i>Appl Microbiol Biotechnol.</i> 78: 449-454. (2008)

Comparison of succinate production

Microorganism	Titer (g/l)	Productivity (g/l/h)	Reference
<i>A. Succiniciproducens</i>	50	2.1	United States Patent 5143834 (1992)
<i>A. Succiniciproducens</i>	84	10.4	<i>Biotechnol Bioeng.</i> 99:129-135. (2008)
<i>A. succinogens</i> FZ53	106	1.4	United States Patent 5573931 (1996)
<i>E. coli</i> NZN111	28	0.7	<i>Appl Environ Microbiol.</i> 73:7837-7843. (2007)
<i>E. coli</i> AFP111/pTrc99A-pyc	99	1.3	<i>J Ind Microbiol Biotechnol.</i> 28:325-332. (2002)
RITE bioprocess (case 1)	146	3.2	<i>Appl Microbiol Biotechnol.</i> 81:459-464. (2008)
RITE bioprocess (case 2)	83	11.8	<i>Appl Microbiol Biotechnol.</i> 81:459-464. (2008)

Amino acids production by RITE bioprocess



Metabolic engineering
of *C. glutamicum* R

RITE bioprocess

- High productivity
- No energy loss for growth
- Simple system

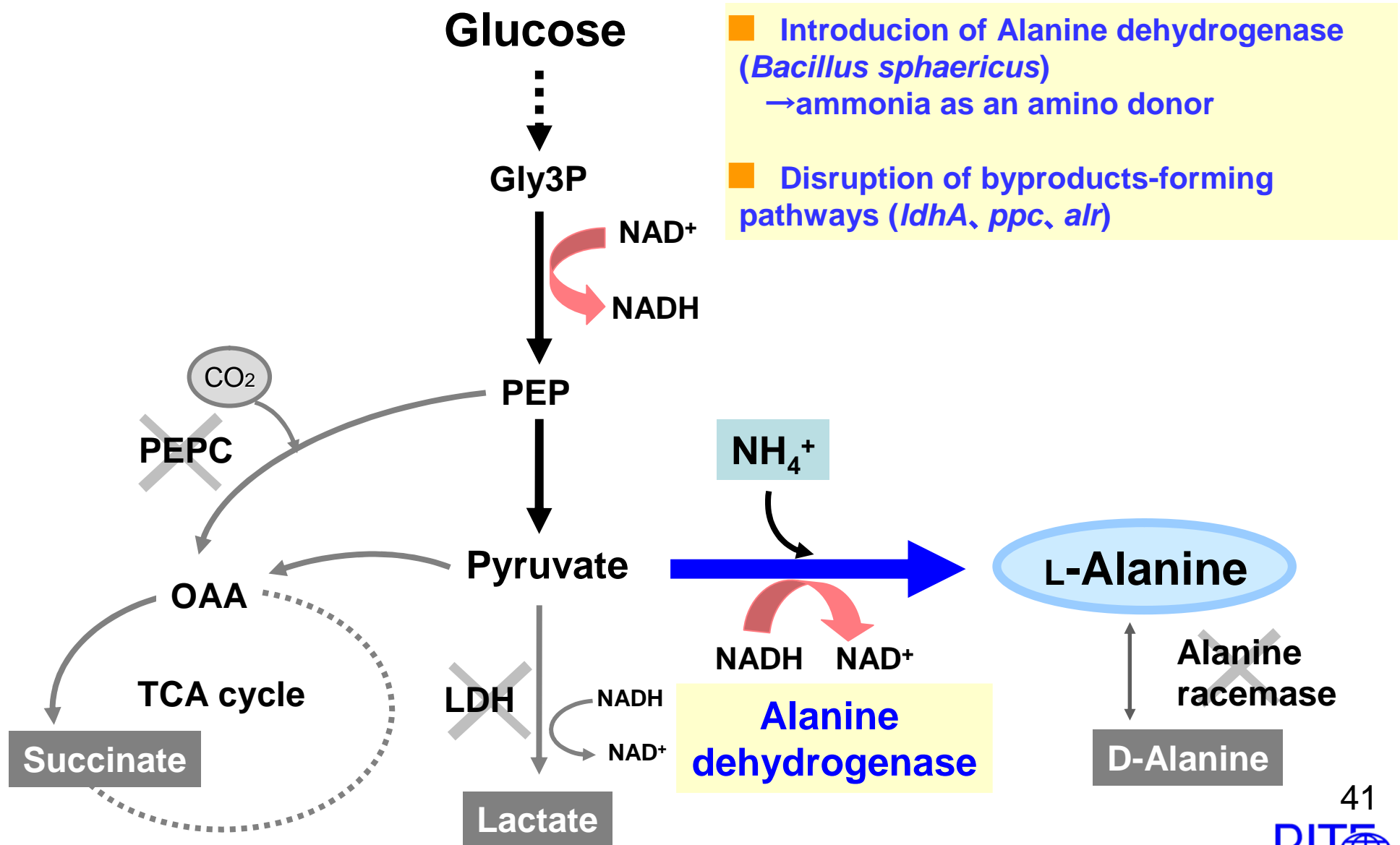
No growth

No aeration

Amino acids

L-Alanine: First trial of amino acid
production by RITE bioprocess

Metabolic engineering for L-Alanine production



Summary

- We have developed a novel process “RITE bioprocess” where we can utilize *C. glutamicum* cells like a catalyst.
- RITE bioprocess shows high productivity in bioethanol and biochemicals production.
- Genetically-engineered *C. glutamicum* consumes mixed sugar simultaneously.
- RITE-bioprocess is tolerant to fermentation inhibitors.
- We continue to improve productivity of butanol.

Host vector system

- *Agric. Biol. Chem.* **54**:443-447. 1990.
- *J. Industrial. Microbiol.* **5**:159-165. 1990.
- *Appl. Environ. Microbiol.* **57**:759-764. 1991.
- *Res. Microbiol.* **144**:181-185. 1993.
- *Biosci. Biotech. Biochem.* **57**:2036-2038. 1993.
- *Appl. Microbiol. Biotechnol.* **81**:1107-1115. 2009.

Gene expression system

- *FEMS Microbiol. Lett.* **131**:121-126. 1995.
- *Appl. Microbiol. Biotechnol.* **82**:491-500. 2009.

Physiology of corynebacteria

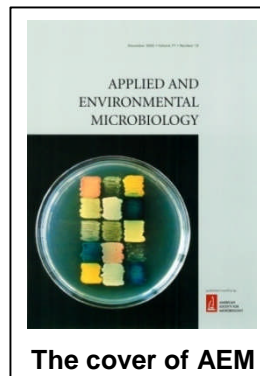
- *DNA seq.* **11**:383-394. 2000.
- *Biochem. Biophys. Res. Commun.* **289**:1307-1313. 2001.
- *J. Biosci. Bioeng.* **92**:502-517. 2001.(Review)
- *Mol. Gen. Genomics.* **271**:729-741. 2004.
- *J. Mol. Microbiol. Biotechnol.* **8**:91-103. 2004.
- *Microbiology* **153**:1042-1058. 2007.
- *Appl. Microbiol. Biotechnol.* **75**:889-897. 2007.
- *Microbiology* **153**:2190-2202. 2007.
- *Appl. Microbiol. Biotechnol.* **76**:1347-1356. 2007.
- *Microbiology* **154**:264-274. 2008.
- *Mol. Microbiol.* **67**:597-608. 2008.
- *J. Mol. Microbiol. Biotechnol.* **15**:264-276. 2008.
- *Appl. Microbiol. Biotechnol.* **78**:309-318. 2008.
- *J. Bacteriol.* **190**:3264-3273. 2008.
- *Appl. Environ. Microbiol.* **74**:5290-5296. 2008.
- *Microbiology* **154**:3073-3083. 2008.
- *Appl. Microbiol. Biotechnol.* **81**:291-301. 2008.
- *J. Bacteriol.* **191**:968-977. 2009.
- *Microbiology* **155**:741-750. 2009.
- *Appl. Microbiol. Biotechnol.* **83**:315-357. 2009.
- *J. Bacteriol.* **191**:2964-2972. 2009.
- *Appl. Environ. Microbiol.* **75**:3419-3429. 2009.
- *Appl. Environ. Microbiol.* **75**:3461-3468. 2009.
- *J. Bacteriol.* **191**:4251-4258. 2009.
- *J. Biol. Chem.* **284**:16736-16742. 2009.
- *Microbiology* **155**:3652-3660. 2009.



C. glutamicum R



The cover of MM



The cover of AEM

Gene transformation methods

- *Mol. Microbiol.* **11**:739-746. 1994.
- *Mol. Gen. Genet.* **245**:397-405. 1994.
- *Biotech. Lett.* **17**:1143-1148. 1995.

RITE bioprocess

- *Microbiology* **149**:1569-1580. 2003.
- *J. Mol. Microbiol. Biotechnol.* **7**:182-196. 2004.
- *J. Mol. Microbiol. Biotechnol.* **8**:243-254. 2004.
- *Appl. Microbiol. Biotechnol.* **68**:475-480. 2005.
- *Appl. Environ. Microbiol.* **72**:3418-3428. 2006.
- *Appl. Environ. Microbiol.* **73**:2349-2353. 2007.
- *Microbiology* **153**:2491-2504. 2007.
- *Appl. Microbiol. Biotechnol.* **77**:853-860. 2007.
- *Appl. Microbiol. Biotechnol.* **77**:1053-1062. 2008.
- *Appl. Microbiol. Biotechnol.* **78**:449-454. 2008.
- *Appl. Environ. Microbiol.* **74**:5146-5152. 2008.
- *Appl. Microbiol. Biotechnol.* **81**:459-464. 2008.
- *Appl. Microbiol. Biotechnol.* **81**:505-513. 2008.
- *Appl. Microbiol. Biotechnol.* **81**:691-699. 2008.
- *Appl. Microbiol. Biotechnol.* **85**: 105-115. 2009
- *Appl. Microbiol. Biotechnol.* **85**:471-480. 2010.(Mini-Review)

Chromosome engineering methods

- *Appl. Environ. Microbiol.* **71**:407-416. 2005.
- *Microbiology* **151**:501-508. 2005.
- *Appl. Microbiol. Biotechnol.* **67**:225-233. 2005.
- *Appl. Environ. Microbiol.* **71**:3369-3372. 2005.
- *J. Mol. Microbiol. Biotechnol.* **8**:243-254. 2005.
- *Appl. Environ. Microbiol.* **71**:7633-7642. 2005.(Review)
- *Appl. Environ. Microbiol.* **71**:8472-8480. 2005.
- *Appl. Microbiol. Biotechnol.* **69**:151-161. 2005.
- *Appl. Environ. Microbiol.* **72**:3750-3755. 2006.
- *Appl. Microbiol. Biotechnol.* **74**:1333-1341. 2007.
- *Biosci. Biotechnol. Biochem.* **71**:1683-1690. 2007.
- *Appl. Microbiol. Biotechnol.* **77**:871-878. 2007.
- *Appl. Microbiol. Biotechnol.* **79**:519-526. 2008.(Mini-Review)

Thank you for your attention

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