



# **Fructophilic Lactic Acid Bacteria for Honey Bee Health**

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# What is fructophilic LAB (FLAB)???

Fructophilic --- fructose + philic = fructose loving

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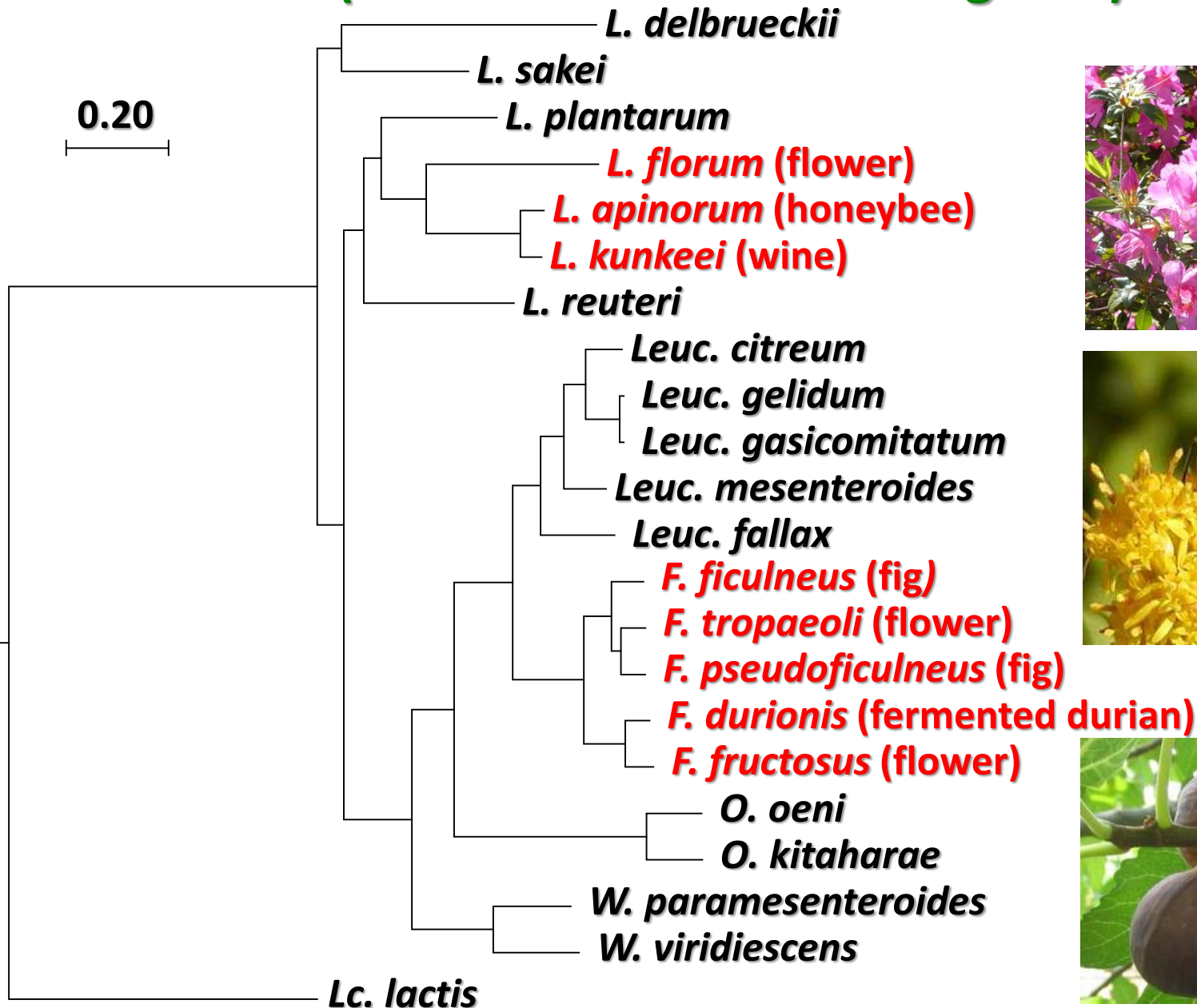
Reclassification of the genus *Leuconostoc* and proposals of *Fructobacillus fructosus* gen. nov., comb. nov., *Fructobacillus durionis* comb. nov., *Fructobacillus ficulneus* comb. nov. and *Fructobacillus pseudoficulneus* comb. nov.

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## **Description of *Fructobacillus* gen. nov.**

*Fructobacillus* (Fruc.to.ba.cil'lus. N.L. masc. n. *Fructobacillus* arbitrarily derived from fructose and *Lactobacillus*, intended to mean fructose-loving lactic acid-producing bacillus).

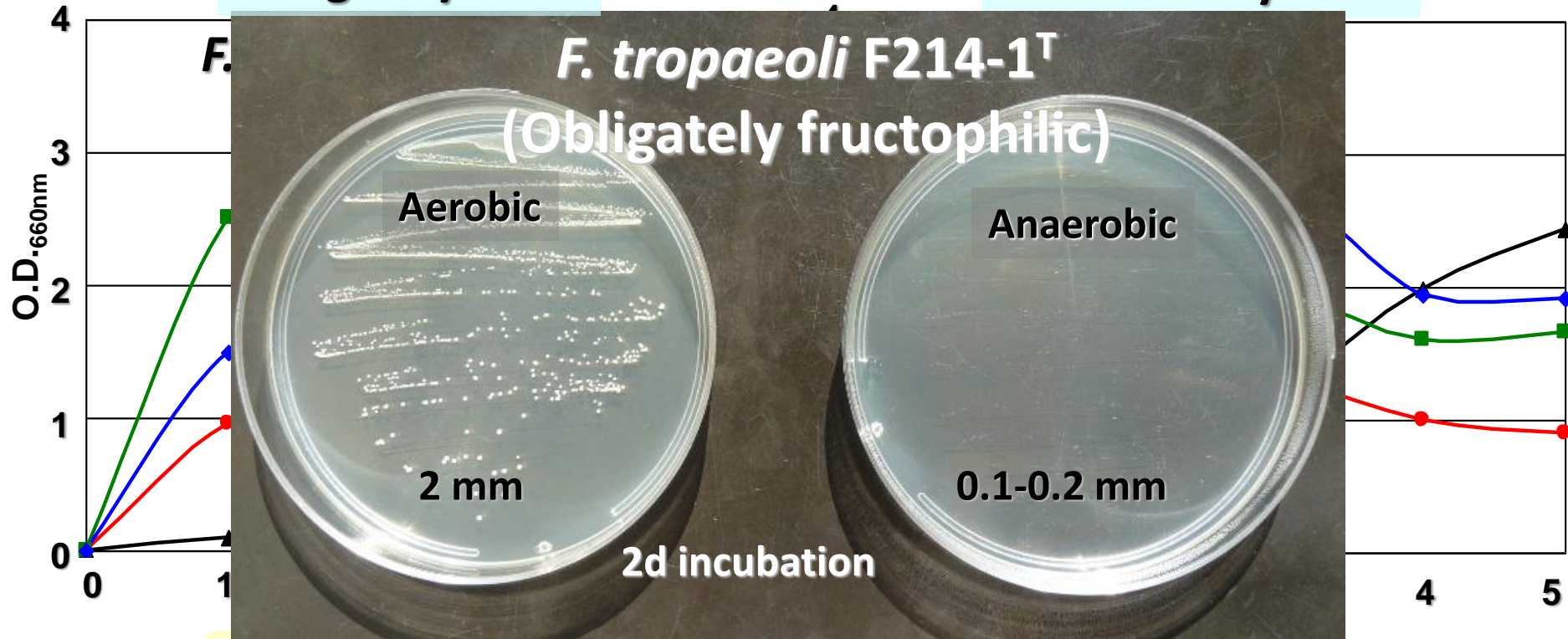
# Phylogenetic relationships between FLAB and other LAB (based on 235 conserved genes)



# Growth characteristics of FLAB

## Obligately FLAB

## Facultatively FLAB

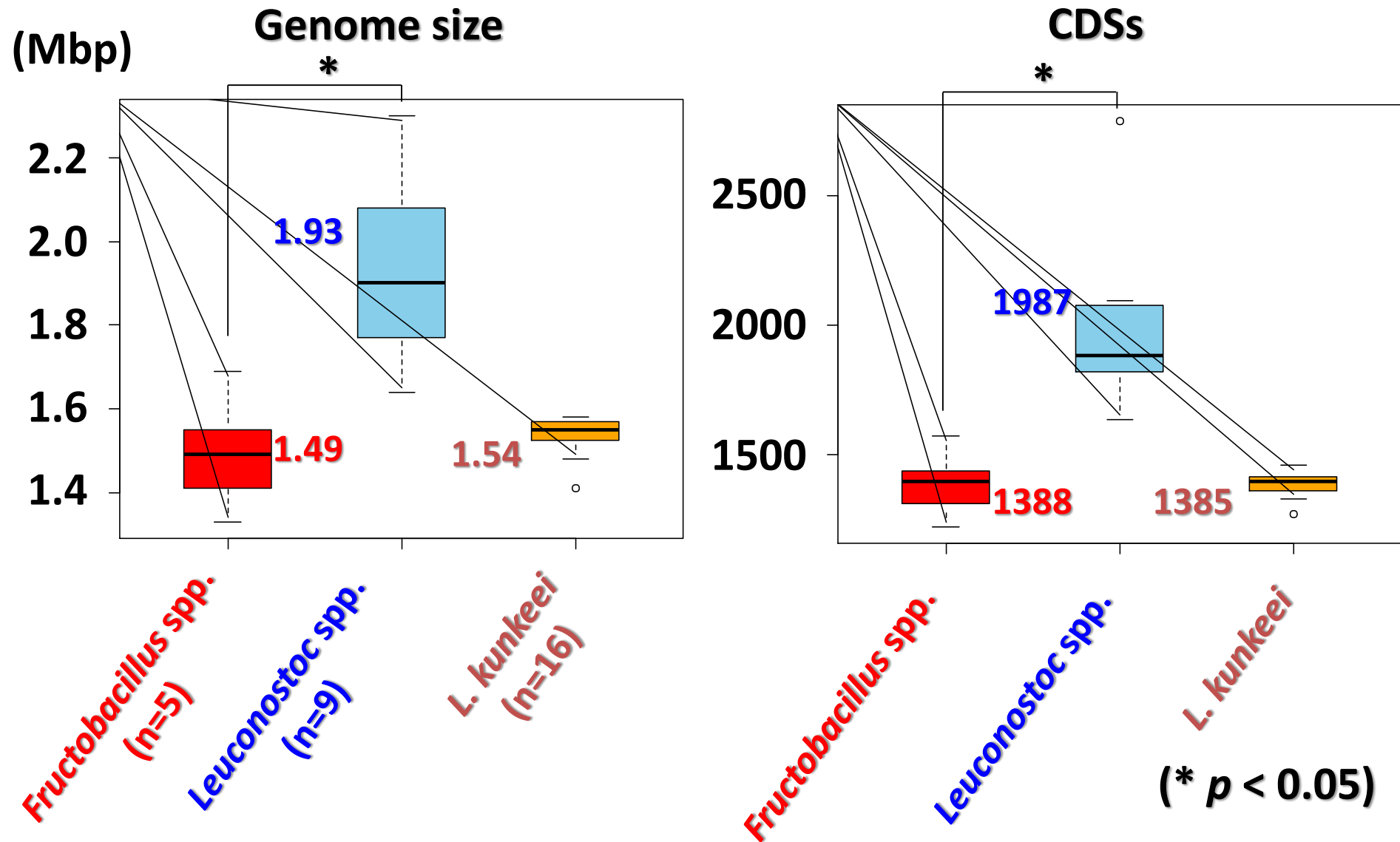


*Fructobacillus* spp.  
*Lactobacillus kunkeei*  
*Lactobacillus apinorum*

*Lactobacillus florum*

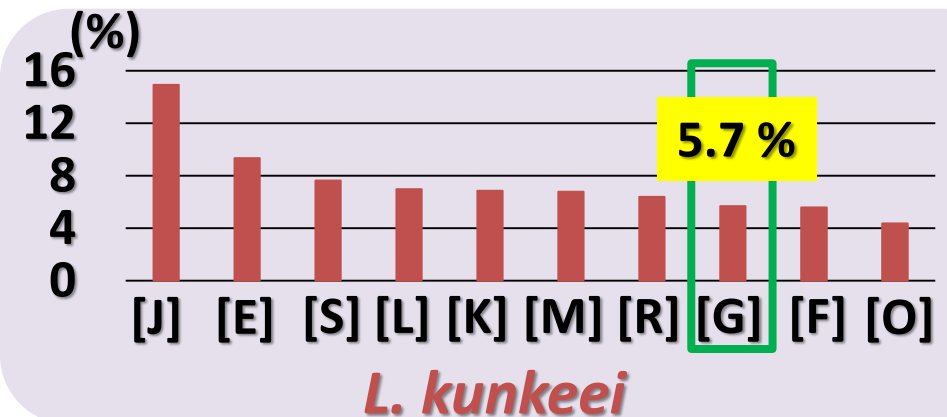
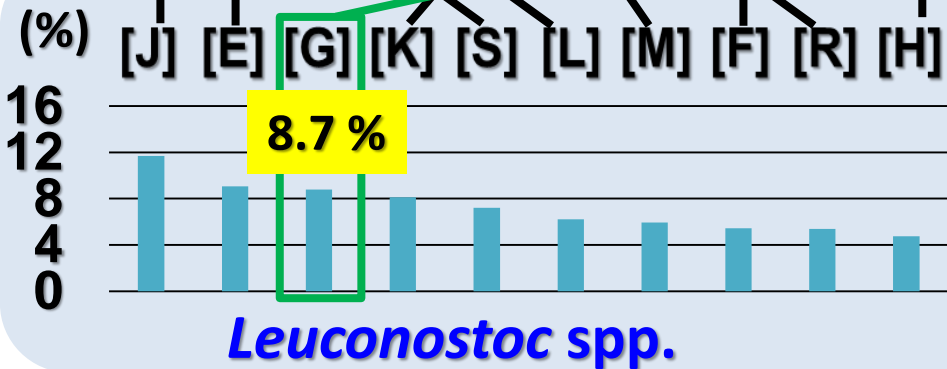
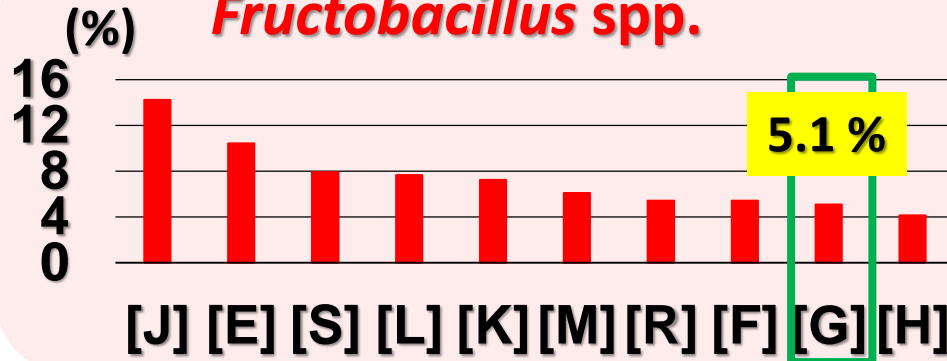


# Genome size and number of CDSs in *Fructobacillus* spp., *Leuconostoc* spp. and *L. kunkeei*



# Top 10 COG classes among the groups

## *Fructobacillus spp.*



## *L. kunkeei*

- [C] Energy production and conversion
- [D] Cell cycle control, cell division etc.
- [E] Amino acid transport and metabolism
- [F] Nucleotide transport and metabolism
- [G] Carbohydrate transport and metabolism**
- [H] Coenzyme transport and metabolism
- [I] Lipid transport and metabolism
- [J] Translation and ribosomal structure etc.
- [K] Transcription
- [L] Replication, recombination and repair
- [M] Cell wall membrane etc.
- [N] Cell motility
- [O] Posttranslational modification etc.
- [P] Inorganic ion transport and metabolism
- [Q] Secondary metabolites biosynthesis etc.
- [R] General function prediction only
- [S] Function unknown
- [T] Signal transduction mechanisms
- [U] Intracellular trafficking, secretion etc.
- [V] Defense mechanisms
- [X] Mobilome: prophages, transposons

# Discriminative pathways among the groups

*illus spp.*

*c spp.*

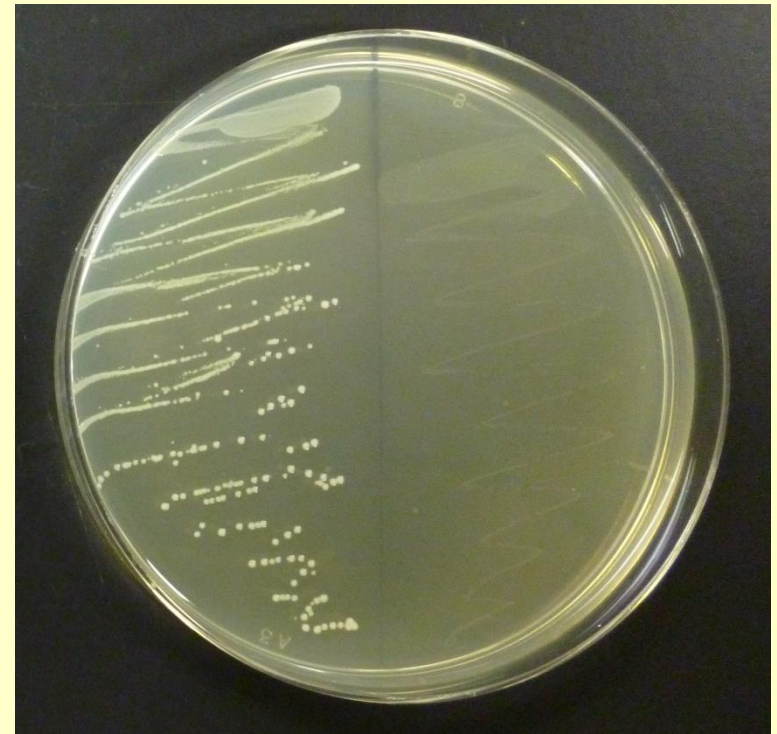
## Aerobic

*Leu. mesenteroides*    *F. fructosus*



## Anaerobic

*Leu. mesenteroides*    *F. fructosus*



n

# Fructobacillus-specific- & Leuconostoc-specific-genes

## Fructobacillus-specific genes

4 ≤ : *Fructobacillus* (n=5)

2 ≥ : *Leuconostoc*

16 genes

- Smaller genome
- Less CDS
- Simple metabolic systems

Alcohol dehydrogenase

NAD(P)H dehydrogenase

Chloride peroxidase

Levansucrase

Acylaminoacyl-peptidase

Xanthine oxidase

etc.

## Leuconostoc-specific genes

1 ≥ : *Fructobacillus* (n=5)

7 ≤ : *Leuconostoc* (n=9)

114 genes

Pyruvate dehydrogenase E1 component

Pyruvate dehydrogenase E2 component

Transketolase

β-fructofuranosidase

aminopeptidase

Xaa-Pro dipeptidase

etc.

Acetaldehyde/alcohol dehydrogenase (*adhE*)

PTS components

Teichoic acid transport system components

etc.

**Reductive evolution in *Fructobacillus* spp.**



# Different habitats among FLAB species

**Flowers**

**Grapes and wine**

*L. florum*

*L. kunkeei*

*F. fructosus*

**Insects**

*L. apinorum*

*F. tropaeoli*

*F. pseudoficulneus*

*F. ficulneus*

*F. durionis*

**Fruits and fruits fermentation  
(other than grapes and wine)**



# Isolation of FLAB from honeybees and beehives



66 strains of LAB  
in the initial  
screening



All were FLAB.

Only 14/354 (4%) strains  
from flowers and fruits  
were FLAB.

**Genetic group  
by rep-PCR**

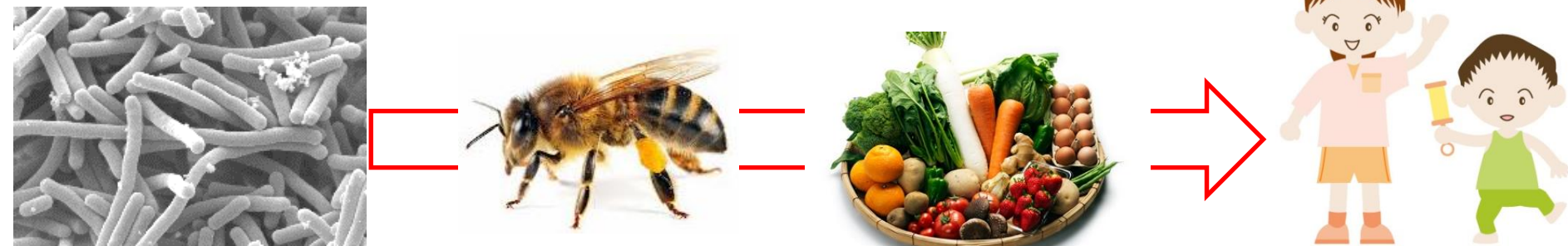
**Origin of grouped strains**

|     |  |                             |
|-----|--|-----------------------------|
| I   | flower (10)  | <i>F. fructosus</i> (99.7%) |
| II  | <i>A. mellifera</i> -SW (3), hybridized bee-SW (2), honey (7), larvae (7), bee bread (4), <i>A. ligustica</i> -Å (6)   | <i>L. kunkeei</i> (100%)    |
| III | <i>A. mellifera</i> -SW (3)  | <i>F. fructosus</i> (99.2%) |
| IV  | <i>A. mellifera</i> -SW (2), <i>A. ligustica</i> -SW (4), hybridized bee-SW (2), honey (4), larvae (11), bee bread (8) | <i>L. kunkeei</i> (100%)    |
| V   | honey (2)  | <i>L. kunkeei</i> (99.9%)   |
| VI  | honey (1)  | <i>L. kunkeei</i> (99.6%)   |

# Summary

## FLAB are

- newly discovered microorganisms.
- adapted to their habitats at the genomic level.
- members of core microbiota in crop and diet of honeybees.
- promising candidates for bee probiotics.
- promising symbionts for a vector of paratransgenesis.



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