



Composition and diversity of **bacterial communities** in the gut of black soldier fly larvae (*Hermetia illucens* L.) **reared** **on organic wastes**

Presented By

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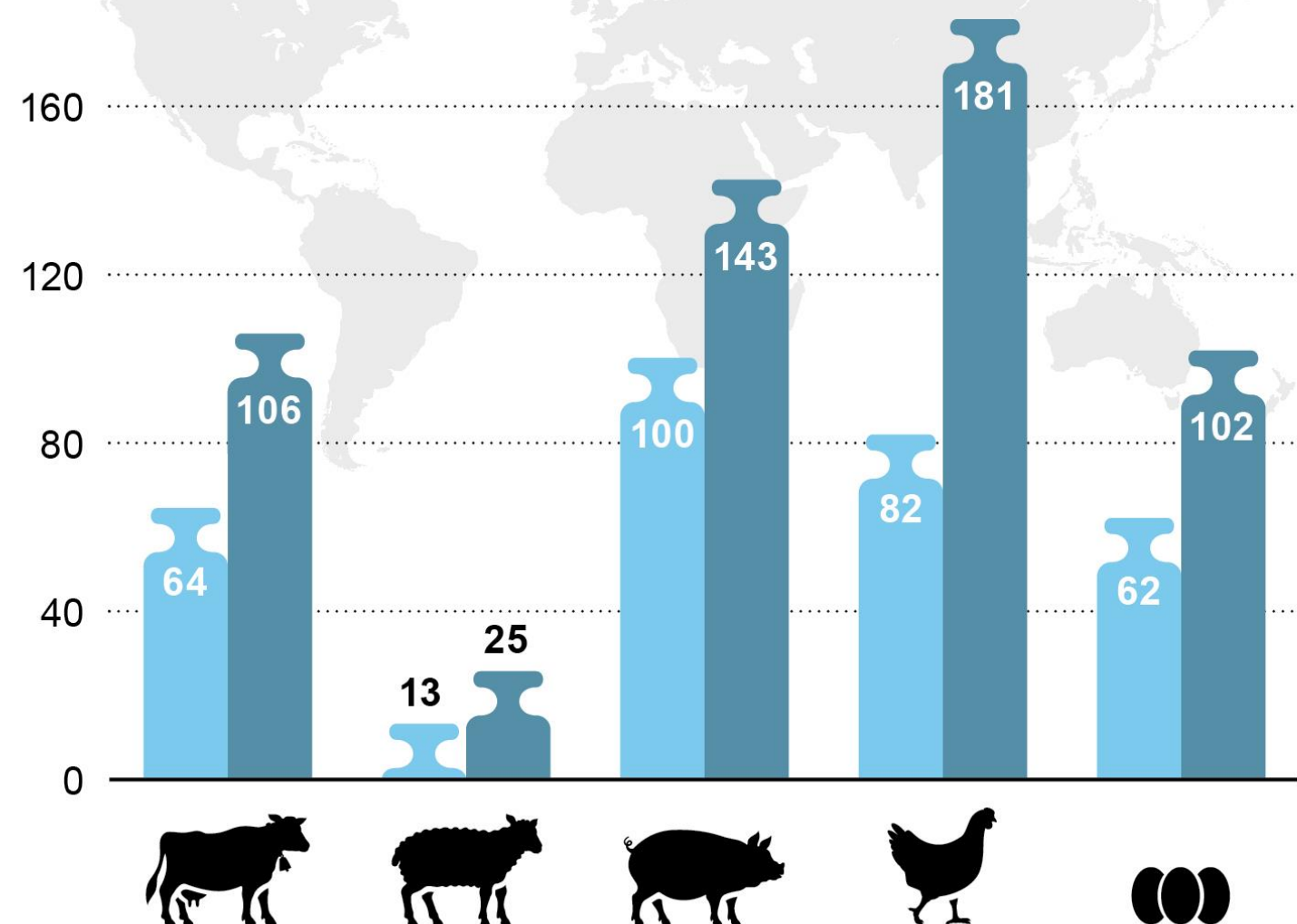
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INTRODUCTION

Global demand for meat
2005 versus 2050 in million tonnes



Source: Food and Agriculture Organization of the United Nations,
ESA Working paper No. 12-03, p. 131

© blue horizon

Source: <https://bluehorizon.com/insight/high-increase-in-demand-per-meat-category-by-2050-show-chicken-with-the-largest-increase-and-total-volume/>

➤ **Rising global population drives higher consumption of animal-based protein.** (United Nations, 2019)

➤ **Increased livestock & feed demand emphasize the need for sustainable feed sources.** (Lu et al., 2022)



Source: <https://www.four-paws.org/campaigns-topics/topics/farm-animals/10-amazing-facts-about-pigs>



Source: <https://www.tinytailstoyou.com/animals/chicks>



Source: <https://beef.unl.edu/beefwatch/2020/how-many-pounds-meat-can-we-expect-beef-animals/>



Source: <https://www.farmsanctuary.org/news-stories/ten-facts-about-sheep/>



Protein-rich edible insects: a sustainable protein alternative.

(Nowak et al., 2016)

BSFL; Black soldier fly larvae



Saprophagous insect:

It converts **low-value organic waste**.



Source: https://www.livinginsider.com/profile/65246/member_info.html



Source: <https://www.sanook.com/women/229989/>



Source: <https://www.scghome.com/living-ideas/articles>



Source: <https://growlandscapes.com.au/product/chook-manure/>



... Nutrient-rich **“insect biomass”**



BSFL; Black soldier fly larvae



Source: https://www.freepik.com/premium-photo/close-up-black-soldier-fly-bsf-larvae-maggot-hermetia-illucens_133636932.htm

- **Protein** 35-49%
- **Fat** 29-35%
- **Energy** 2,900 Kcal/Kg
- **Amino acid profile** = fish meal

“Dietary supplement” for livestock animals.



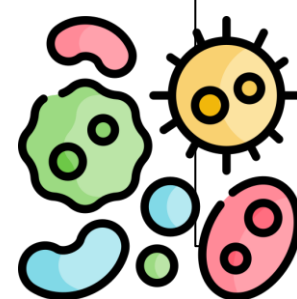
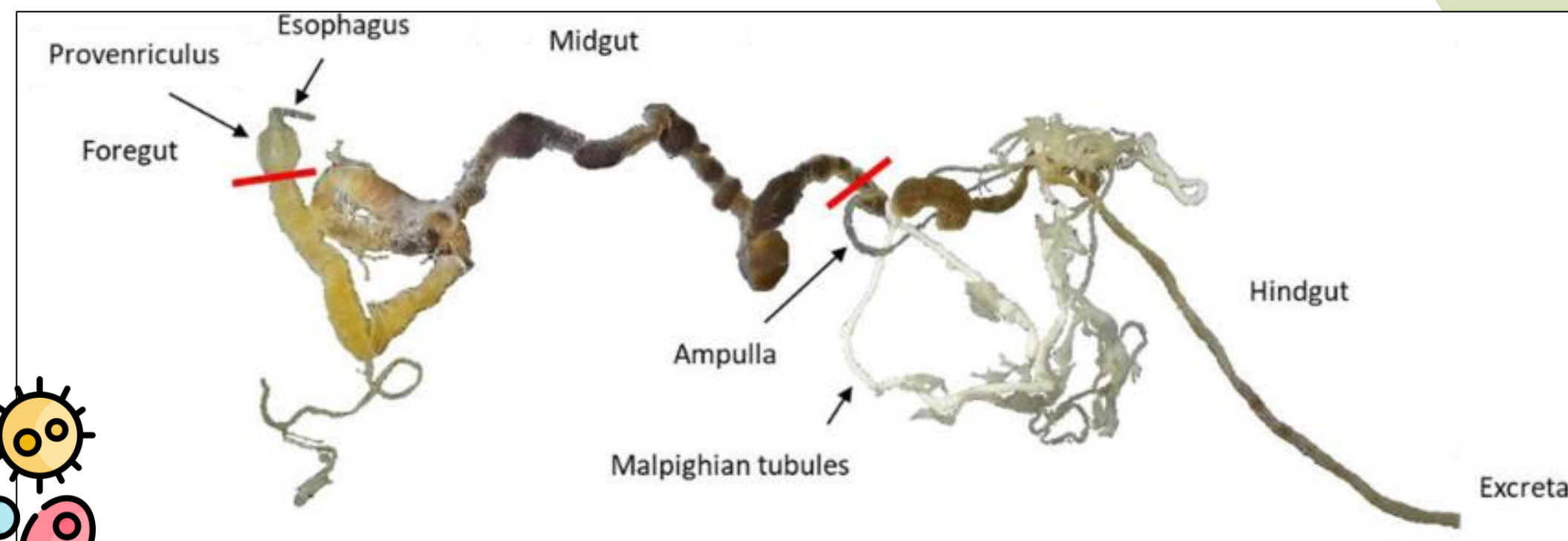
Effective growth of BSFL as a protein source depends on “substrates”.

(Gligorescu et al., 2018)

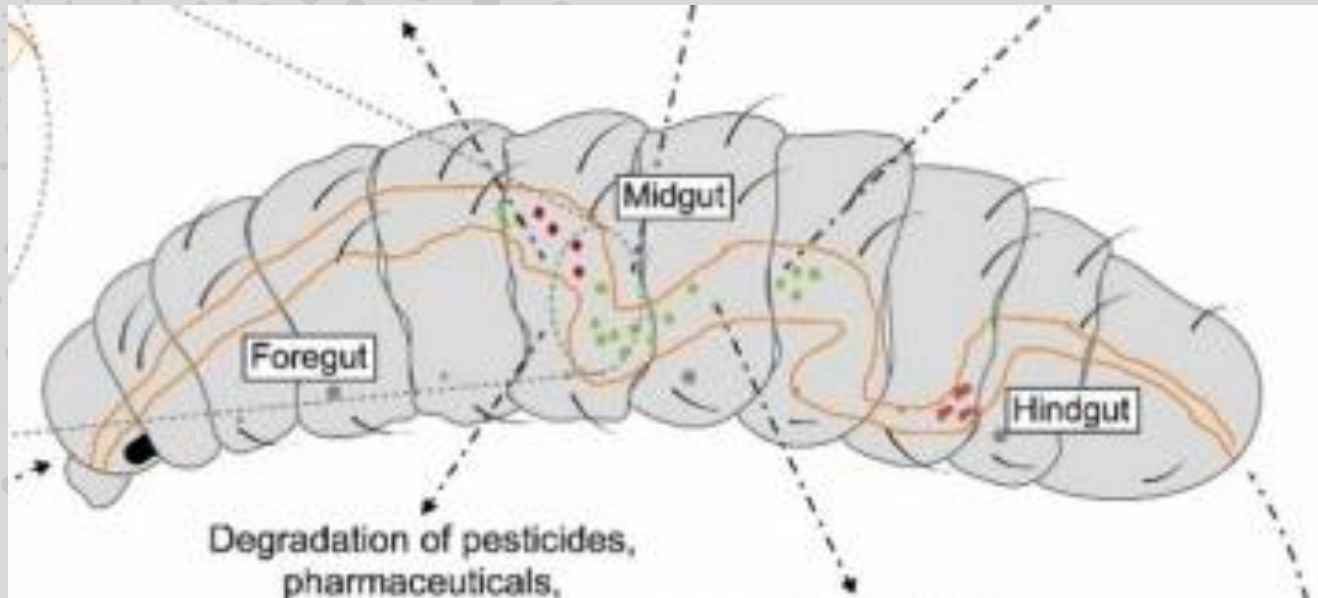
✓ Gut microbiota (Chen et al., 2023)



Source: <https://flukerfarms.com/reptile-u/blog/black-soldier-fly-larvae-soldierworm-grubs-/>



- Digestion
- Antimicrobial peptide (AMPs) synthesis
- Overall insect health (Silvaraju et al., 2024)



Other factors

●
affect **changes** in gut microbiota.

➤ Biotic factors

- Developmental **stage**
- **Genetic** background
- **Gut regions**
- Initial **substrate microbes**

➤ Abiotic factors

- Geographic **locations**
- **Temperature**
- Substrates **pH**
- **Density**



- ✓ **AMPs properties** of BSFL & its gut microbiota: lead to alternative valorization of the BSF system.



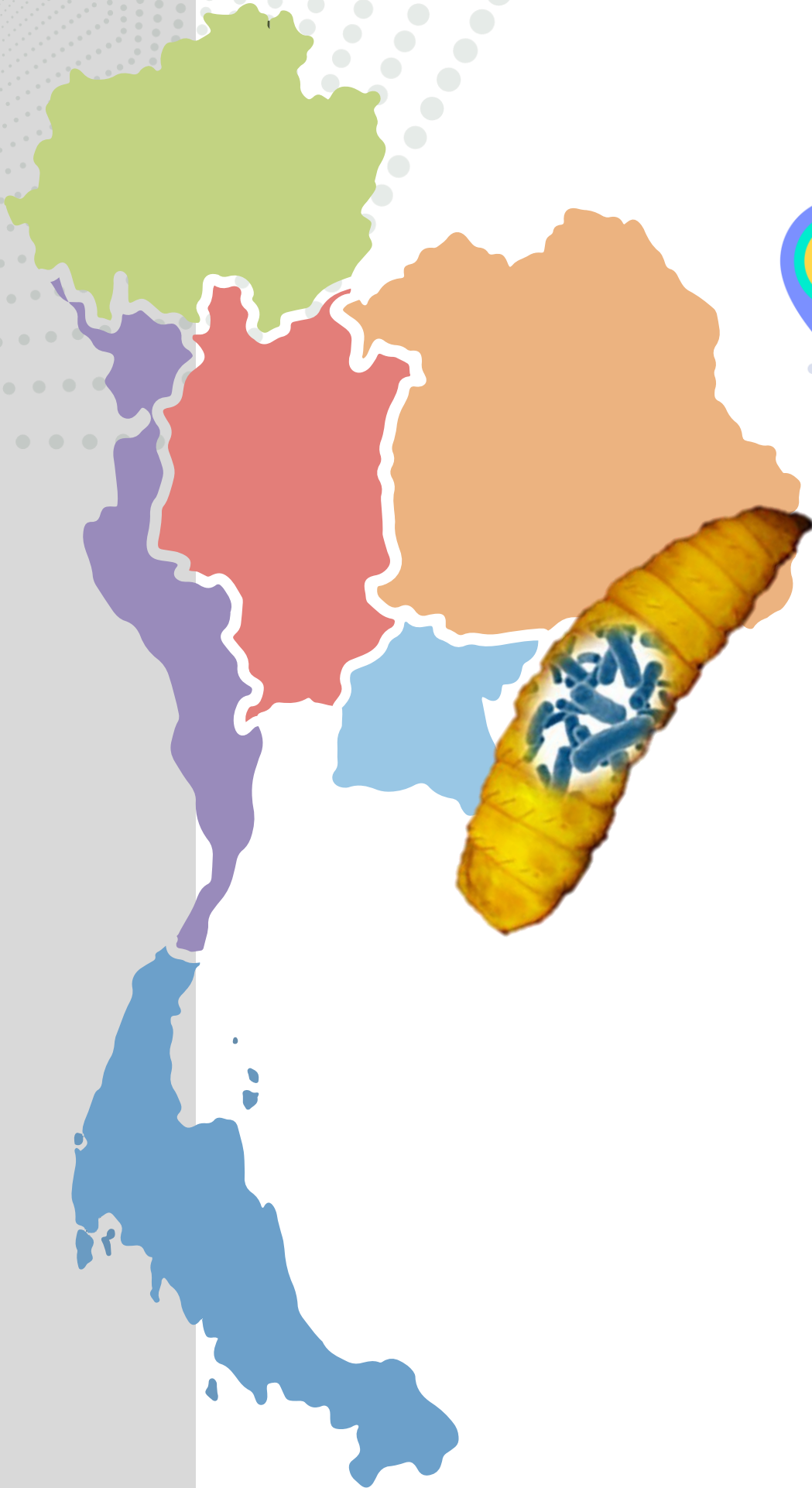
Bruno et al. (2019) & Khamis et al. (2020)

- **BSFL** can serve as reservoirs for various significant **foodborne pathogens**.

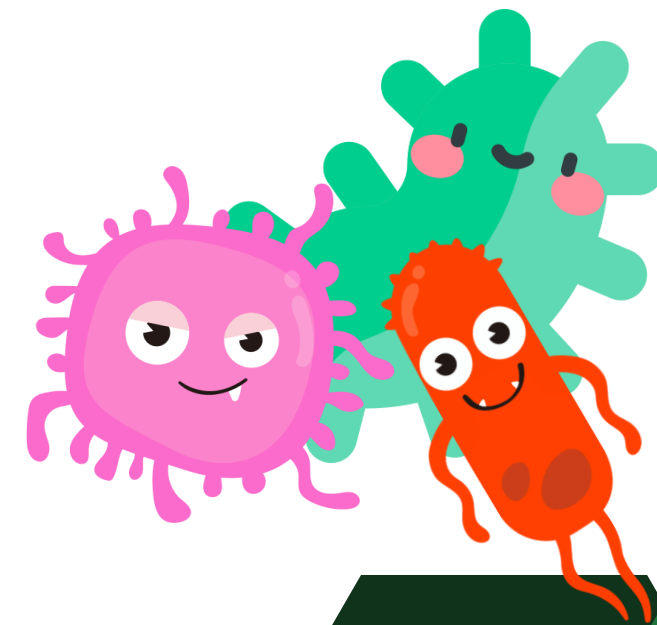


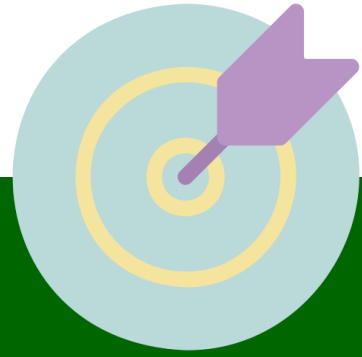
Biosafety assessments should be conducted.

- **Substrates** = “**Microbial contamination**” resulted in a significant shift in the **BSFL gut microbiota**. (Chen et al., 2023)



Bacterial communities in the gut of BSFL reared on organic wastes that locally available in **Thailand remains unexplored.**





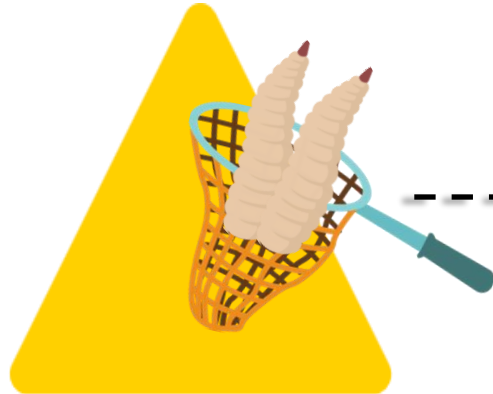
OBJECTIVES



To investigate effect of substrates on the **gut bacterial communities** of BSFL.



METHODOLOGY



1

Collecting **BSFL** samples



2

Bacterial **culture** method



3

Bacterial **non-culture** method

1 Collecting BSFL samples



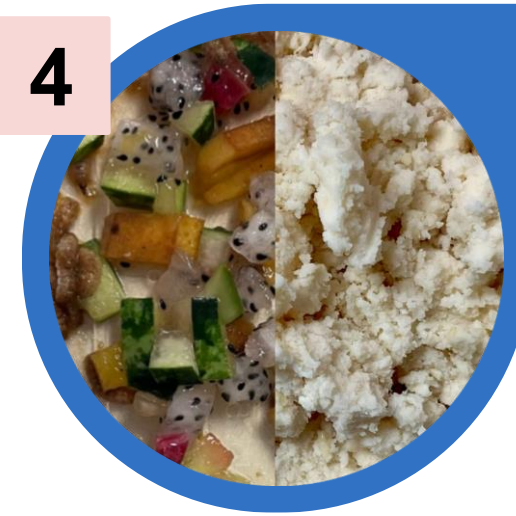
FOW; Food waste



FSM; Food waste 1:
Supermarket waste 1



FRW; Fruit waste



FSB; Fruit waste 1:
Soybean residue 1



CHM; Chicken manure



STD; Control diets
(chicken feed 1: soybean residue 1)

1 Collecting BSFL samples



▶ Larvae **13-17** days old.

▶ **Dark brown** color.

2 Bacterial culture method

2.1 Gut sample preparation



Dissection of BSFL gut
& grinding with solution.

2.2 Bacteria **cultivation**



- 30 °C
- 24–48 h



“**Observation**” of colony morphology

- Form
- Margin
- Elevation
- Color
- Gram staining



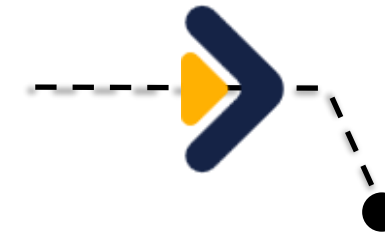
2 Bacterial culture method

2.3 Bacteria DNA Extraction



Bacteria DNA Extraction

- Invitrogen PureLink™ Genomic DNA Mini Kit



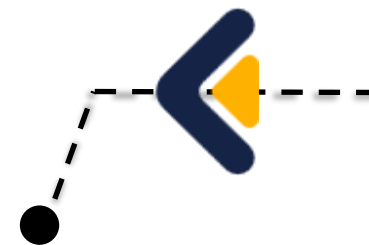
PCR



Target gene: **16s rRNA region**

Specific primers:

- **27F:** 5-AGAGTTTGATCMTGGCTAG-3
- **1513R:** 5-ACGGYTACCTTGTTACGACTT-3



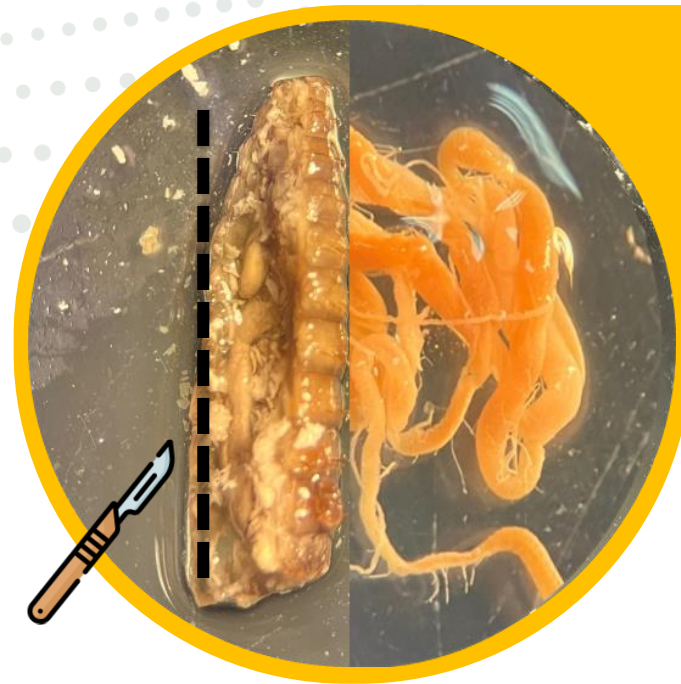
Bacterial identification

Nucleotide sequence analysis

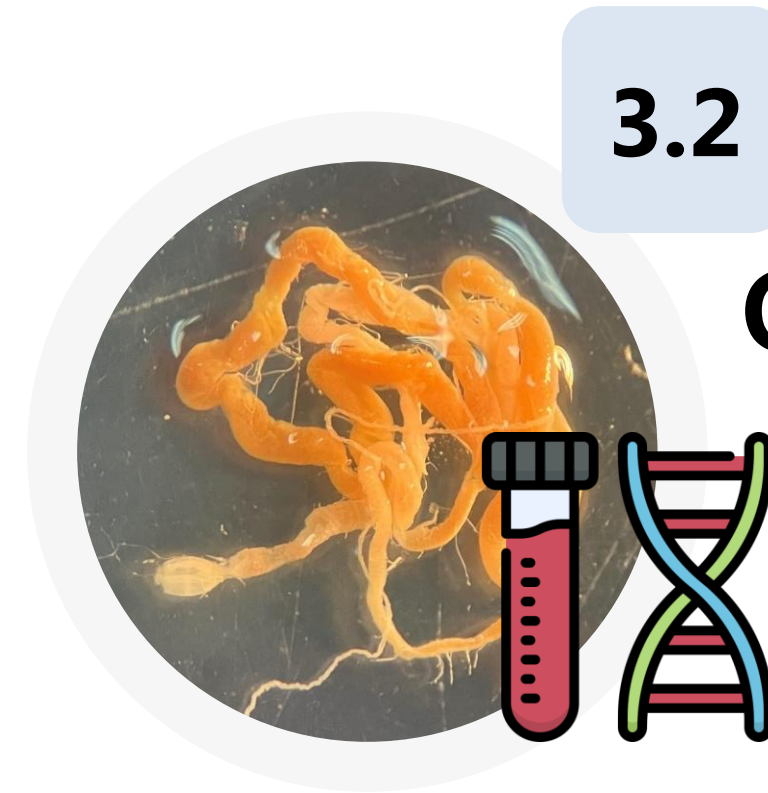
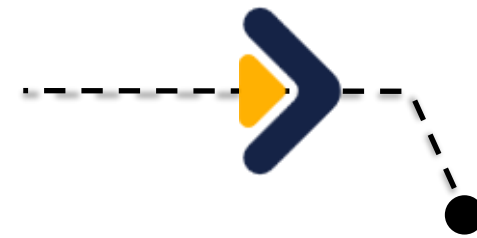


Source: <https://blast.ncbi.nlm.nih.gov/Blast.cgi>

3 Bacterial non-culture method



3.1 Dissection of BSFL gut



3.2

Gut **DNA Extraction**

- Invitrogen PureLink™ Genomic DNA Mini Kit

Metagenomics technique



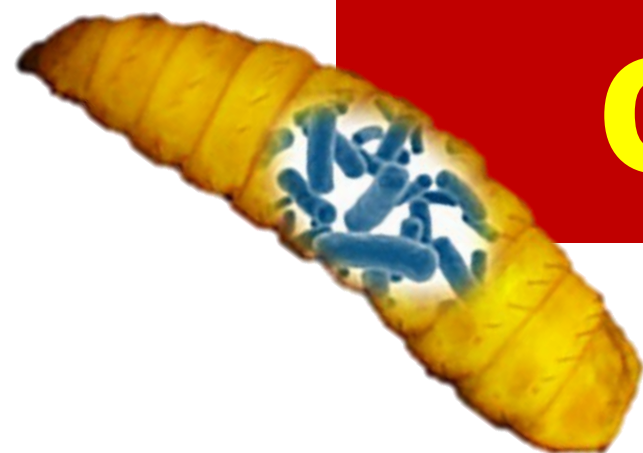
3.3

- Bacterial diversity analysis



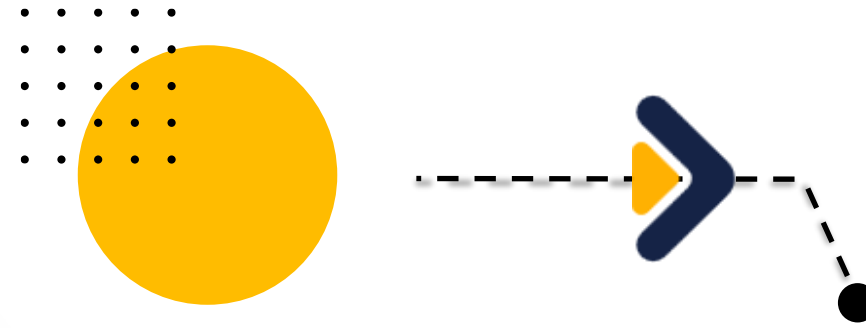
RESULTS & DISCUSSION

“Bacterial communities:
culture method”





“Bacterial culture method”

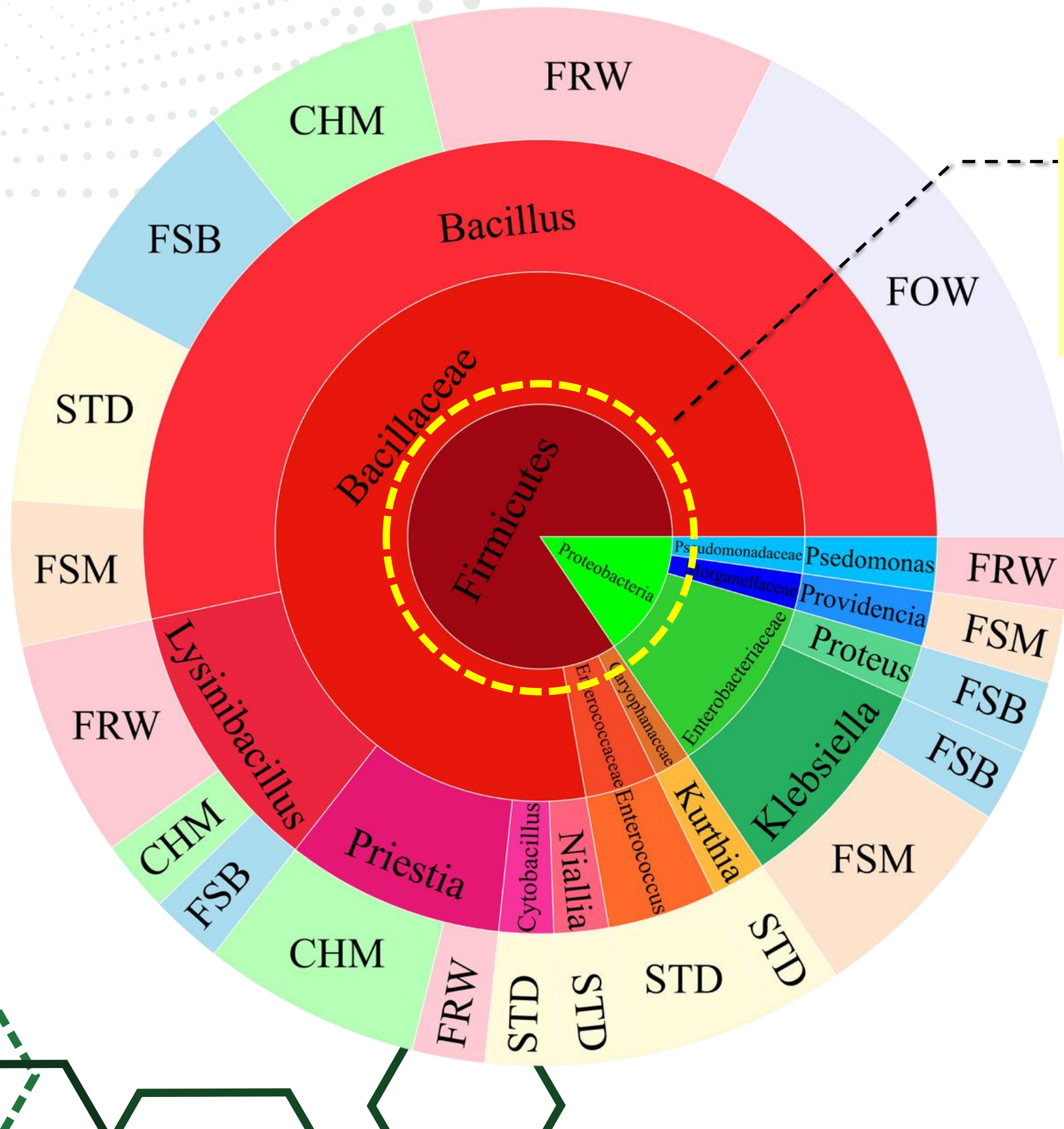


A total of **45 isolates** were obtained from different substrates, selected based on distinct colony morphologies.

- STD & FOW: 8 isolates
- FSM & FSB: 6 isolates
- FRW: 10 isolates
- CHM: 7 isolates



“Bacterial culture method” - Phylum level

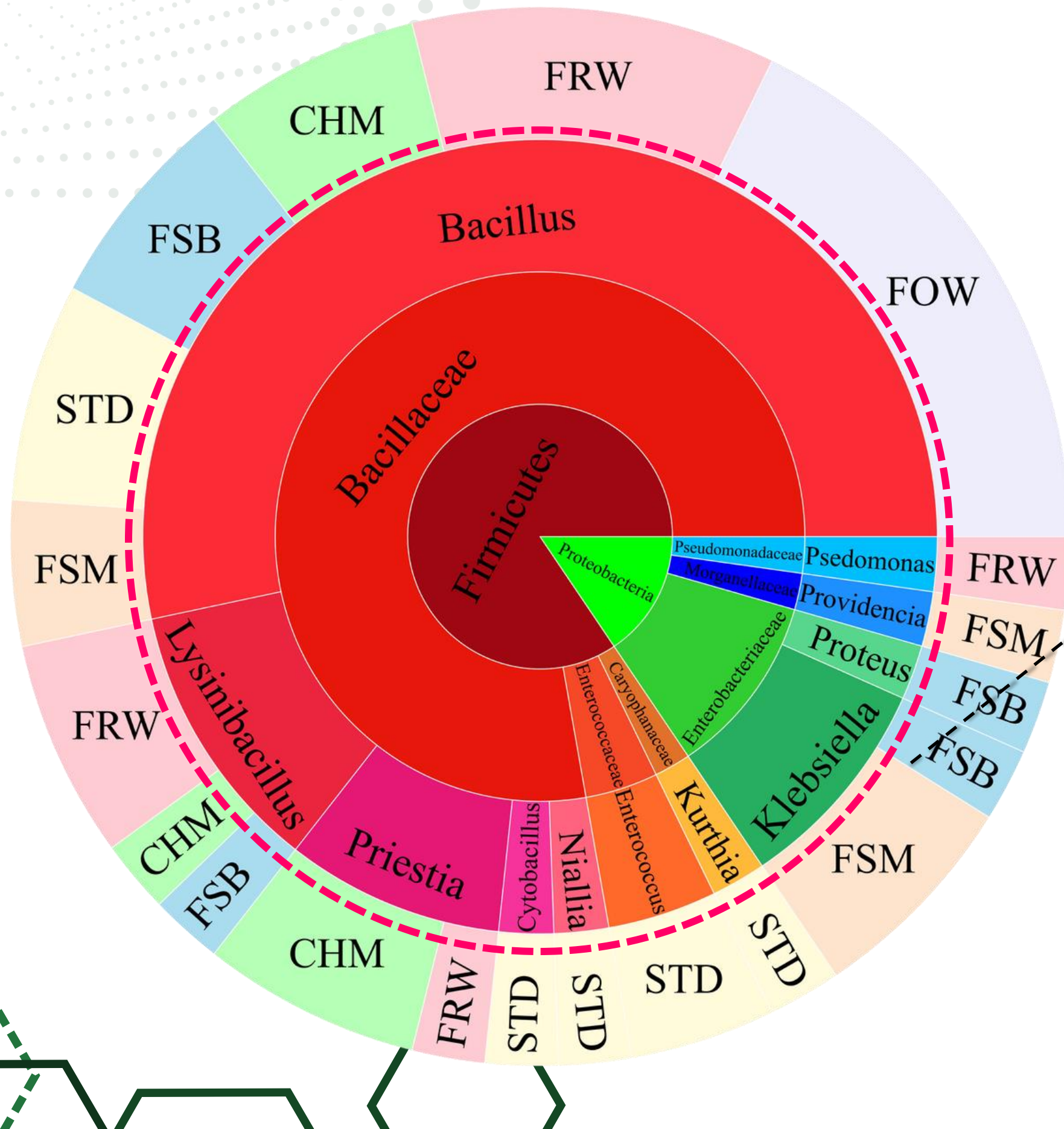


✓ 2 Phyla

- **Firmicutes:** 6 substrates
- **Proteobacteria:** FSM, FRW & FSB



“Bacterial culture method” - Genus level

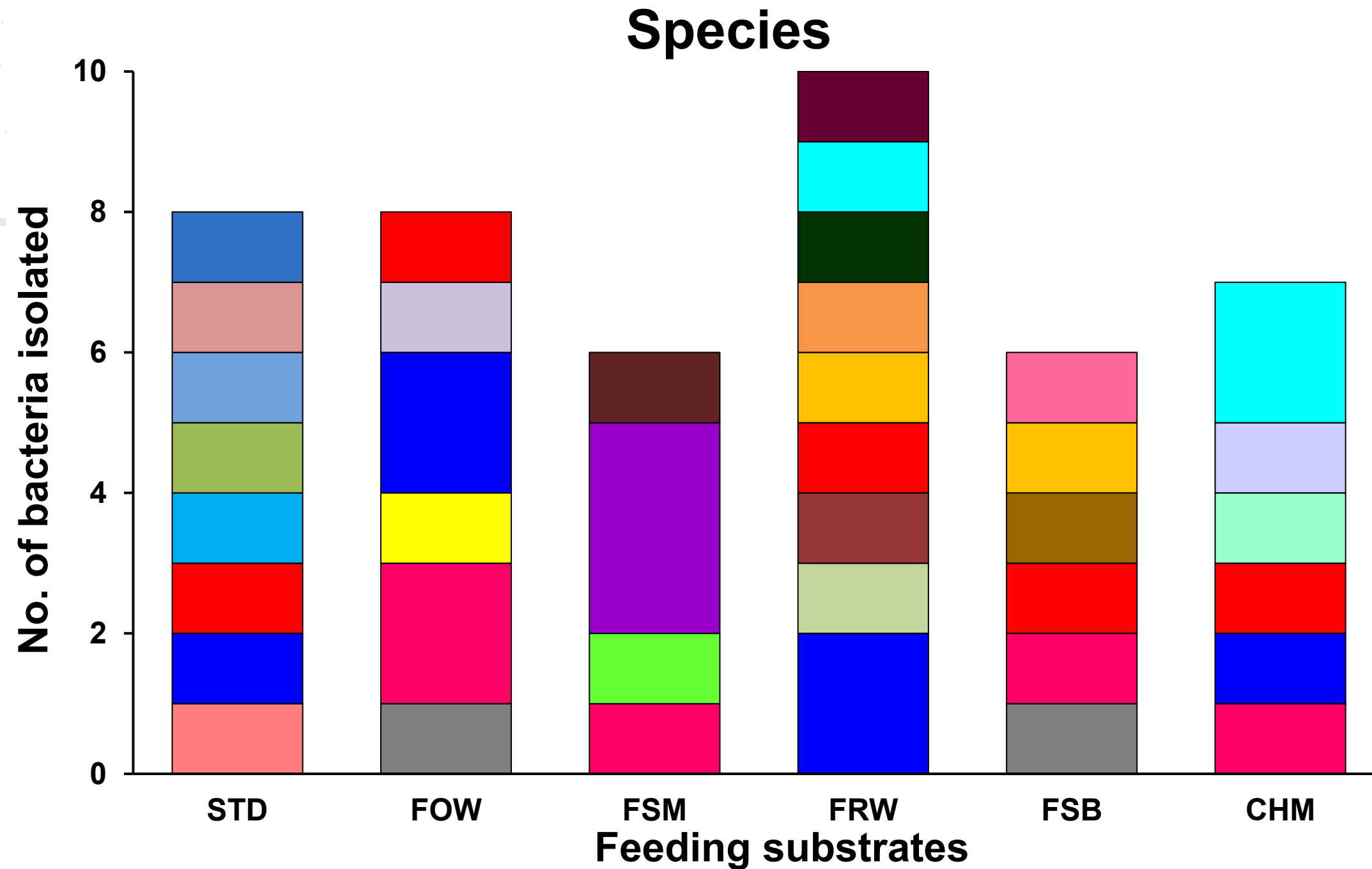


✓ 11 Genera

- ***Bacillus***: 6 substrates
- ***Lysinibacillus***: FRW, FSB & CHM
- ***Klebsiella***: FSM & FSB
- ***Enterococcus***: STD
- ***Providencia***: FSM
- ***Pseudomonas***: FRW



“Bacterial culture method” - Species level



✓ 28 Species

- *B. cereus*
- *B. amyloliquefaciens*
- *B. velezensis*

FRW: the highest diversity. (9 species)

- | | | | |
|-------------------------|-----------------------------|-------------------------|-------------------------|
| <i>B. altitudinis</i> | <i>B. amyloliquefaciens</i> | <i>B. anthracis</i> | <i>B. atrophaeus</i> |
| <i>B. cereus</i> | <i>B. paramycoides</i> | <i>B. safensis</i> | <i>B. subtilis</i> |
| <i>B. thuringiensis</i> | <i>B. velezensis</i> | <i>C. kochii</i> | <i>E. faecalis</i> |
| <i>E. termitis</i> | <i>K. aerogenes</i> | <i>K. pneumoniae</i> | <i>K. gibsonii</i> |
| <i>L. fusiformis</i> | <i>L. macroides</i> | <i>L. mangiferihumi</i> | <i>L. pakistanensis</i> |
| <i>N. circulans</i> | <i>P. aryabhattai</i> | <i>P. megaterium</i> | <i>P. mirabilis</i> |
| <i>P. rettgeri</i> | <i>P. putida</i> | | |

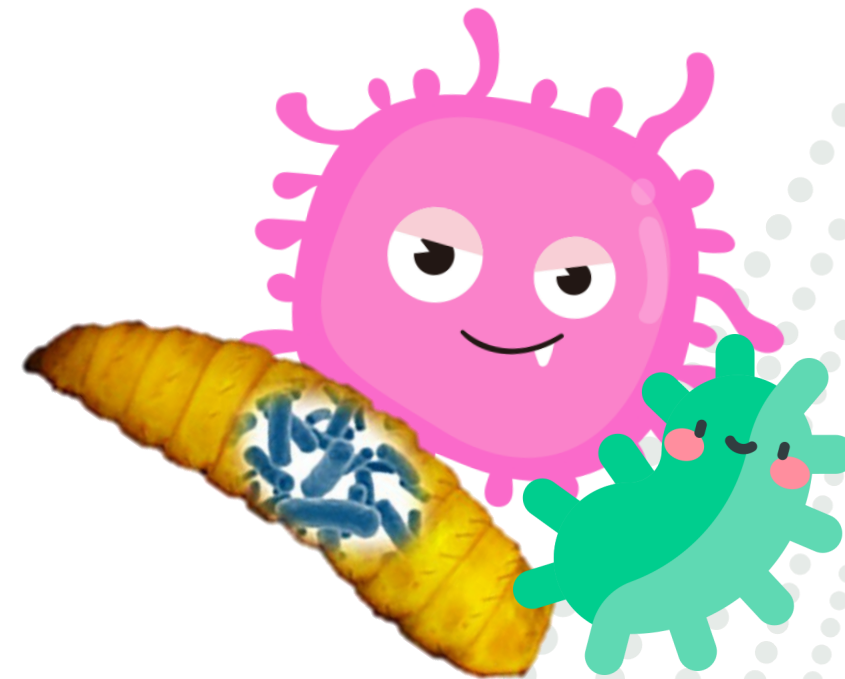
Dominate bacteria....



genus *Bacillus*

(Shumo et al., 2021; Pei et al., 2024; Van Kessel et al., 2024)

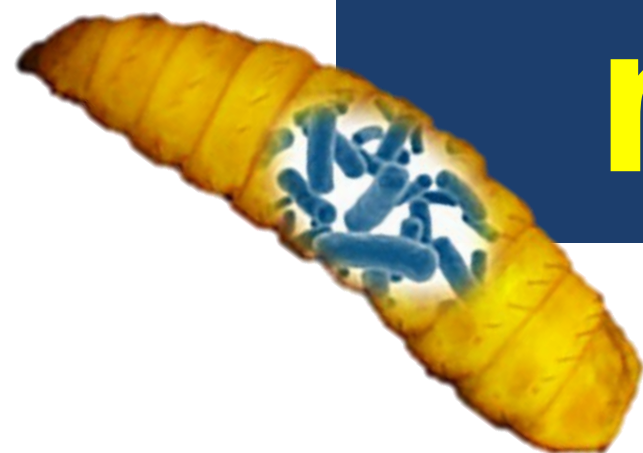
- **Enhances production of digestive enzymes.**
(protease, amylase, cellulase & lipase)
- **Antibiotic synthesis**





RESULTS & DISCUSSION

“Bacterial communities:
non-culture method”





“Bacterial non-culture method”

Table 1 Summary on raw data processing and alpha diversity of the gut bacterial communities of BSFL fed on different substrates.

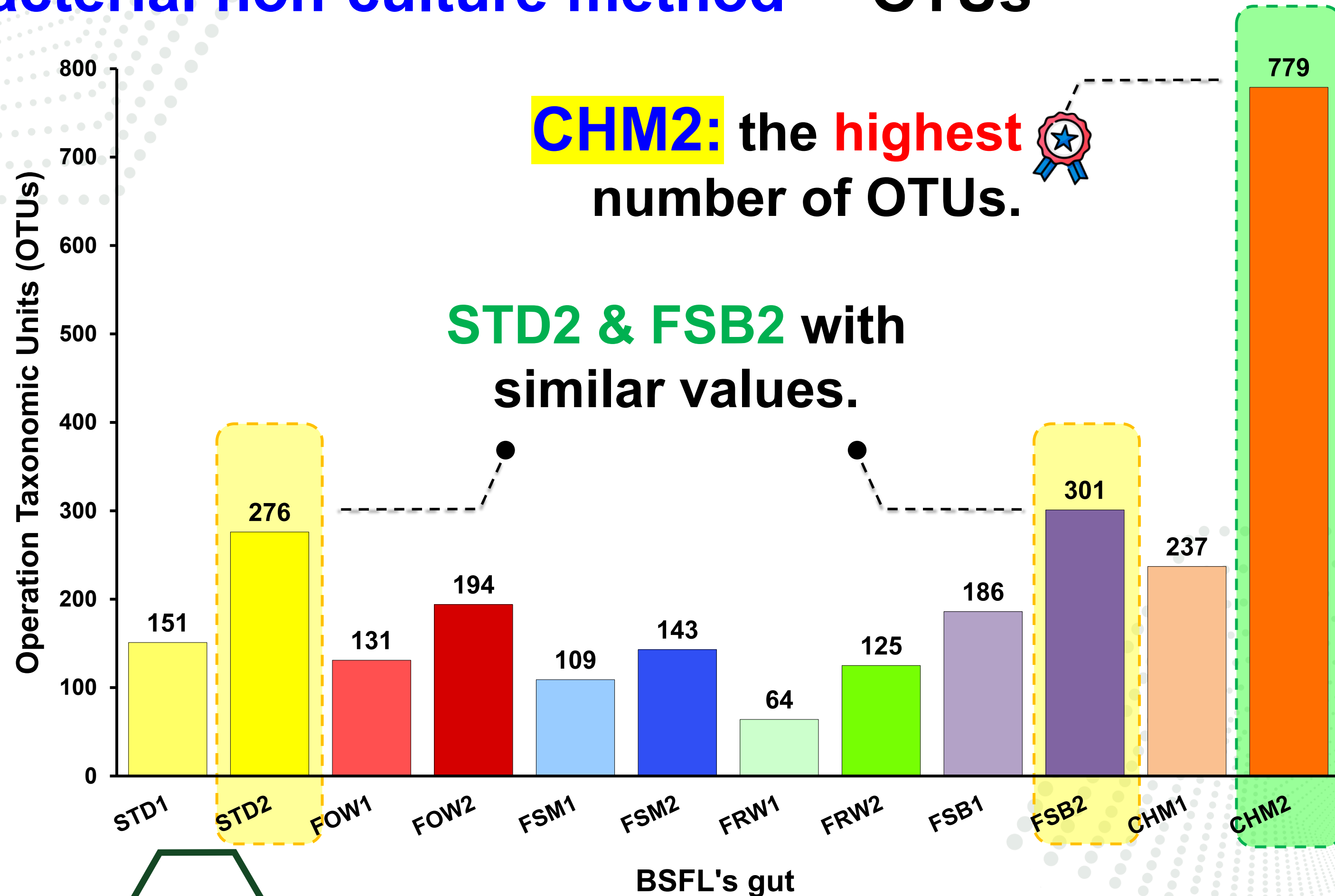
Feeding substrates	Raw reads	Clean reads	Shannon	Simpson	Choa1
STD1	79,876	74,408	4.70	0.94	151.27
✓ STD2	80,363	75,172	5.12	0.95	276.02
FOW1	80,019	73,436	2.12	0.53	131.18
FOW2	79,837	74,809	2.79	0.76	194.00
FSM1	80,406	74,215	2.13	0.55	109.03
FSM2	79,960	75,158	3.19	0.83	143.03
FRW1	80,186	74,136	2.09	0.55	64.00
FRW2	79,898	75,265	3.35	0.75	125.04
FSB1	79,893	73,494	4.88	0.92	186.03
✓ FSB2	80,234	75,521	3.70	0.74	301.01
CHM1	79,886	73,818	3.69	0.79	237.00
★ CHM2	80,141	76,095	6.20	0.97	779.00

CHM2: the **highest** gut bacterial diversity.



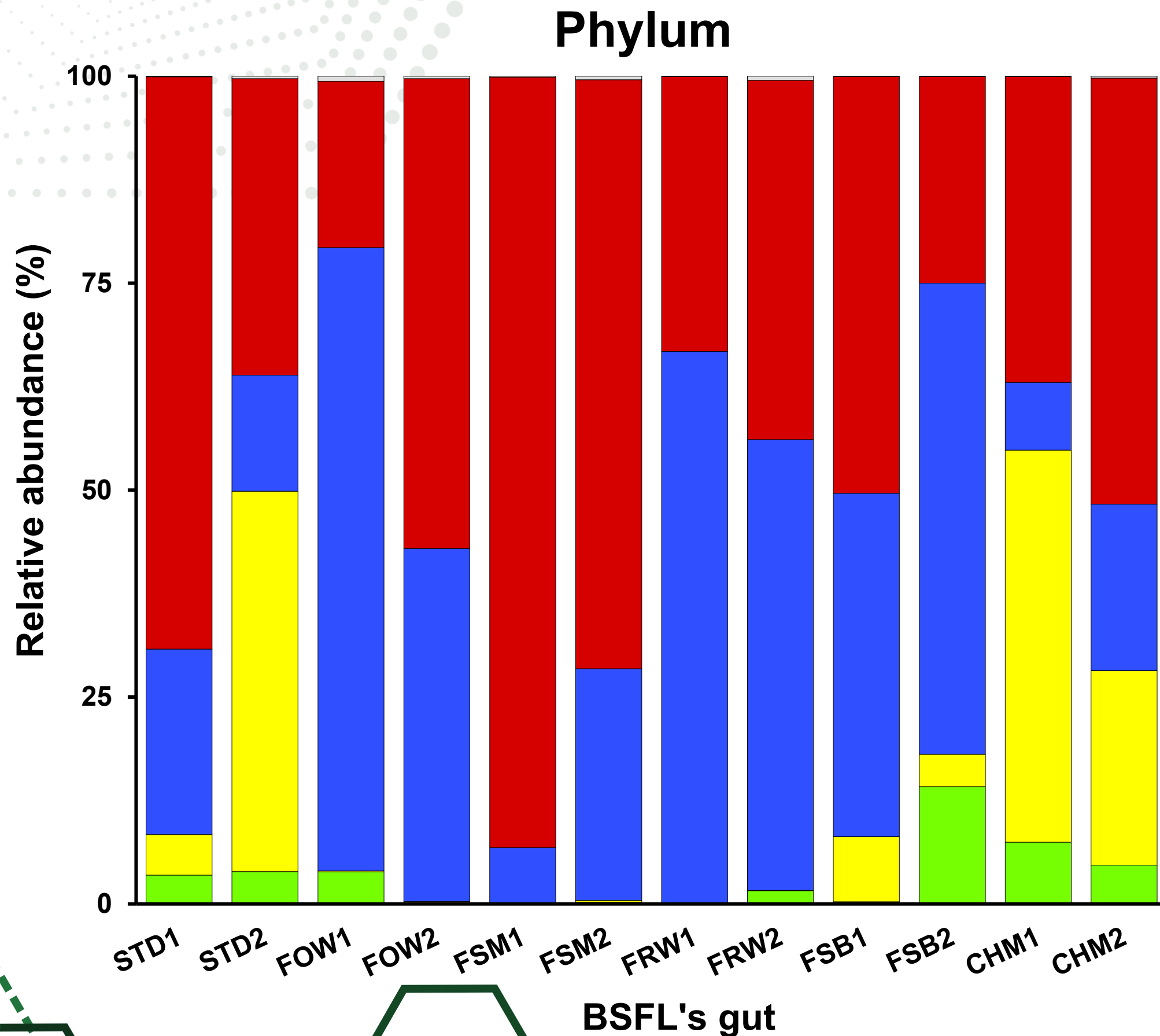


“Bacterial non-culture method” - OTUs





“Bacterial non-culture method” - Phylum level



All substrates



- Proteobacteria
- Firmicutes

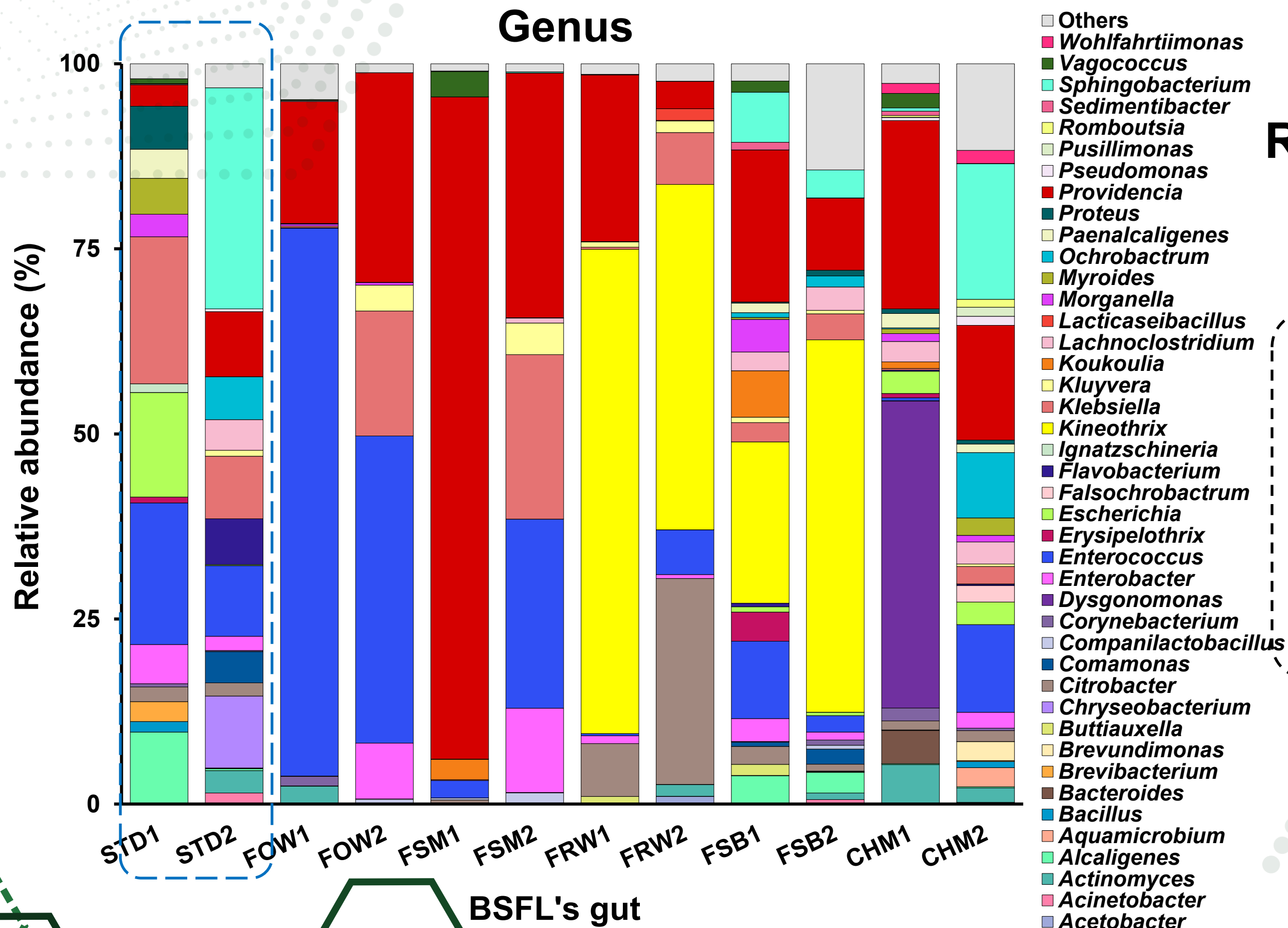
Some rearing substrates

- Bacteroidota
- Actinobacteriota

- Others
- Proteobacteria
- Firmicutes
- Bacteroidota
- Actinobacteriota

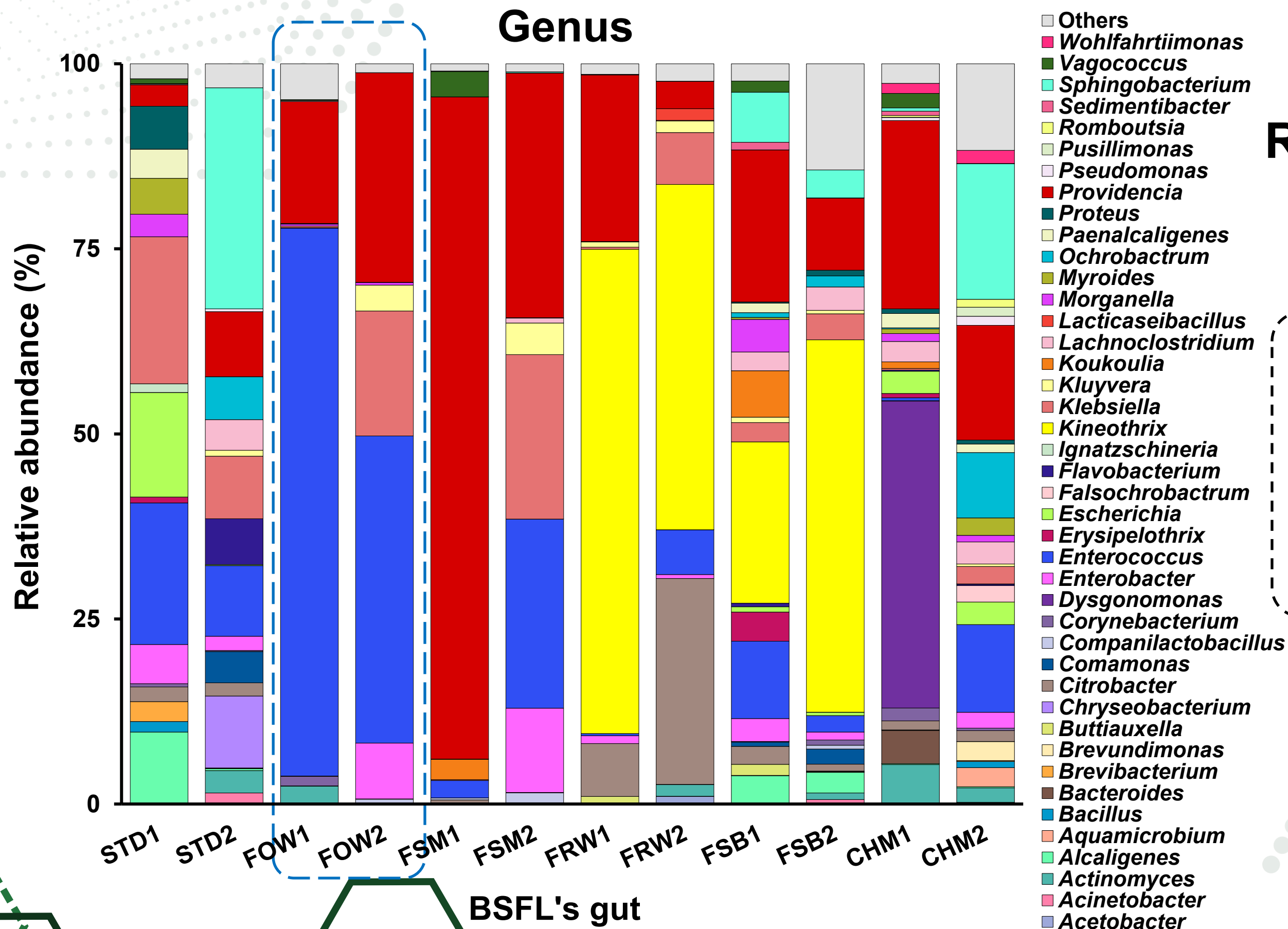


“Bacterial non-culture method” - Genus level





“Bacterial non-culture method” - Genus level



Relative abundance $\geq 1\%$

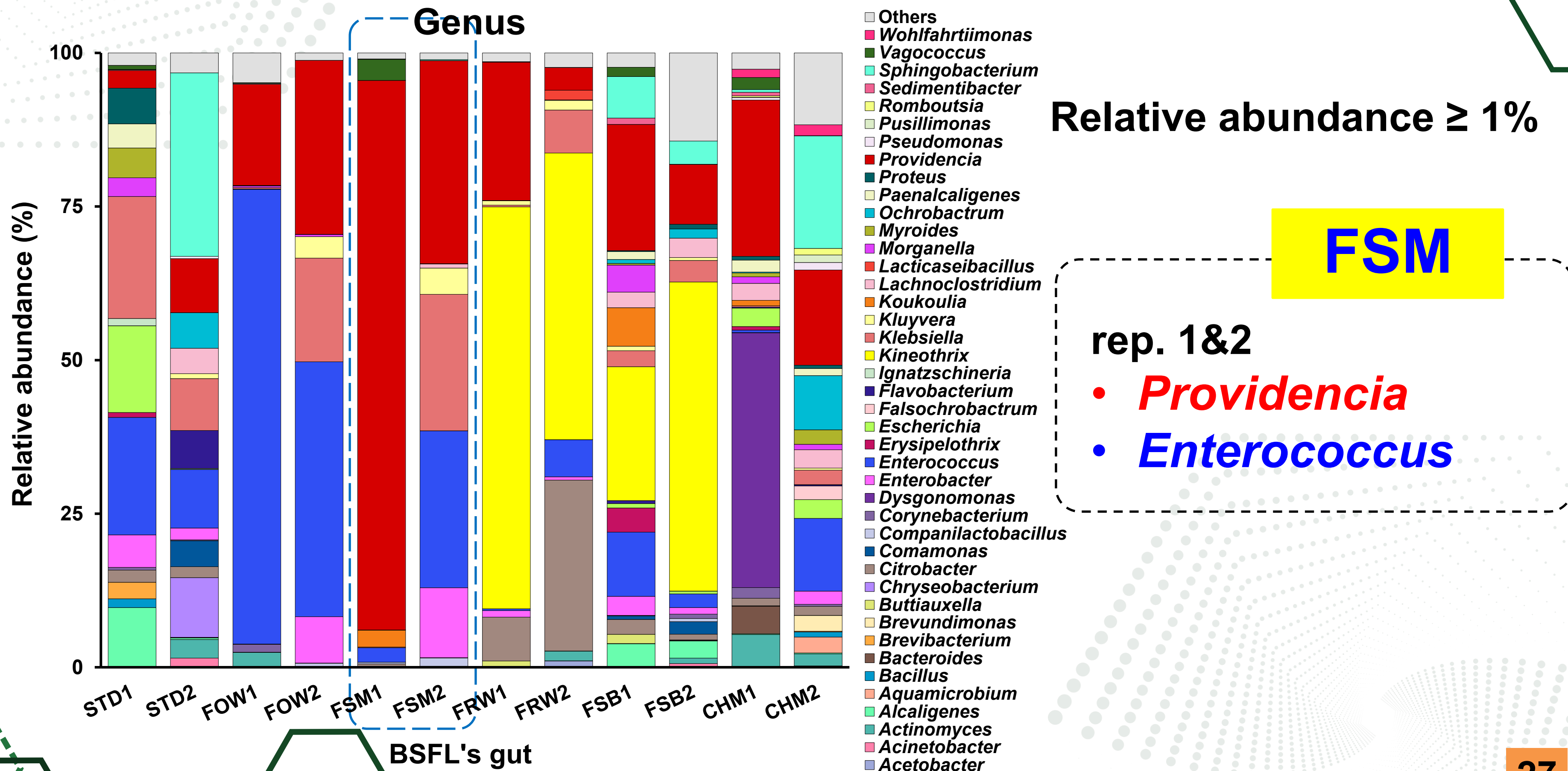
FOW

rep. 1&2

- *Enterococcus*
- *Providencia*

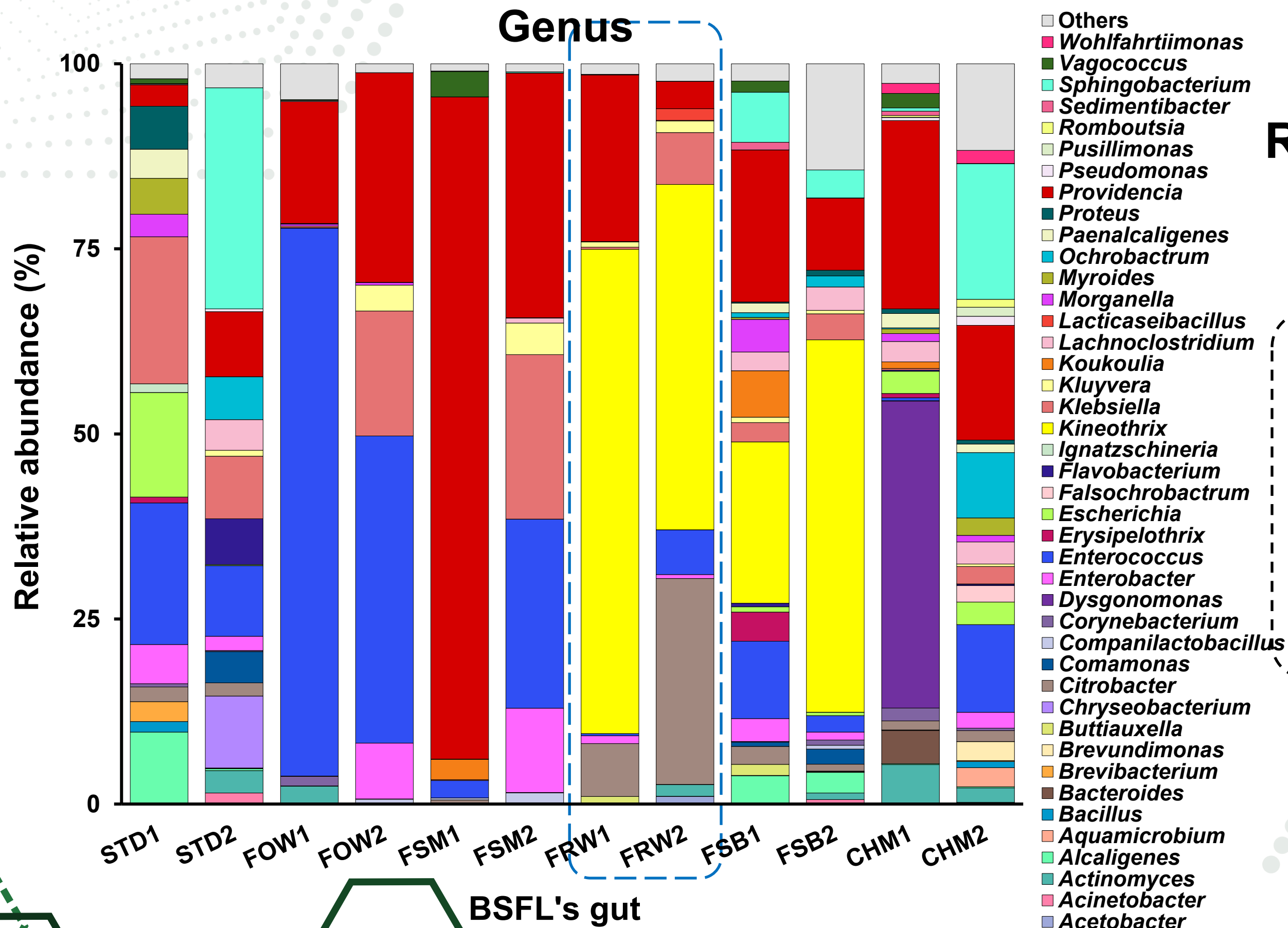


“Bacterial non-culture method” - Genus level





“Bacterial non-culture method” - Genus level



Relative abundance $\geq 1\%$

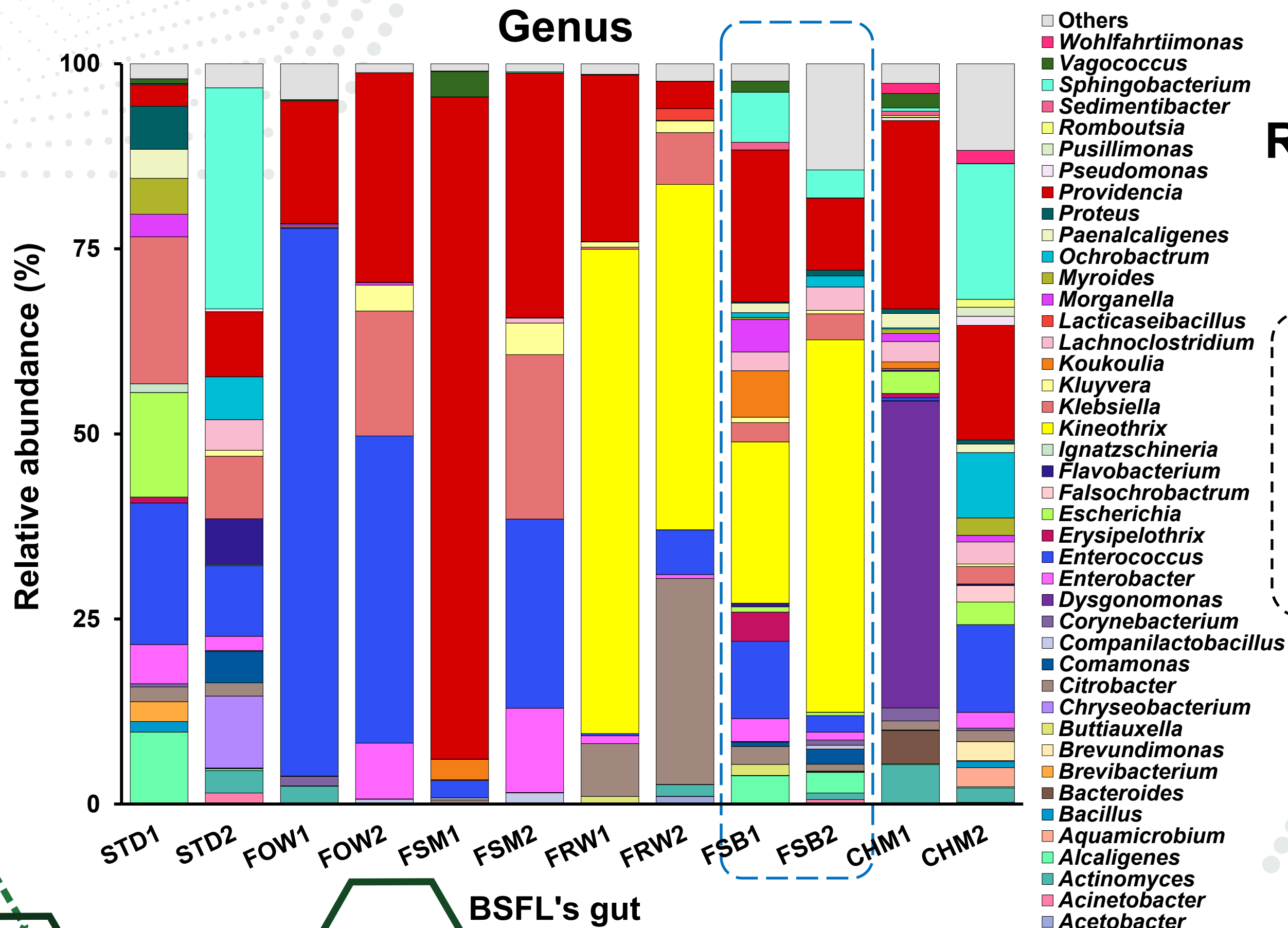
FRW

rep. 1&2

- *Kineothrix*
- *Providencia*
- *Citrobacter*



“Bacterial non-culture method” - Genus level



Relative abundance $\geq 1\%$

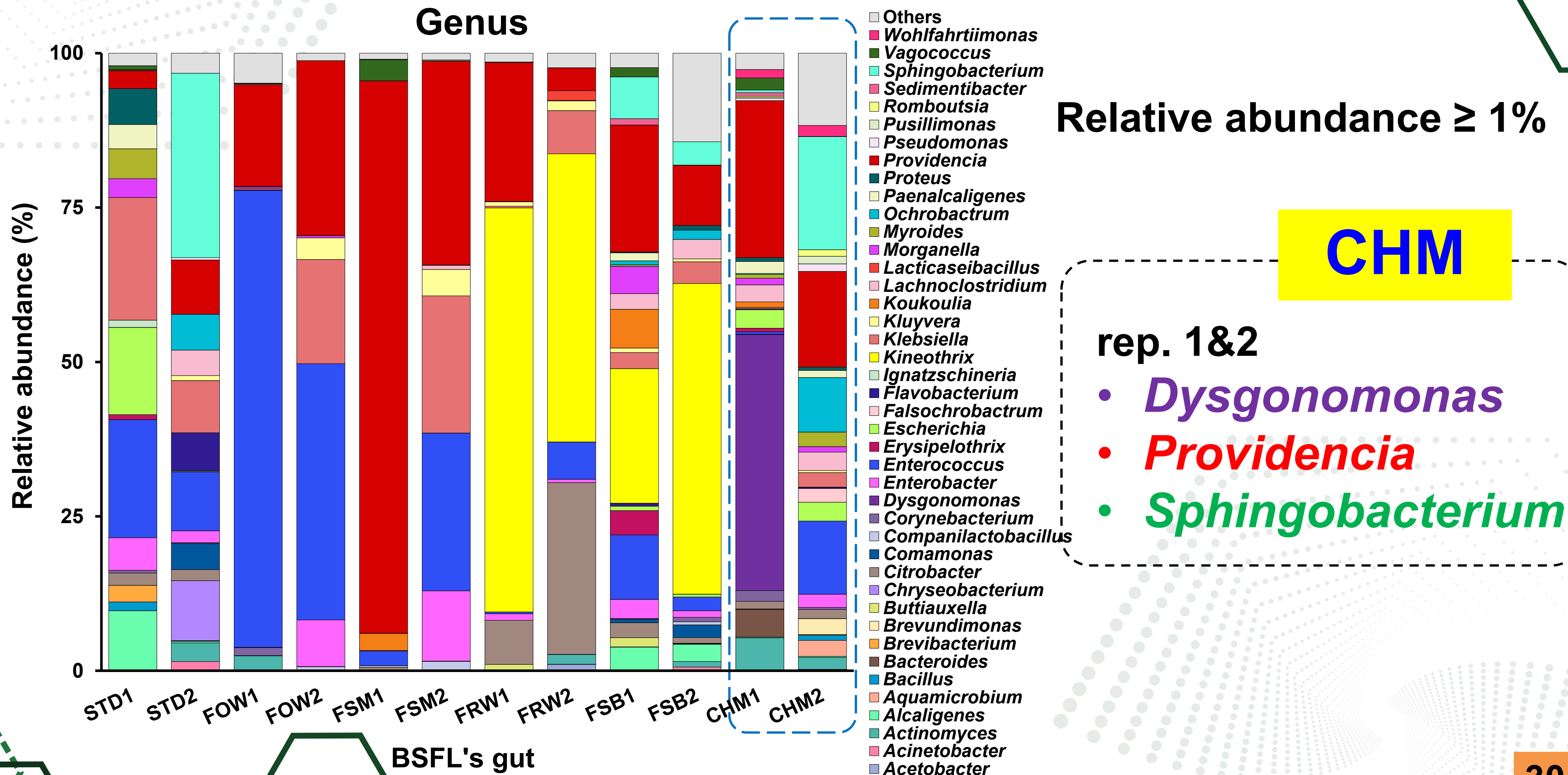
FSB

rep. 1&2

- *Kineothrix*
- *Providencia*



“Bacterial non-culture method” - Genus level



- 4 main phyla: **Actinobacteriota**, **Bacteroidota**, **Firmicutes** & **Proteobacteria** were found in this study.



= Commonly found as **dominant bacteria** in the **larval gut**. (Bruno et al., 2019; Tanga et al., 2021; Schreven et al., 2022)



The **abundance of each bacterial phyla** differ across studies.



Klammsteiner et al. (2020)

- phylum **Actinobacteriota & Bacteroidota**



Gorrens et al. (2021)

- phylum **Proteobacteria & Firmicutes**

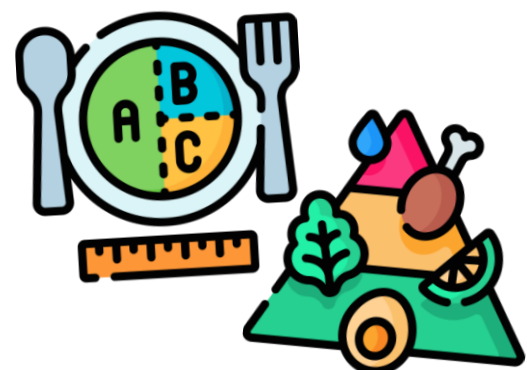


Gut bacteria varies depending on the “**substrates.**”



Substrates microbes

- It can **enter the gut & become part** of the larval gut microbiota. (Tanga et al., 2021)



Nutrient composition

• **Protein, Fat & Carbohydrate.**

- It influences bacterial diversity & **enzyme production** for digestion. (Chen et al., 2023; Bonelli et al., 2020)



Bacteria that degrade **protein- & fat-rich substrates**

- *Providencia*
- *Enterococcus*
- *Morganella*



Bacteria that degrade **carbohydrate- & fiber-rich substrates**

- *Providencia* (Hemicellulose)
- *Klebsiella* (Cellulose, Hemicellulose & Pectin)
- *Enterococcus*
- *Dysgonomonas* (Cellulose& Hemicellulose)

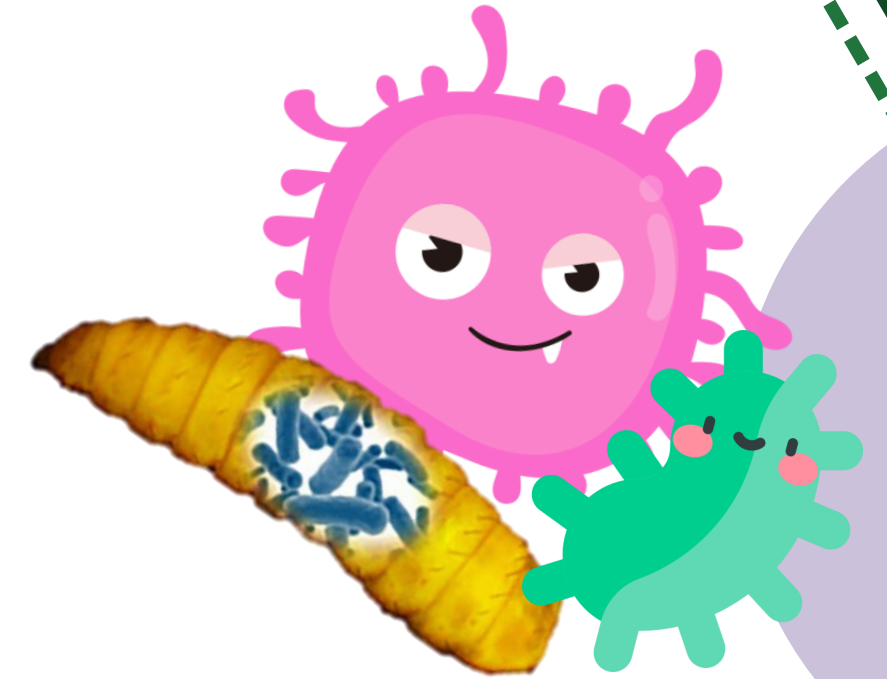
= “**Core microbiota**”

(Tanga et al., 2021; Tegtmeier et al., 2021)

“Biosafety perspective”



Bacteria + Positive association with BSFL.



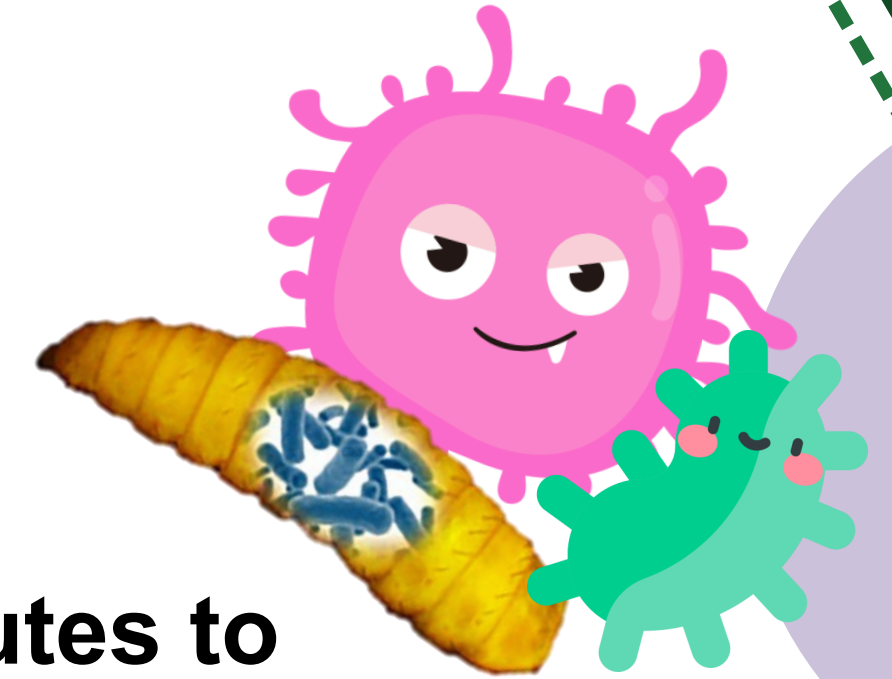
> genus *Providencia & Morganella*

+ It can stimulate oviposition & be vertically transmitted in BSF. (Shumo et al., 2019)

> genus *Enterococcus*

+ • Increasing the BSF egg-hatching rate & larval growth rate. (Shumo et al., 2019)

“Biosafety perspective”



> genus *Klebsiella*

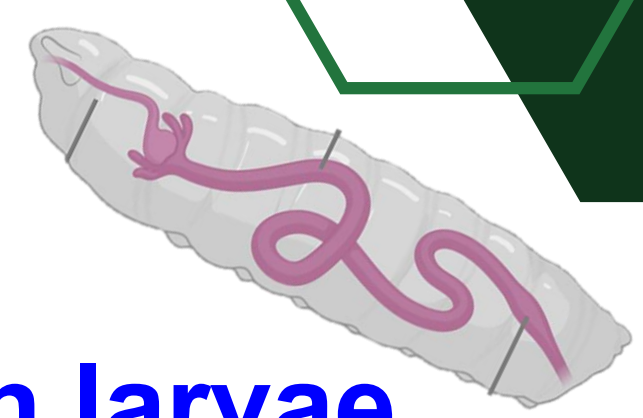
+ It supports **early larval growth** & contributes to **pre-pupal weight gain.** (Li et al., 2022b)

> genus *Dysgonomonas*

+ It promotes **faster larval growth** & enhances **pupation efficiency.** (Li et al., 2022b)



“CONCLUSION”



All substrates induce shifts gut bacteria in larvae.



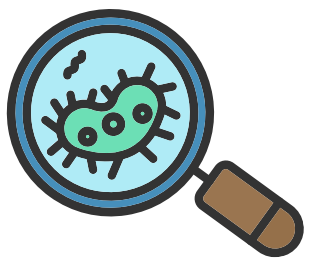
The **culture** method

- genus *Bacillus* in all BSFL guts.
- **FRW** had the **highest** diversity.



The **non-culture** method

- genus *Enterococcus*, *Klebsiella* & *Providencia* in all BSFL guts.
- **CHM** had the **highest** diversity.



Some species may be “pathogenic to humans.”



“RECOMMENDATION”



“Ensuring Safety in **BSFL Utilization**”

✓ Implement proper **feed management**.

- Hygienic feed storage
- Immediate use after preparation

✓ **Postharvest** treatments

- Heating
- Drying
- Blanching
- Defatting
- Storage in hermetic bag



Reduce microbial contaminants.



“RECOMMENDATION”

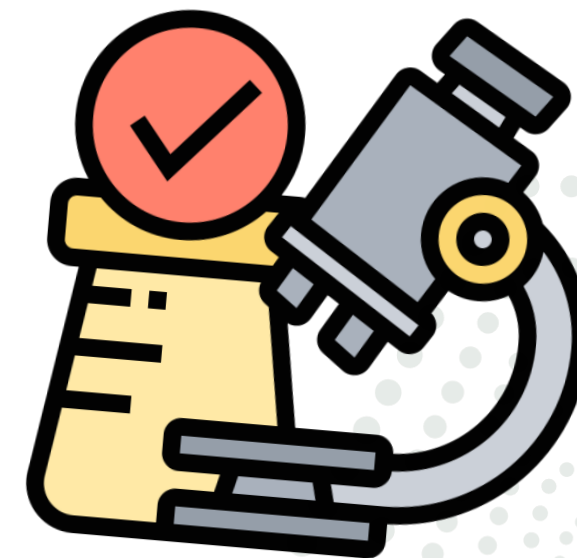


“Research further”



Detection of the **main foodborne** pathogen strains in **BSFL’ s gut** and **substrates**.

- *Campylobacter jejuni*
- *Clostridium botulinum*
- *Escherichia coli*
- *Listeria monocytogenes*
- *Salmonella spp.*
- *Staphylococcus aureus*



THANK
YOU

