

# Microbial communities of homemade fermented vegetables

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#### ▶ To cite this version:

Anne Thierry, Marie-Noëlle Madec, Anne-Sophie Bage, Victoria Chuat, Laurent Marché, et al.. Microbial communities of homemade fermented vegetables. 16ème colloque National SFM, Sep 2021, Nantes, France. , 2022. hal-03789330

HAL Id: hal-03789330

https://hal.inrae.fr/hal-03789330

Submitted on 27 Sep 2022

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Fraternité



Microbial communities of home-made fermented vegetables

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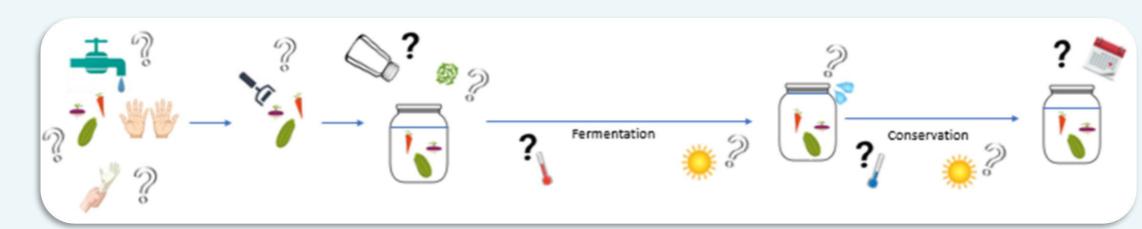


#### **CONTEXT & AIM**

- ✓ Fermentation : an ancestral process of preserving food that has recently received a renewed interest because of its natural image, nutritional interest, and potential for innovation.
- ✓ Excepted sauerkraut, fermented foods issued from vegetables have been little consumed in Western Europe, in contrast with central Europe and Asia.
- ✓ FLEGME "Fermentation des LEGuMEs": a citizen science project that gathers citizens, SMEs, agricultural schools, culinary journalists, researchers, ....
- ✓ One of its goal is to characterize the microbial communities and the safety of home-made fermented vegetables and isolate strains.

### **STRATEGY**

- ✓ Constitution of a community of > 250 citizens that manufacture fermented legumes for their personal consumption
- Home-made fermented vegetables collected from Feb, to Oct 2020, as well as the manufacturing conditions, practices, and recipes via an online survey

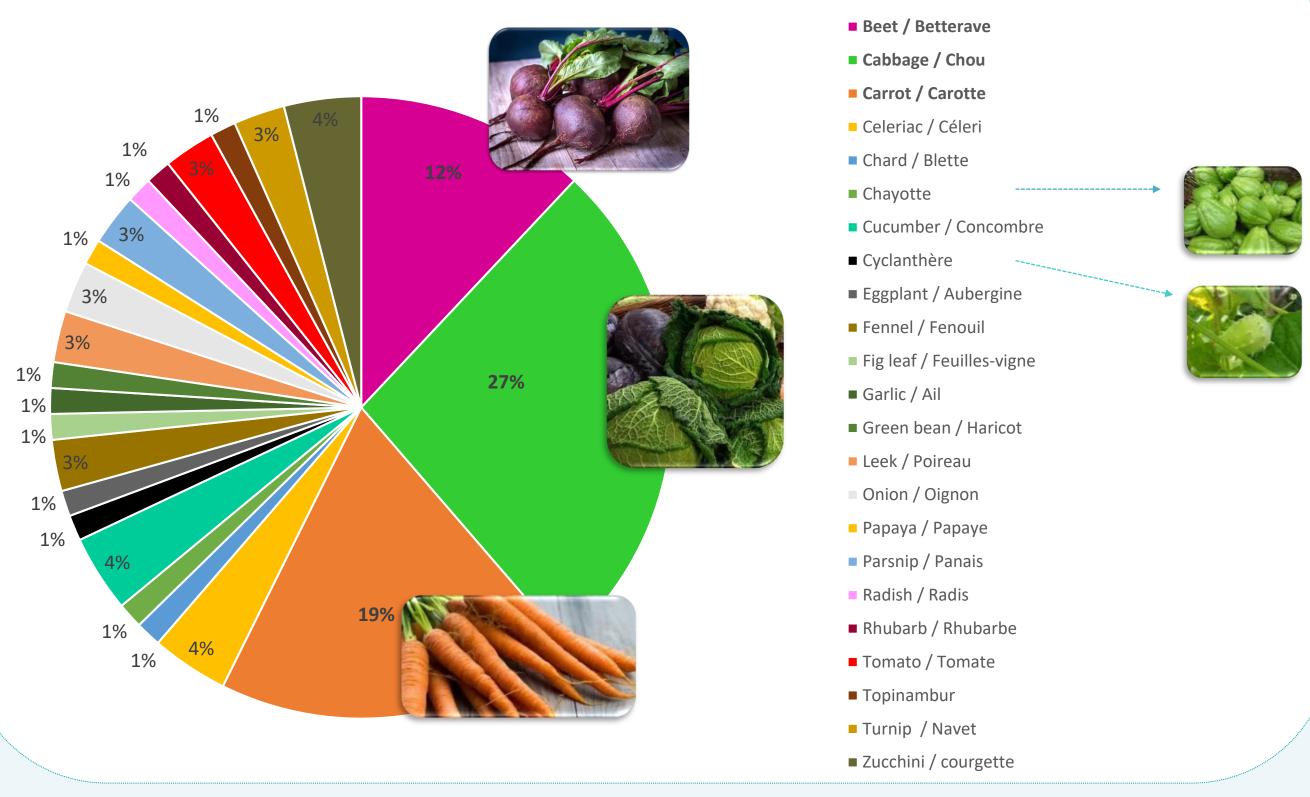


Desirable/undesirable microbial groups assessed by a culture-dependent approach:

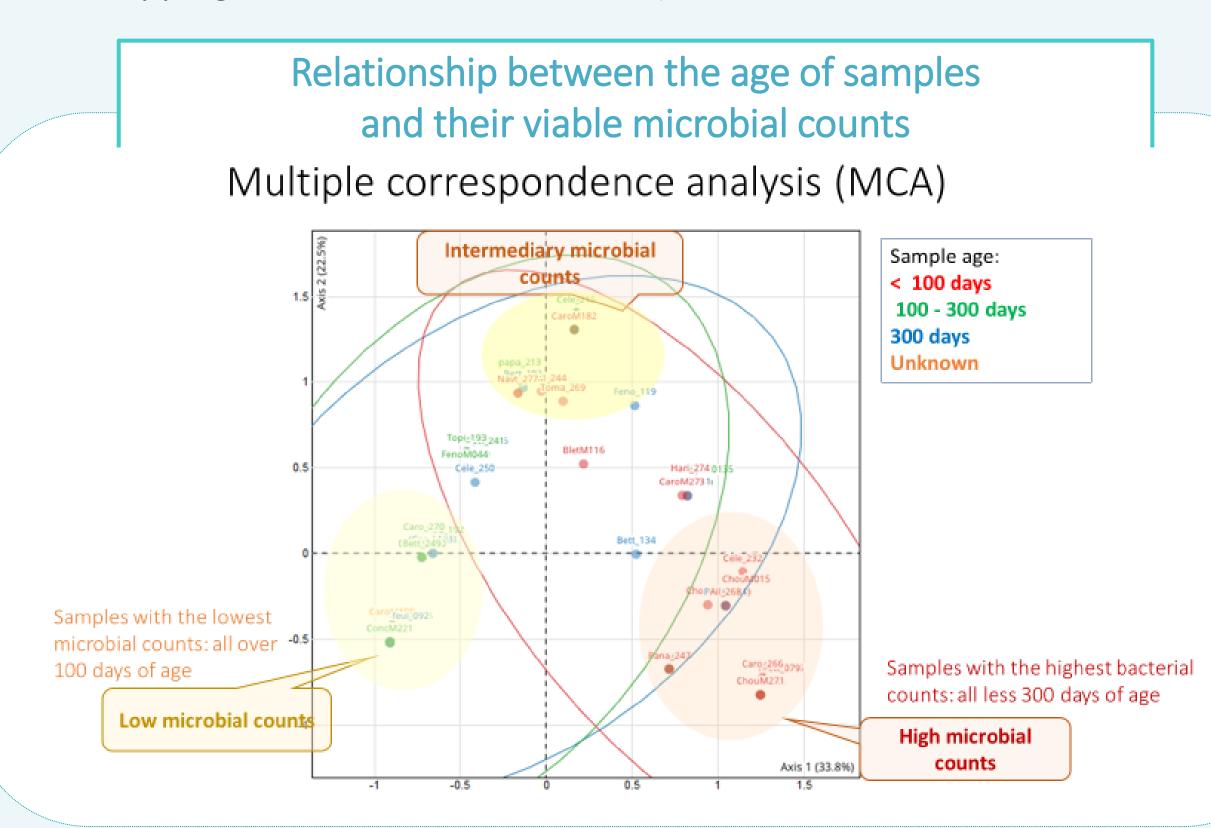
lactic acid bacteria (LAB)

- total aerobic bacteria,
- Enterococcus, Enterobacteriaceae,
  - spore-forming bacteria,
  - pathogenic bacteria

A large diversity of fermented vegetables collected from citizens (n=75) and various manufacturing practices and recipes

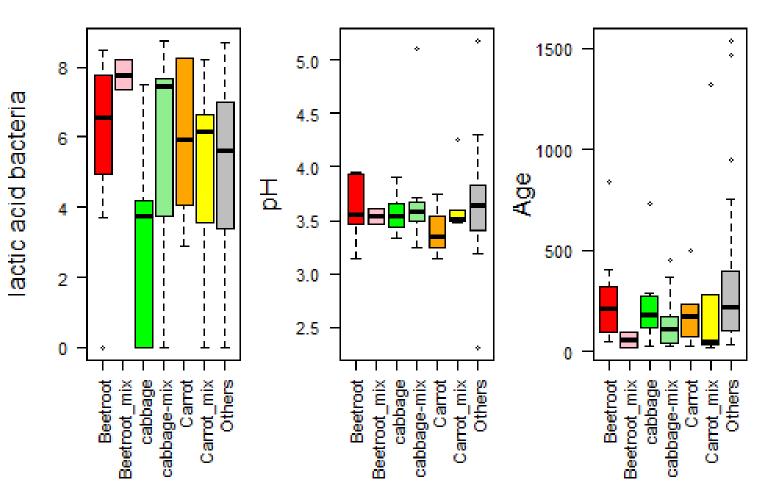


- $\checkmark$  23 types of legumes, the most represented being cabbage (27%, 7 varieties), carrots (19%) and beets (12%), most of them coming from from organic commercial production or garden vegetables
- ✓ Various manufacturing practices: vegetables peeled or not, various washing practices and slicing preparations (in pieces, sliced, minced, grated)
- ✓ Recipes include in general 1 to 2 spices and/or seasonings (garlic, coriander seeds, pepper, thyme, cumin, gingers, bay leaf being the most used)
- ✓ Spontaneous fermentation for 71 out of 75 samples (for the 4 others, 3 using backslopping and 1 kefir for inoculation)



## **RESULTS**

# Several microbial groups, low pH and no pathogenic bacteria



LAB counts: median: 7.5x10<sup>4</sup> CFU/g

• from non-detectable to 6x10<sup>8</sup> CFU/g

pH: median pH: 3.6

- 87% of samples between 3.2 and 4.0
- pH>4.5 for only 2 samples

#### **Storage period (sample age):**

- median 6 months
- from 2 weeks to 4 years

Lactococcus lactis

Lactobacillus parabuchnerii

Lactiplantibacillus plantarum/paraplantarun

✓ LAB represented the dominant population

Lentilactobacillus parafarraginis.

- ✓ No pathogenic bacteria detected (*Escherichia coli, Clostridium perfringens*, coagulase positive staphylococci, Salmonella, Listeria monocytogenes)
- ✓ Enterococci detected in only 5 samples, at counts < 10<sup>5</sup> CFU/g
- $\checkmark$  Yeasts detected in almost half of the samples, at counts ranging from  $10^2$  to  $9x10^7$ CFU/g

90 isolates of lactic acid bacteria – 30 species

### Schleiferilactobacillus harbinensis Bacillus velezensis/amyloliquefaciens/.. Weissella viridescens Pediococcus ethanolidurans Pediococcus pentosaceus Enterobacter hormaechei/ Cloacae Liquorilactobacillus satsumensis . Levilactobacillus yonginensis/koreensis Lactiplantibacillus diolivorans Lactiplantibacillus plantarun Levilactobacillus parabrevis Levilactobacillus brevis Lactiplantibacill... rapi/parabuchneri

- $\checkmark$  1 − 3 isolates randomly picked up on culture media dedicated to LAB numeration conducting to the collection of 90 LAB isolates
  - ✓ Clones were identified by 16S sequencing: 31 taxons identified

Lentilactobacillus kefiri

Lentilactobacillus diolivorans

✓ Levilactobacillus brevis and Lactiplantibacillus plantarum/paraplantarum group were the most common, with 21 and 20 % of total number of isolates

## CONCLUSION

- ✓ No significant relationship between the pH, LAB counts and the type of vegetables was observed.
- ✓ However, a significant effect of the age of samples was observed: high LAB counts were most frequent observed in the youngest samples (age < 100 days).
- ✓ This first view of the microbial community of fermented vegetables based on culture-dependent analysis will be completed by 16S rRNA gene metabarcoding
- ✓ Constitution of collection of lactic acid bacteria representative of the LAB diversity of fermented vegetables (90 strains, 30 species) available for research community







