

# DPC 2026 - Freiburg

Distangling contaminants from true intratumoural microbial signals in pancreatic ductal adenocarcinoma – A benchmarking approach

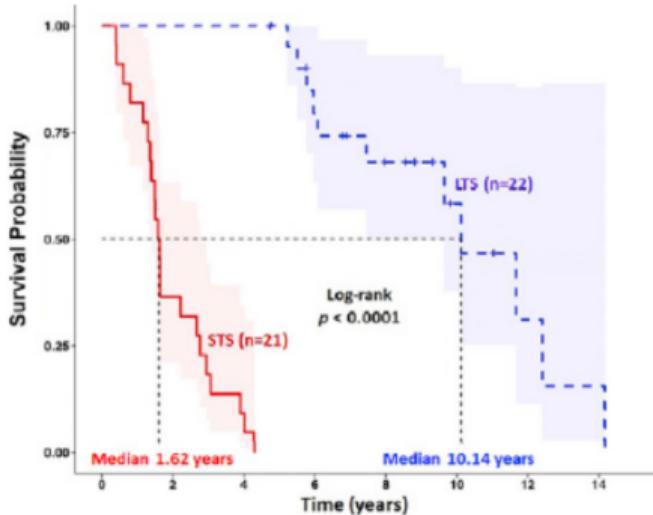
Linh Dang

2026-03-07

# Contamination in Microbiome Study

Cell

## Tumor Microbiome Diversity and Composition Influence Pancreatic Cancer Outcomes



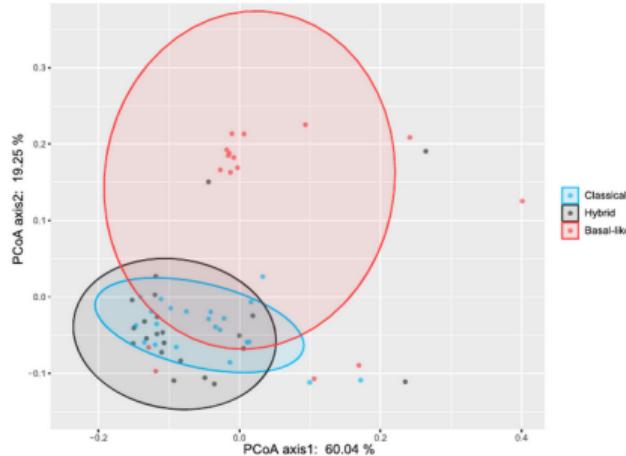
Riquelme et al. Cell 2019

Key taxa: Streptomyces, Pseudoxanthomonas, Saccharopolyspora

Article | Open access | Published: 31 August 2021

## Tumor microbiome contributes to an aggressive phenotype in the basal-like subtype of pancreatic cancer

Wei Guo, Yuchao Zhang, Shiwei Guo, Zi Mei, Huiping Liao, Hang Dong, Kai Wu, Haocheng Ye, Yuhang Zhang, Yufei Zhu, Jingyu Lang, Landian Hu, Gang Jin & Xiangyin Kong



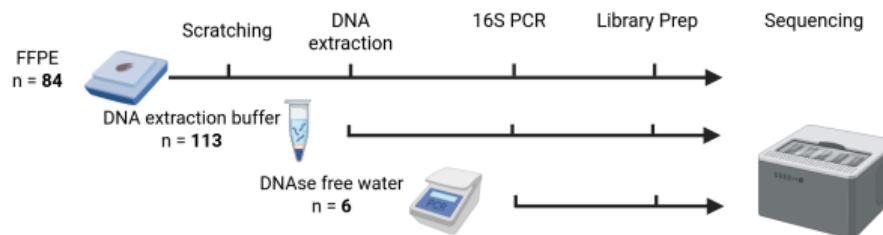
Guo Communication biology 2021

Key taxa: Acinetobacter, Pseudomonas, Sphingopyxis

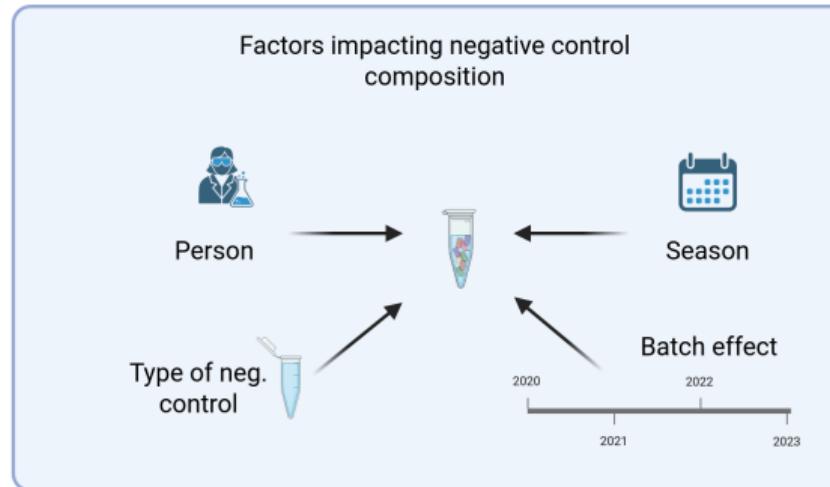
## Section 1

Negative Control Survey in the Lab

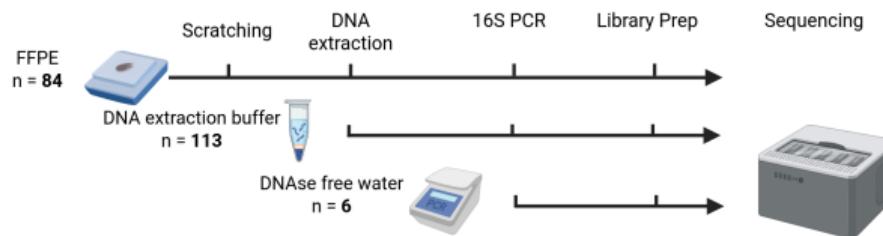
# Negative Sample Collection



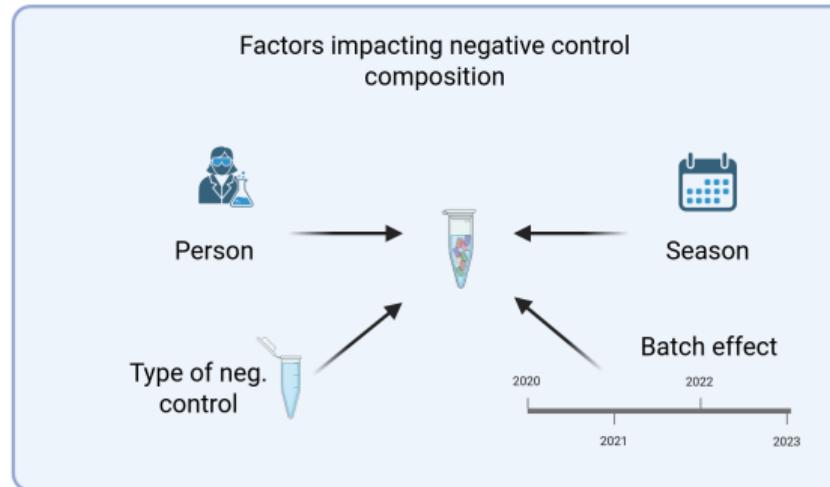
- In total 203 negative control samples over 4 years



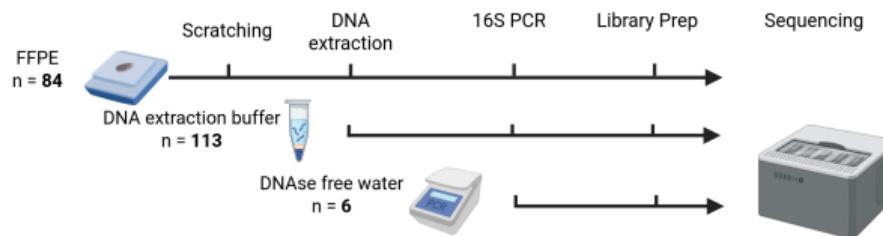
# Negative Sample Collection



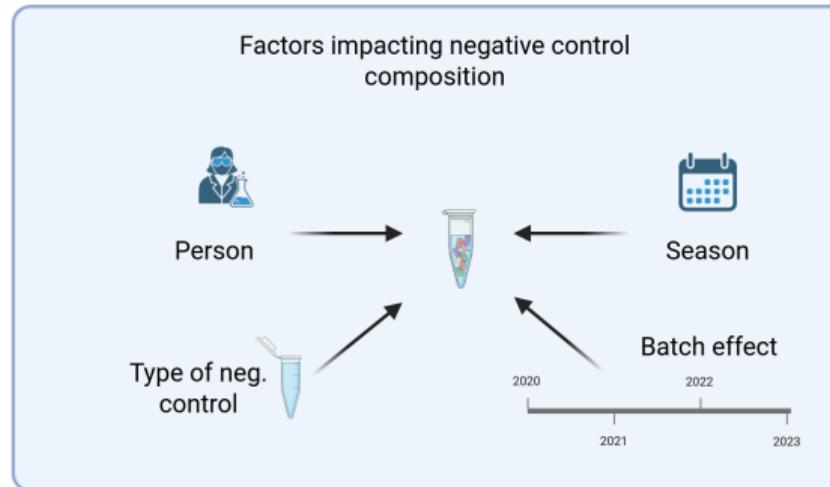
- In total 203 negative control samples over 4 years
  - 113 buffer controls,



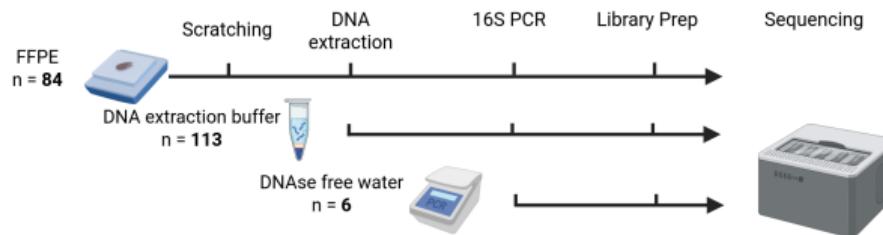
# Negative Sample Collection



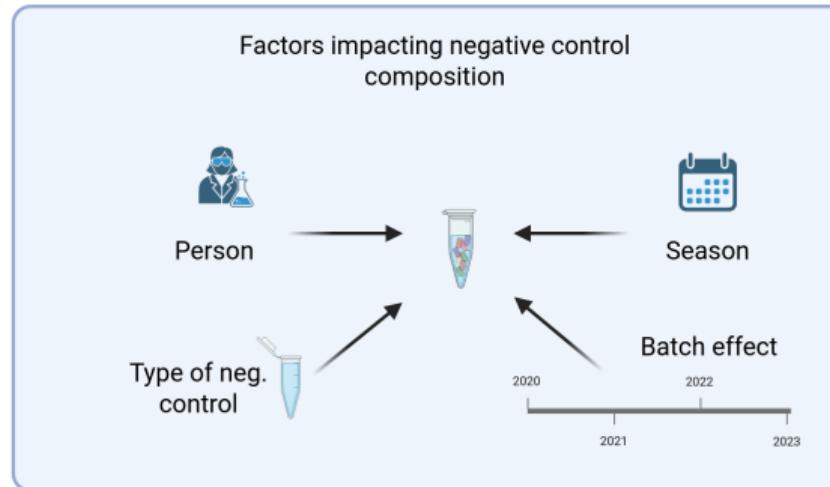
- In total 203 negative control samples over 4 years
  - 113 buffer controls,
  - 84 paraffin controls,



# Negative Sample Collection



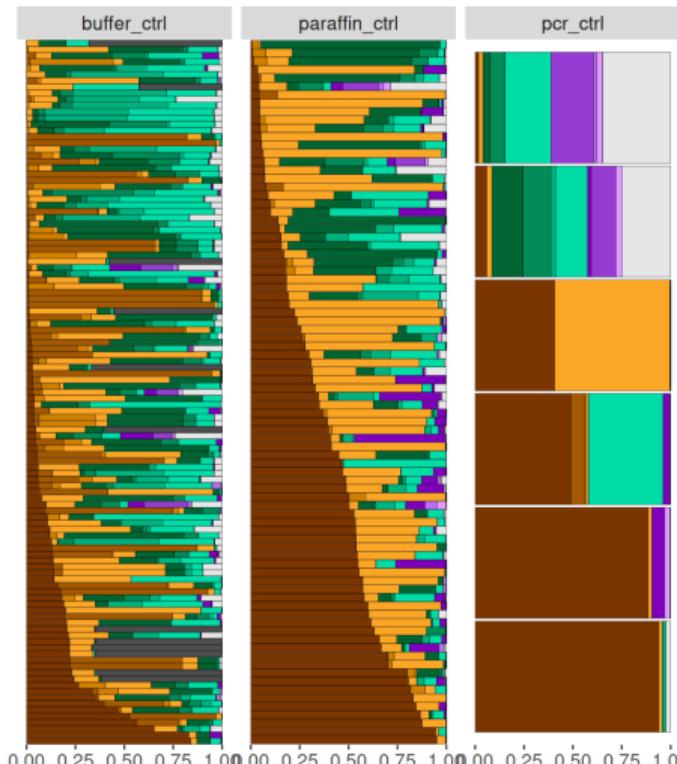
- In total 203 negative control samples over 4 years
  - 113 buffer controls,
  - 84 paraffin controls,
  - 6 PCR control.



# Bacterial Composition

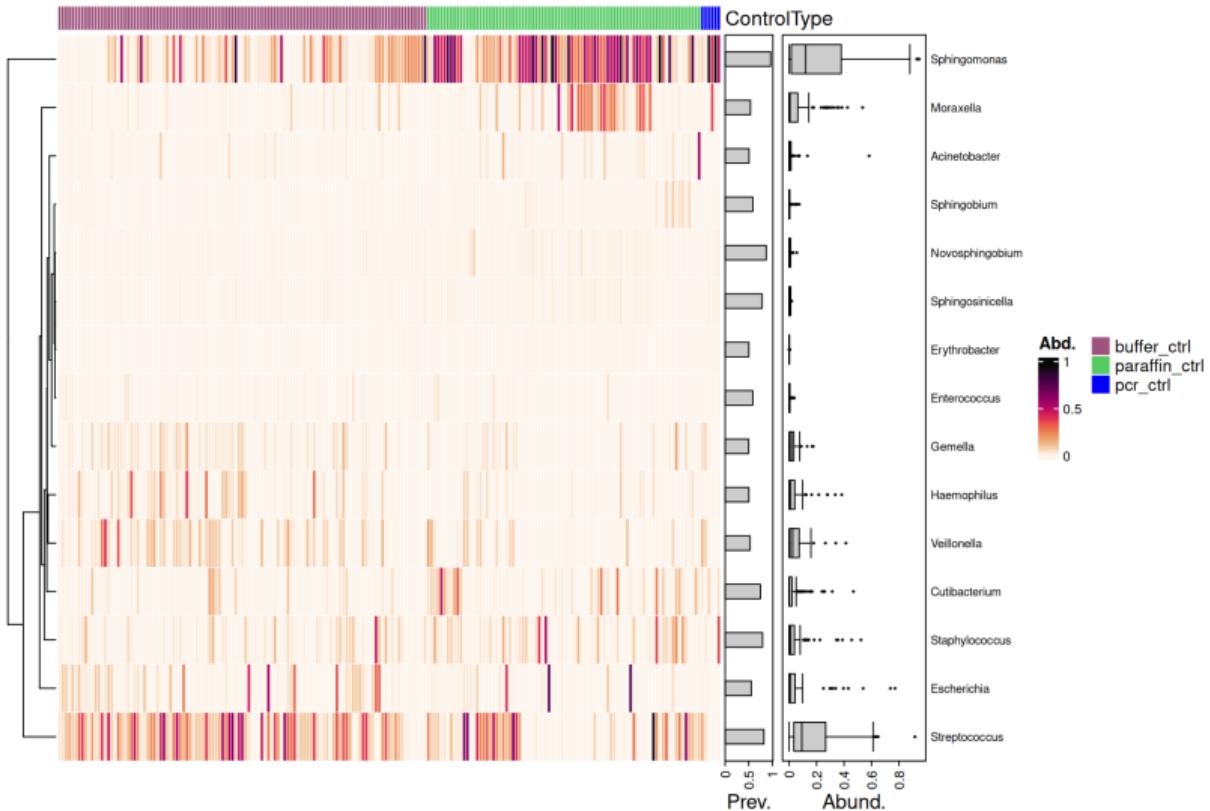
## Composition of NCT Samples

Order by sum



# Heatmap from Negative Controls

Top 15 genera by prev

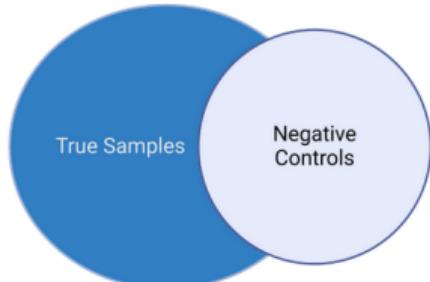


## Section 2

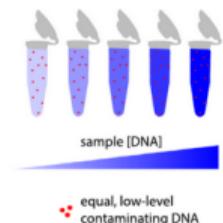
Intratumor Microbiome in PDAC and Assessment of

# Decontamination Strategies

Restrictive

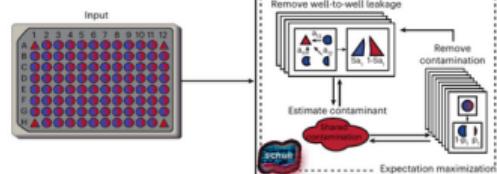


Decontam



contaminant DNA correlates inversely with total DNA

SCRuB

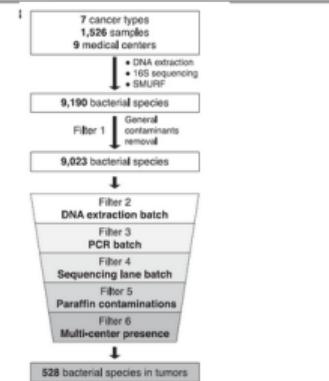


Austin et al. 2023 Nature Biotechnology

Davis et al. 2018 Microbiome

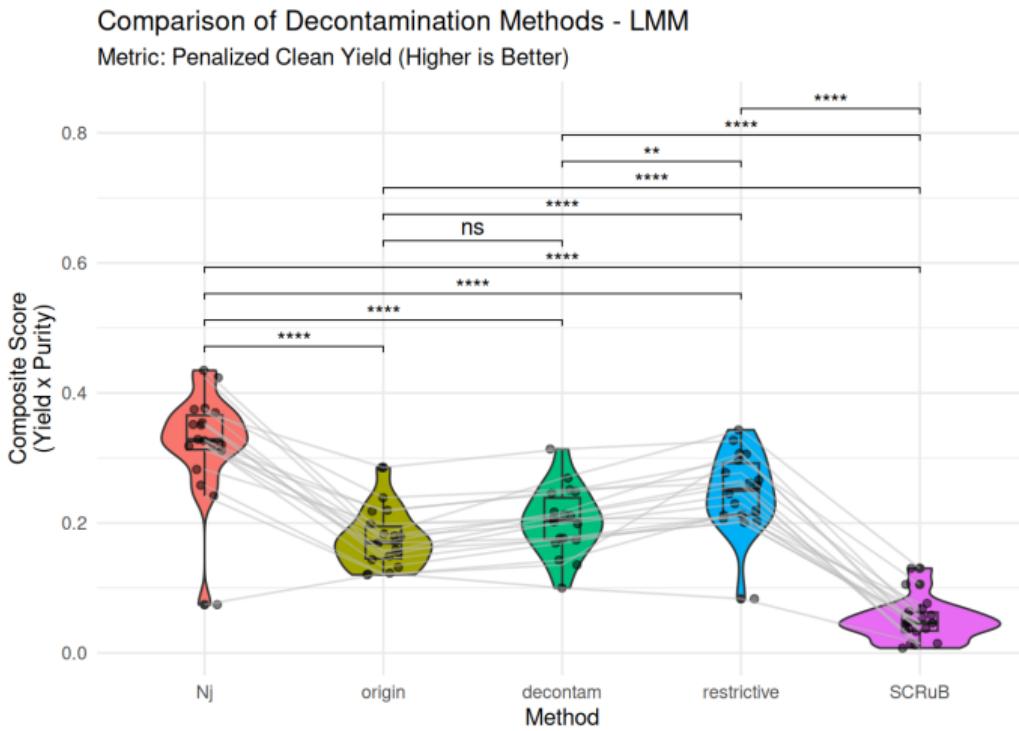


Nejman et al. 2020 Science



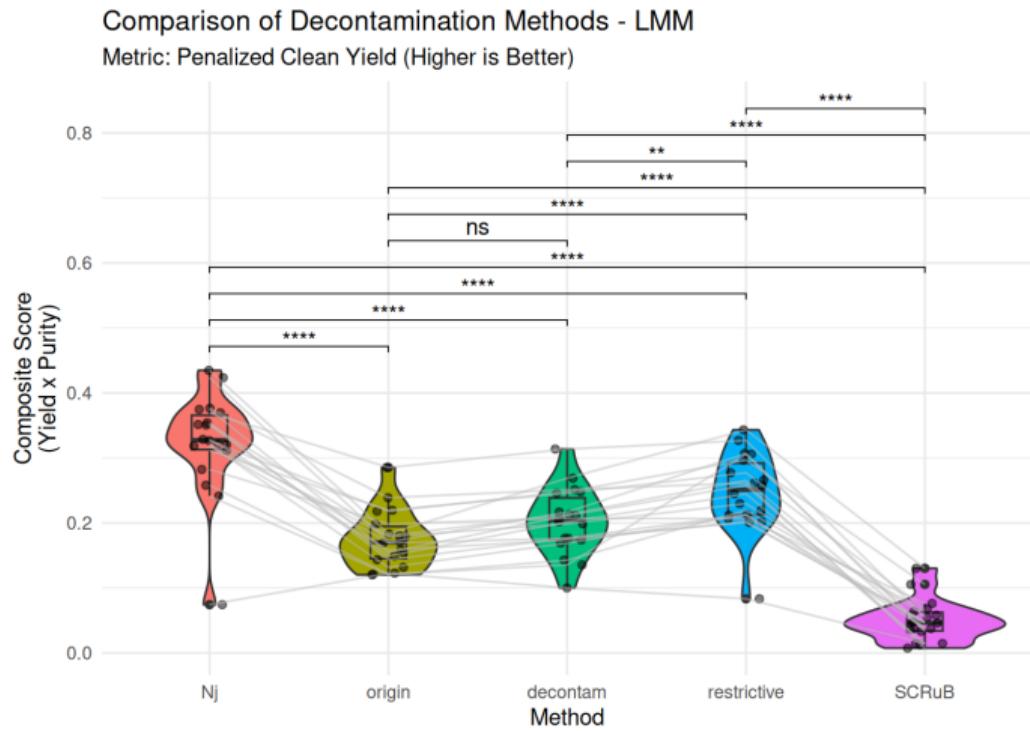
# Assessment of Decontamination Approach

- **Yield:**  
$$\frac{\# \text{ putative true taxa} \notin NCT}{\# \text{observed species}}$$



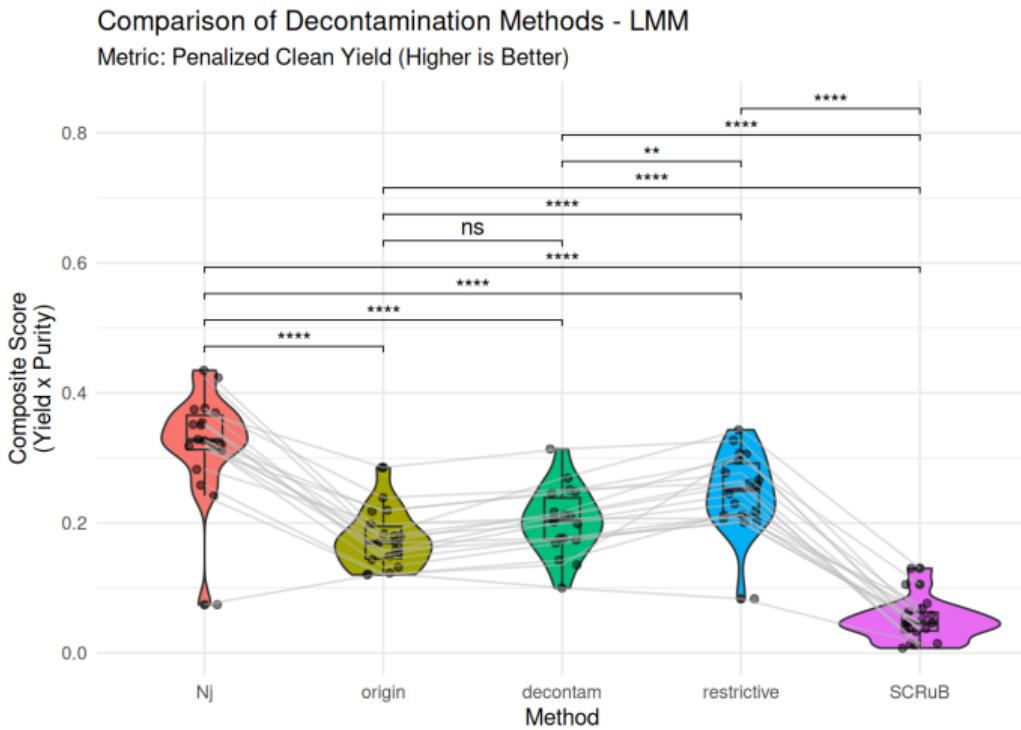
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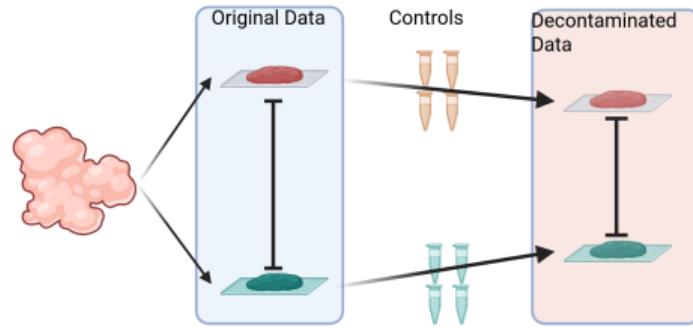
# Assessment of Decontamination Approach

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- **Composite Score =**  
$$\text{Yield} \times \text{Purity}$$



# Integrate Replica

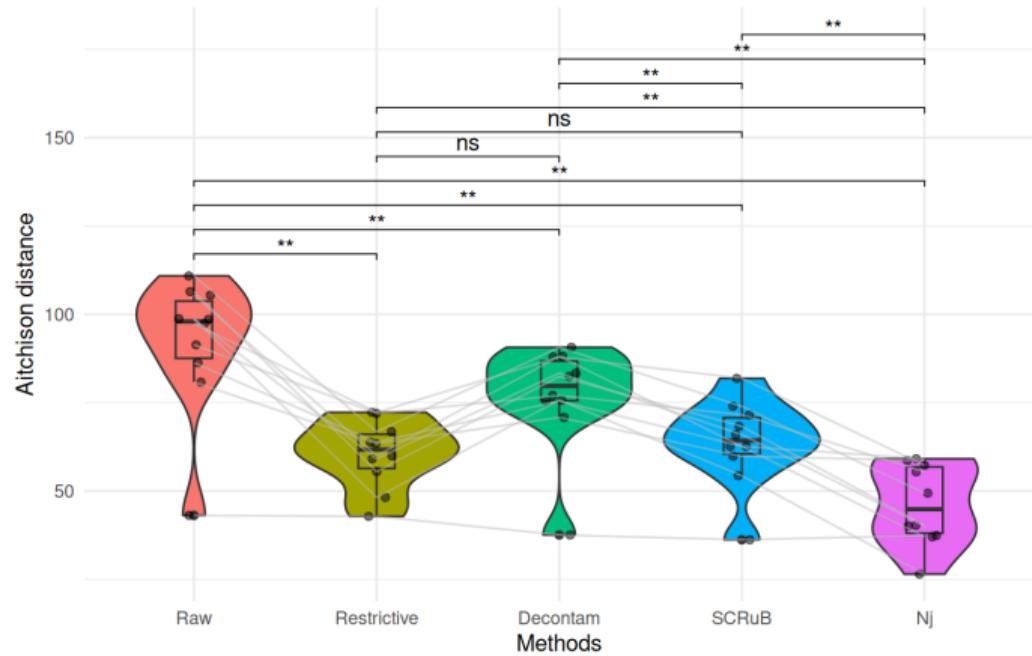
## Utilize Replica for Assessment



- 10 PDAC sample, each with 2 replica

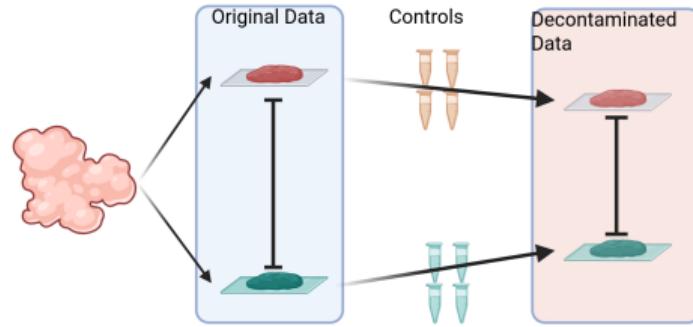
## Paired Wilcox test

The lower the better



# Integrate Replica

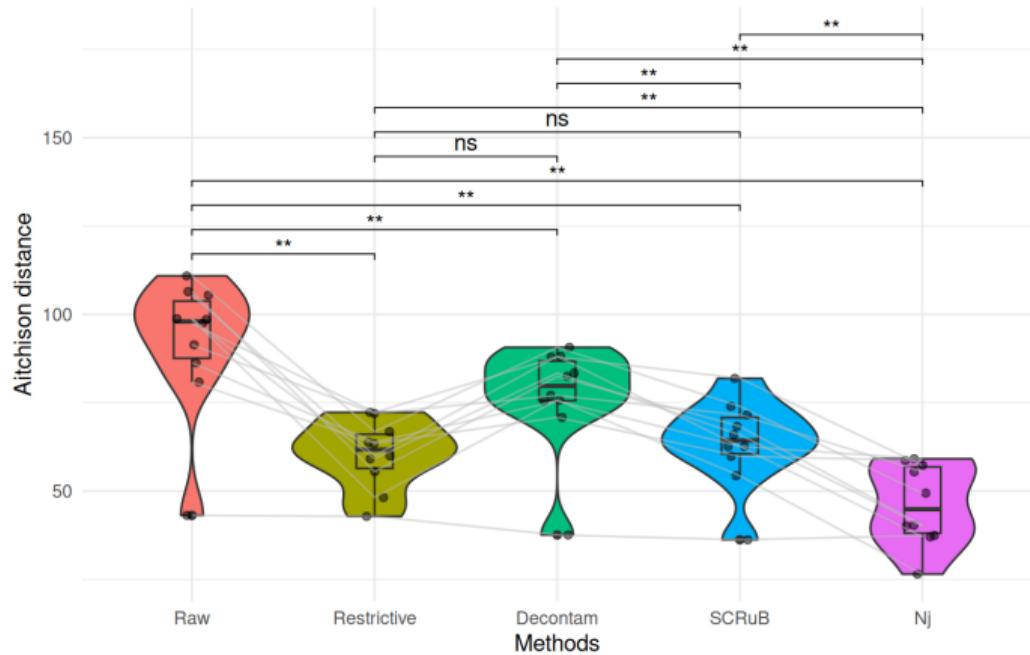
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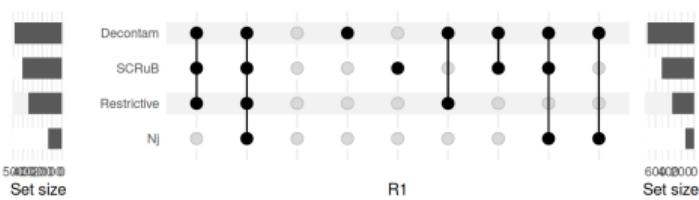
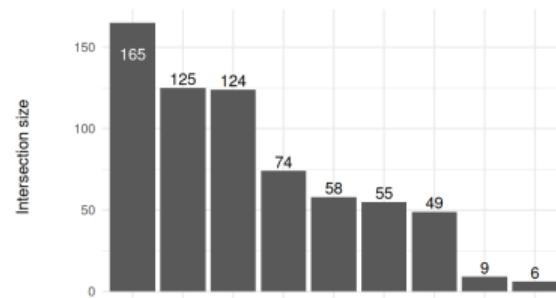
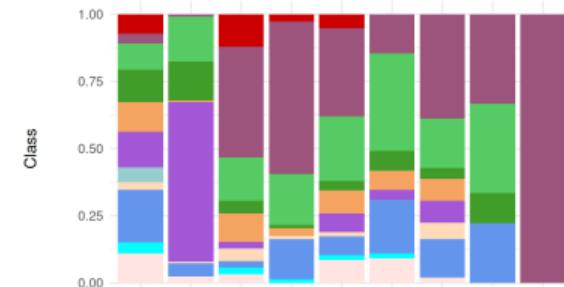
- 10 PDAC sample, each with 2 replica
- Aitchison distance with paired Wilcox test.

## Paired Wilcox test

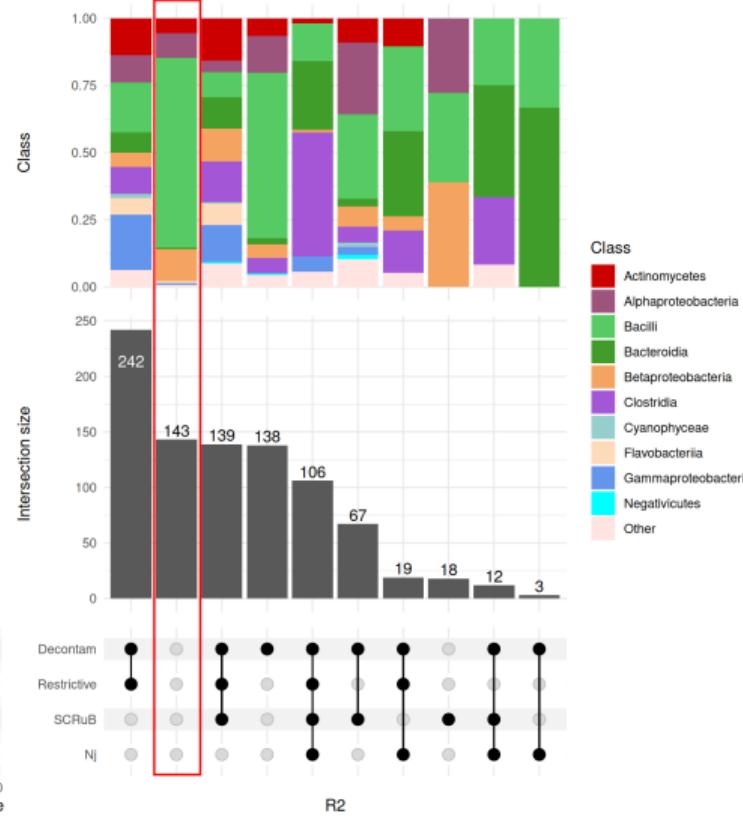
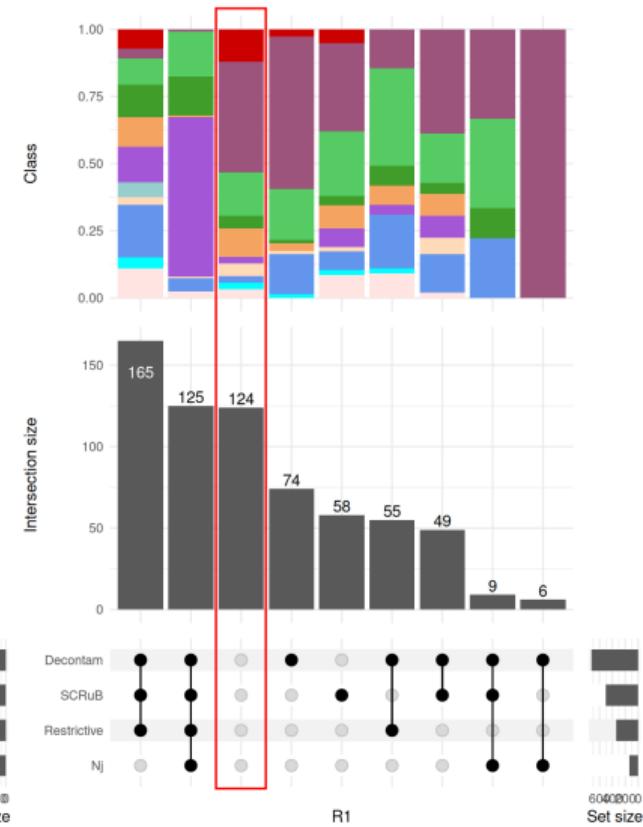
The lower the better



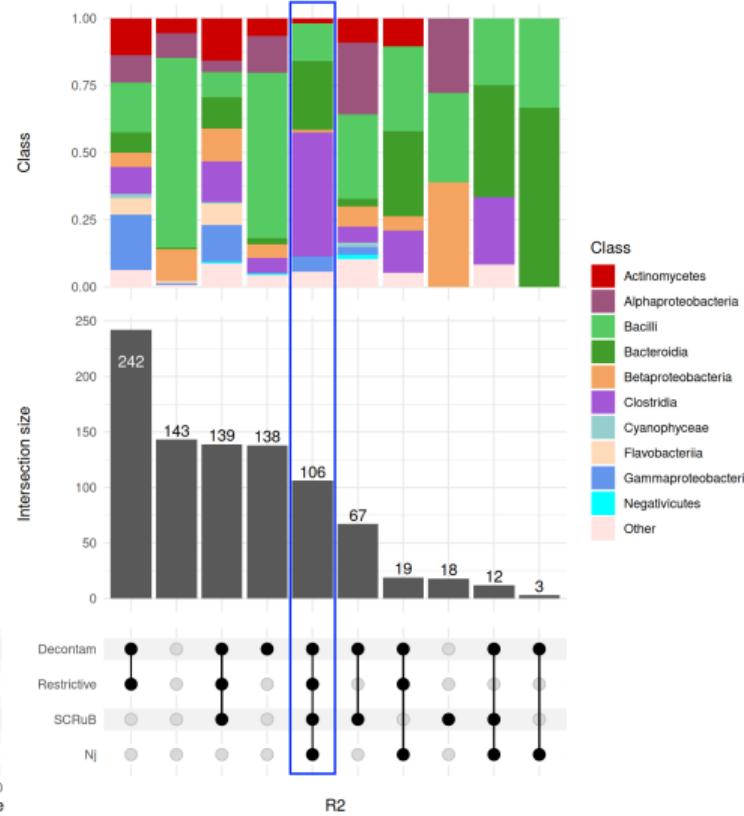
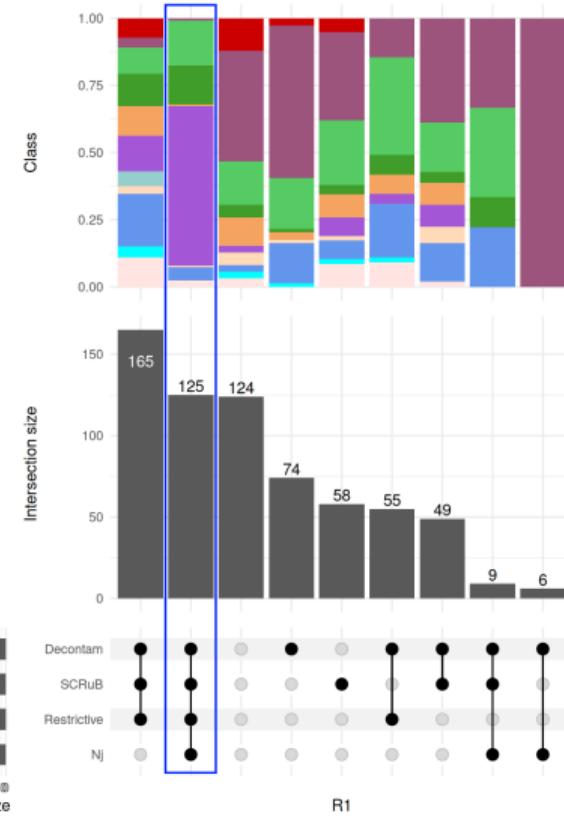
# Overlap - Microbiome profile of PDAC in mice



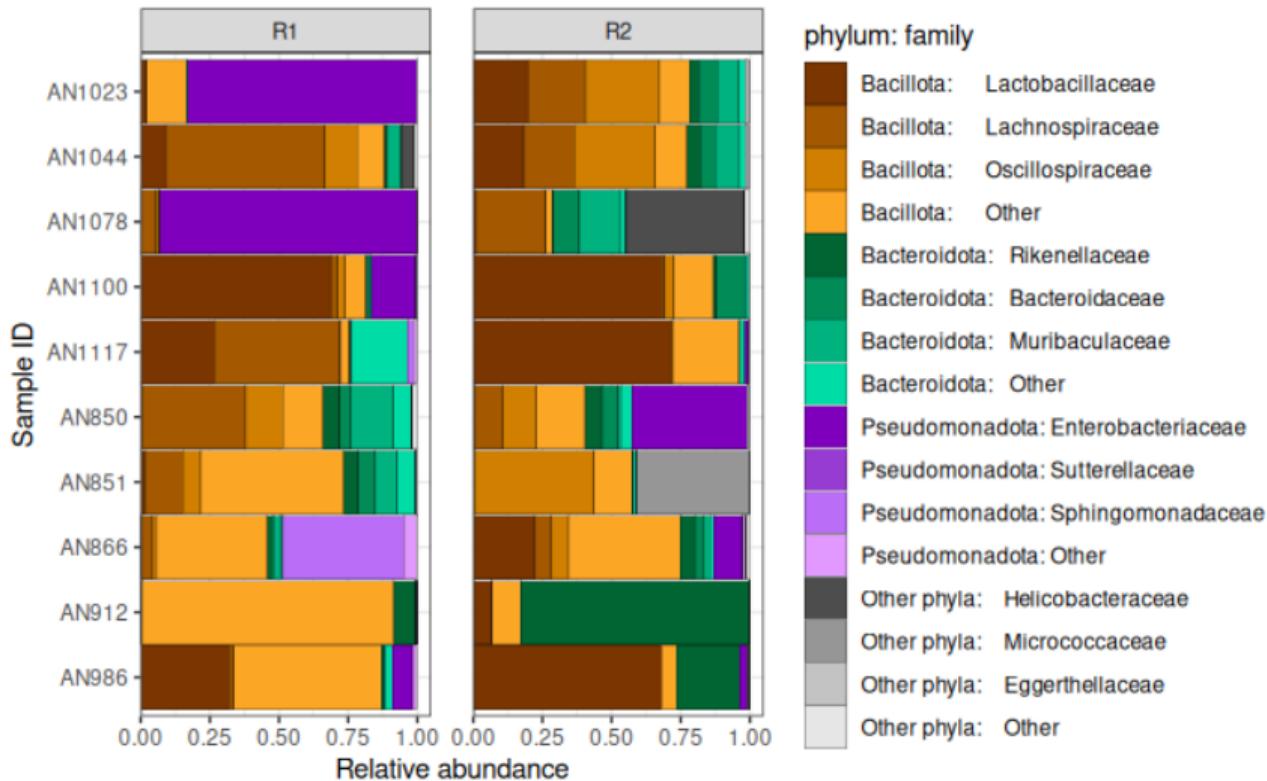
## Overlap - Microbiome profile of PDAC in mice



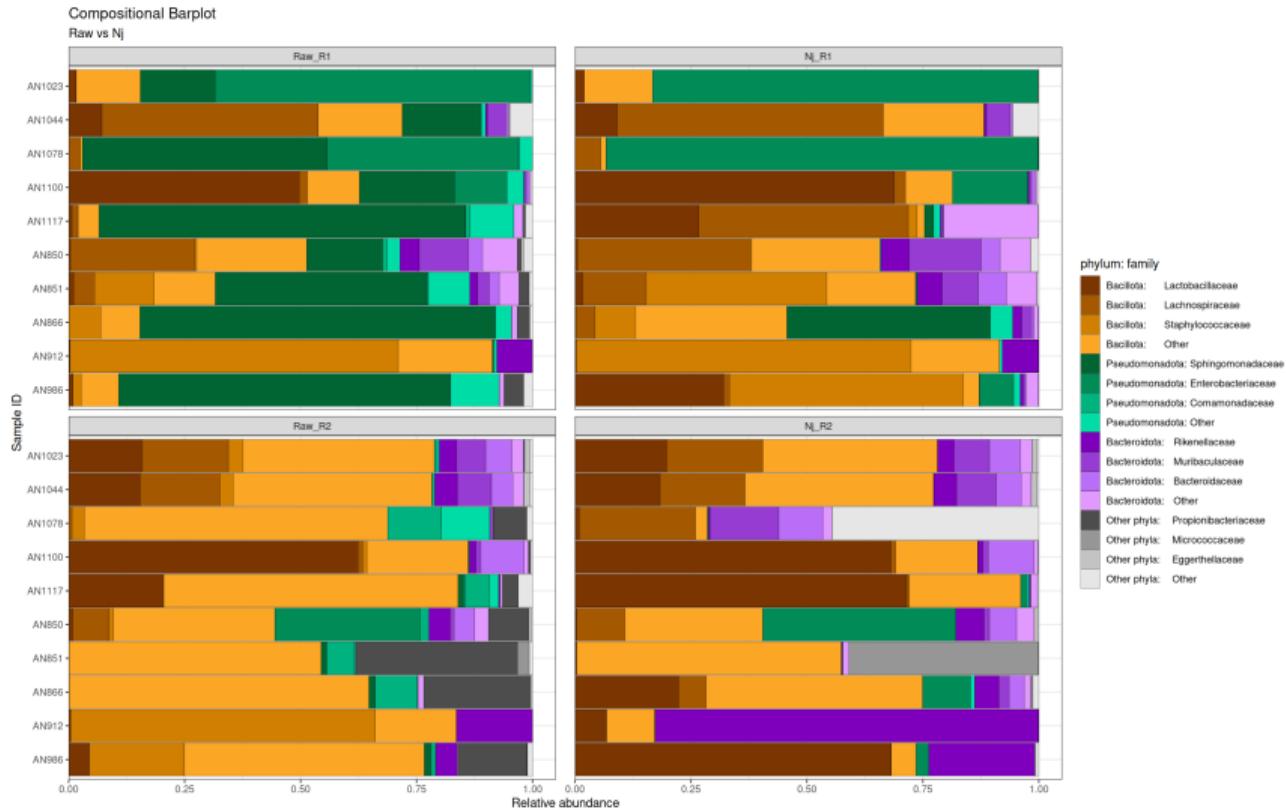
# Overlap - Microbiome profile of PDAC in mice



# Core microbiome after Nj approach



# Core microbiome after Nj approach



# Conclusions

## Decontamination Methods

- Negative control samples are essential to avoid spurious taxa.

## Intratumor microbiome in PDAC

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- Re-sequencing and various decontamination methods with proper negative controls could yield a reliable set of intratumor microbiota.

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- Negative control samples are essential to avoid spurious taxa.
- Each batch has its own contaminant profile.
- In our study, thoroughly remove contaminant such as Nejman et al. procedure outperforms others.

## Intratumor microbiome in PDAC

- Re-sequencing and various decontamination methods with proper negative controls could yield a reliable set of intratumor microbiota.
- Almost identified taxa are well-known gut microbiota.

# Acknowledgements

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## Section 3

Backup Slides

# Core microbiome - All decontamination method - phylum: family

