

# Emergence of biocide resistance in *Salmonella* Typhimurium via directed evolution experiments

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## 1 INTRODUCTION

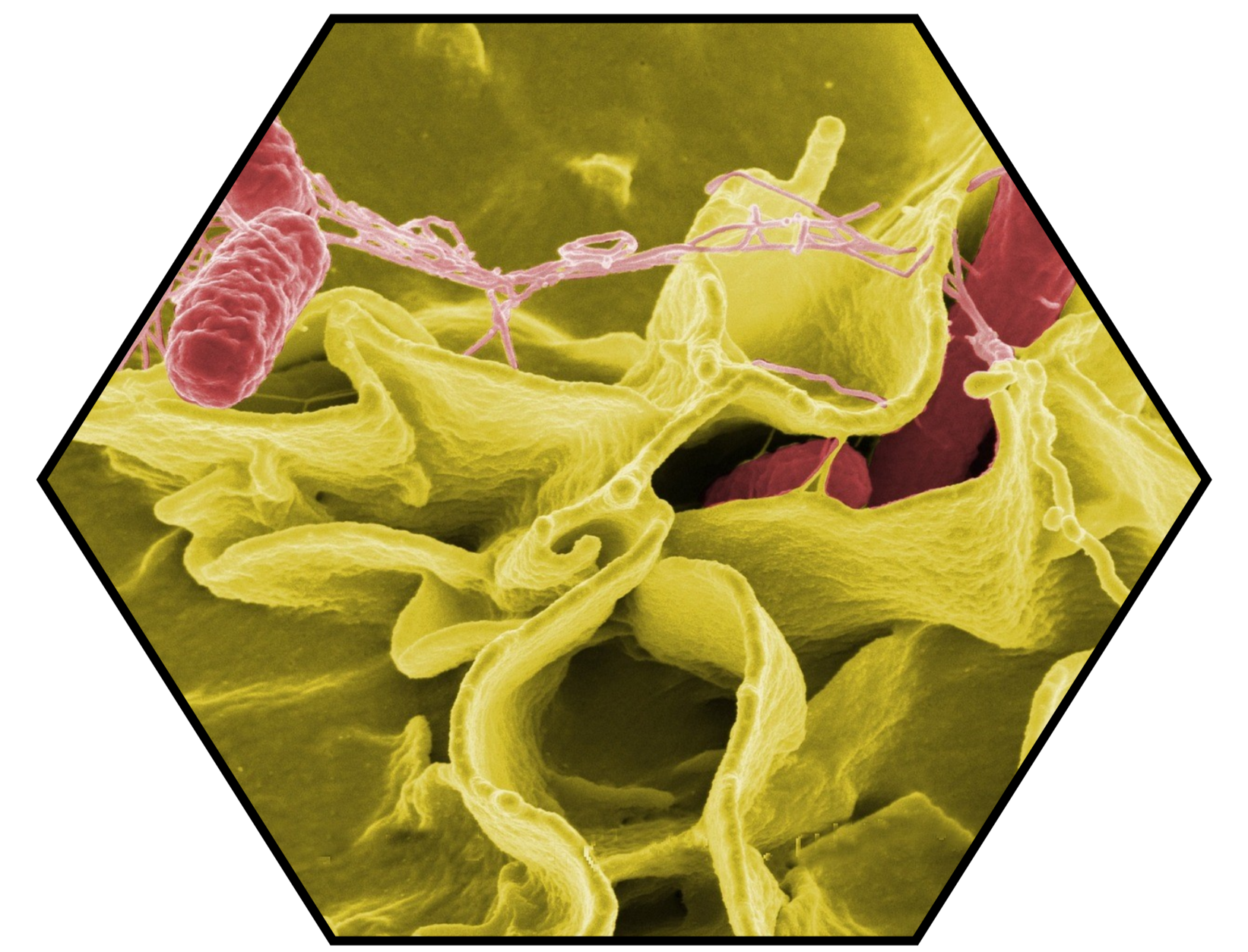
*Salmonella* spp. is a biotic agent responsible for food-borne outbreaks. It is the second most reported toxin-infection, causing more than half of all hospitalizations related to foodborne. The use of antibiotic treatments for clinical cases of salmonellosis may lose effectiveness due to the emergence of antimicrobial resistance.

The phenomenon of antibiotic resistance is considered a global problem due to the threat it poses to public health, so it is being widely studied. However, it is uncertain whether enhanced resistance might occur against other antimicrobial compounds, such as biocides.

Among the available biocides, peracetic acid (PAA) is one of the main biocides used as surface disinfectants in the food industry. The mechanism of its biocidal activity consists of the oxidation of components of its cell membrane components, mainly membrane proteins, which leads to instability and subsequent cell lysis.

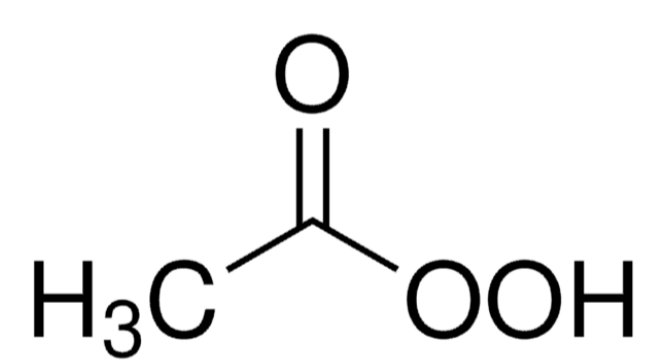
## 2 OBJECTIVE

This study aims to determine whether peracetic acid (PAA) exposure could lead to the emergence of resistant variants in *Salmonella* Typhimurium populations.



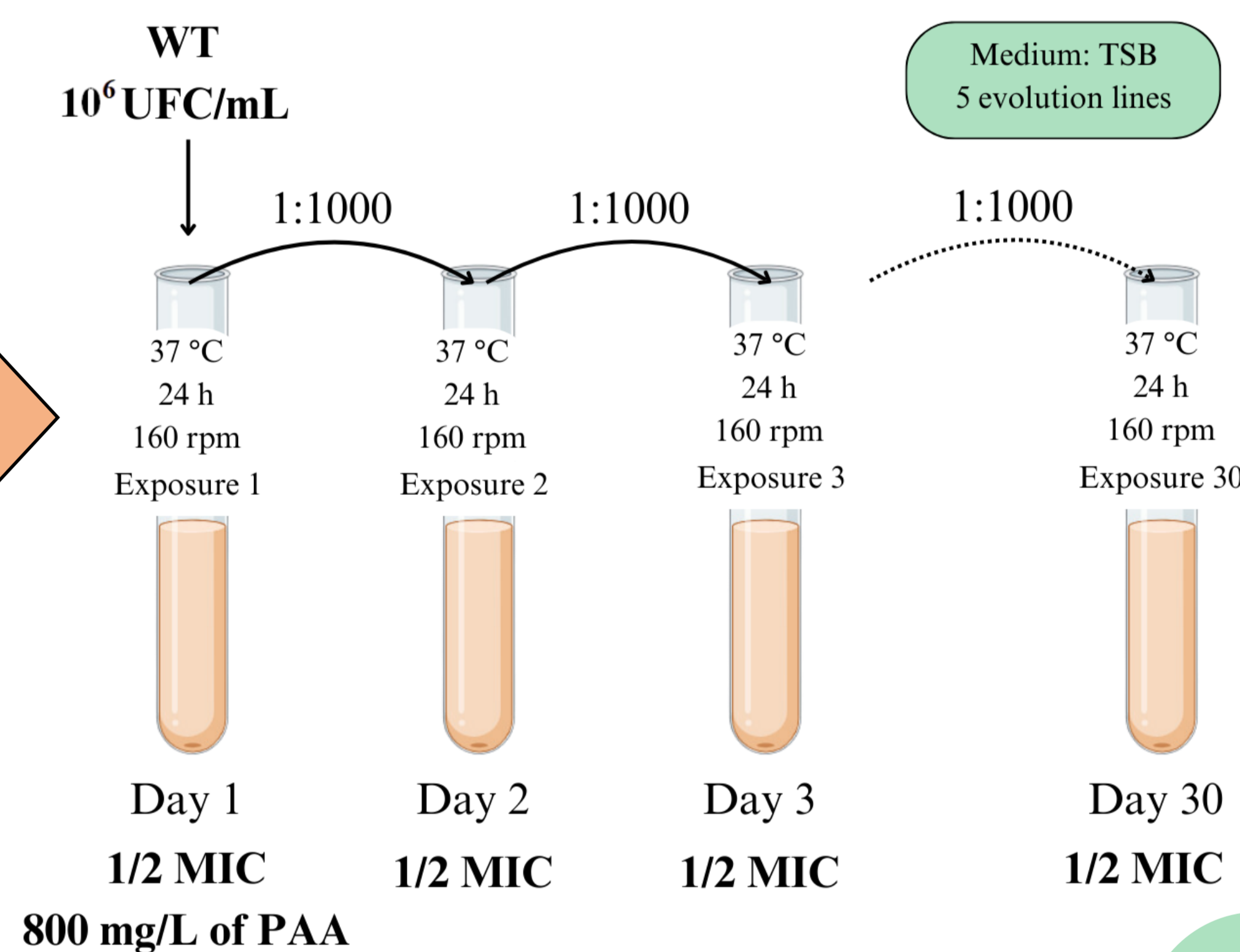
## 3 MATERIAL & METHODS

Peracetic acid  
(Supelco, Alemania)

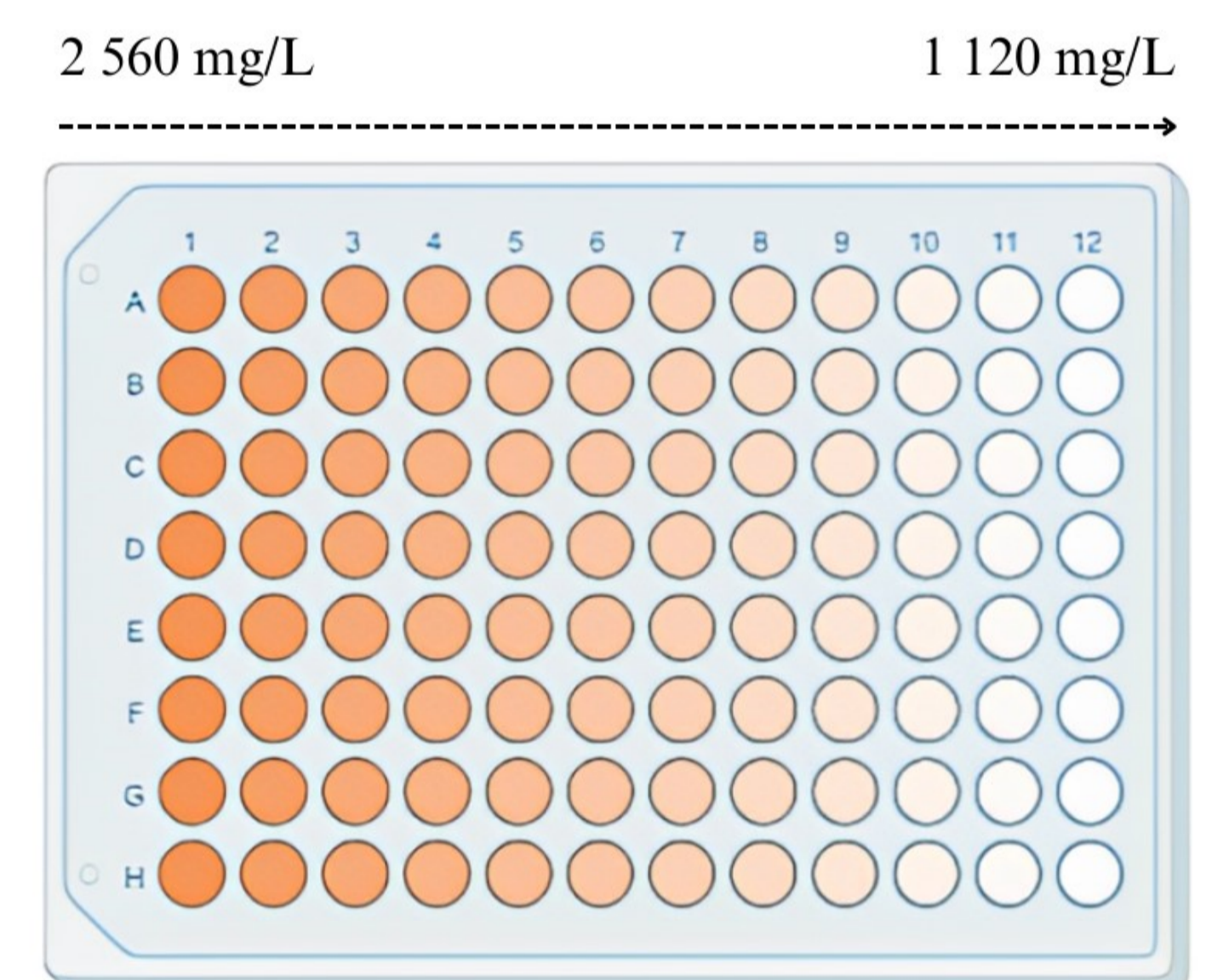


Adaptative  
Laboratory  
Experiment ALE

*Salmonella enterica* subsp. *enterica*  
serovar Typhimurium str. LT2  
(CECT 722)



Minimum  
Inhibitory  
Concentration MIC



Turbidity is measured at 620 nm to determine the MIC  
MIC determination of evolved populations and isolated strains

## 4 RESULTS

**Table 1.** MIC comparison between parental strain of *S.* Typhimurium (WT) and evolved populations.

Bacterial Populations	MIC (mg/L)
<i>S.</i> Typhimurium	1600
1P	1760
2P	1760
3P	1760
4P	1760
5P	1760

The 5 evolved populations showed a 10% increase in direct resistance against PAA after a 30-day exposure to subinhibitory concentrations.

## 5 CONCLUSIONS

Prolonged exposure to subinhibitory doses (1/2 MIC) of PAA leads to increased resistance in populations but this is not translated into the isolation of resistant variants. However, more studies are needed to understand in depth the relationship between the use of PAA and the emergence of resistance to biocides.

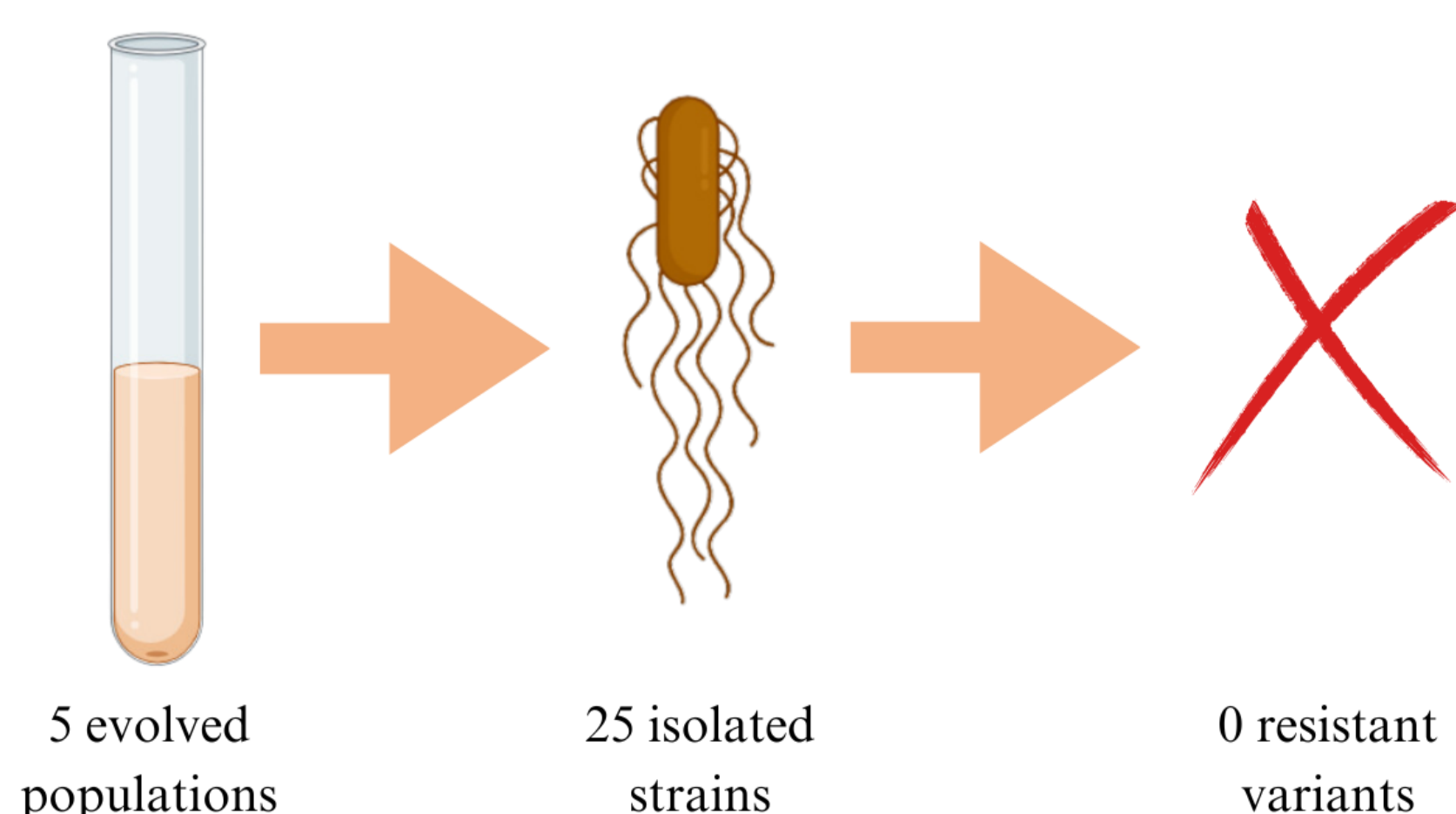
In addition, it would be interesting to evaluate the occurrence of cross-resistance to antibiotics or food preservation methods, as well as other methodologies of adaptive laboratory experiments, such as lethal treatments.

## 6 KEY REFERENCES

Despite the increased resistance observed in the 5 evolved populations, no resistant variants to PPA were detected among 25 isolated strains selected

This phenomenon is consistent with previous findings described by Lopatkin *et al.* (2021).

Berdejo, D., Pagán, E., Merino, N., García-Gonzalo, D. and Pagán, R. (2021). "Evolution Assays for the Isolation of Mutant Bacteria Resistant to Natural Antimicrobials". In: MAGNANI, M. (ed.) *Detection and Enumeration of Bacteria, Yeast, Viruses, and Protozoan in Foods and Freshwater*. New York, NY: Springer US. 65-75 pp.  
European Food Safety Authority (2023). "The European Union One Health 2022 Zoonoses Report". *EFSA Journal*, 21, pp. 30-65.  
Lopatkin, A. J., Bening, S. C., Manson, A. L., Stokes, J. M., Kohanski, M. A., Badran, A. H., Earl, A. M., Cheney, N. J., Yang, J. H. and Collins, J. J. (2021). "Clinically relevant mutations in core metabolic genes confer antibiotic resistance". *Science*, 371(6531), 8 pp.



**Figure 1.** Selection of resistant variants from PAA-exposed populations.