

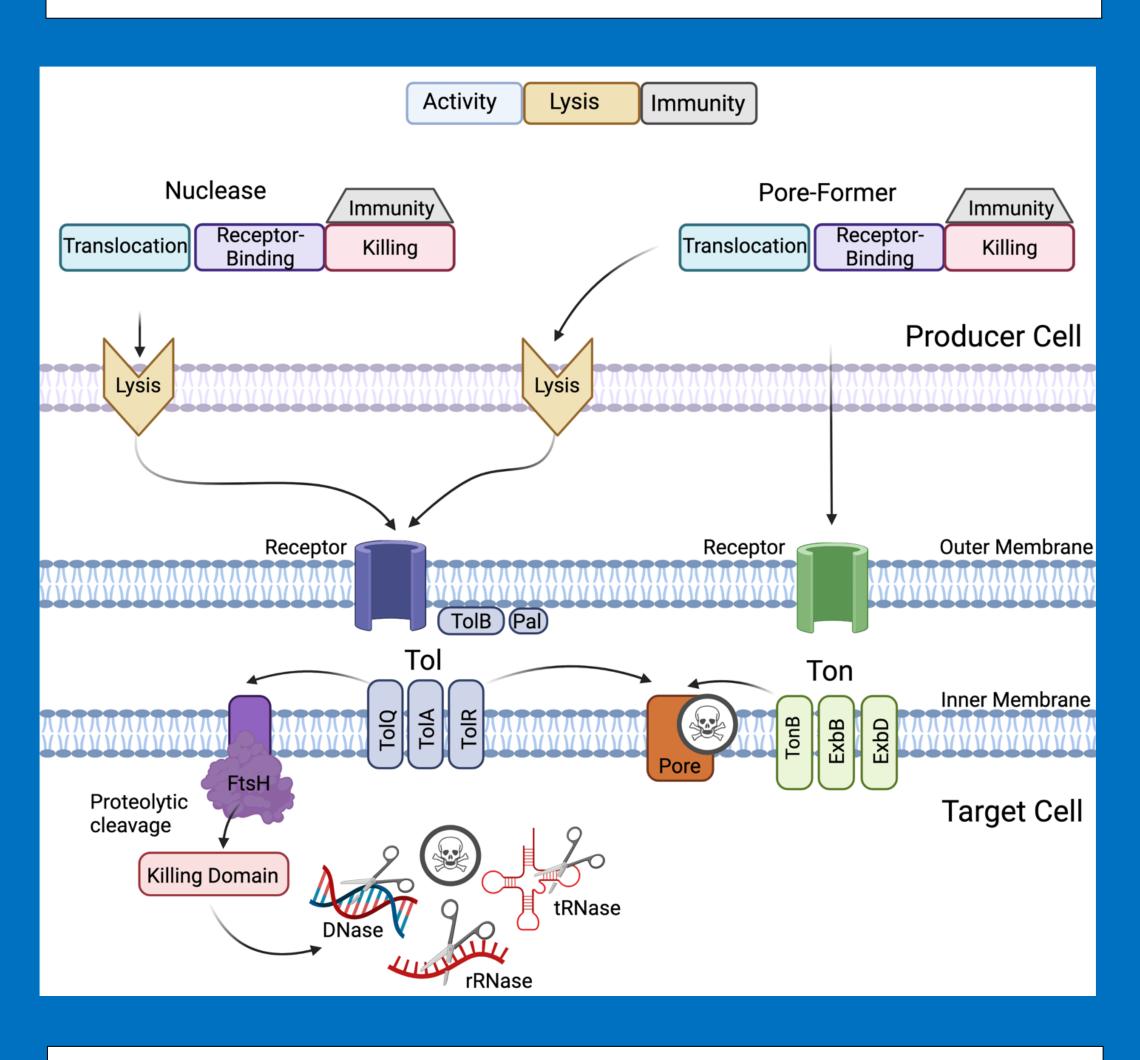


## Introduction

### **Background:**

- Antibiotic resistance is a major problem the world faces.
- Bacteriocins are antimicrobials produced by bacteria that kill other bacteria.
- Colicins are bacteriocins produced by *E*. *coli* and are a model system for bacteriocin evolution.
- Colicins can kill by pore formation or nuclease activity.

**Objective/Goal:** How do colicins evolve, and what does their evolution tell us about bacteriocins as a long-term solution to antibiotic resistance?



# Methods

- Obtained protein sequences from online databases (NCBI and UniProt)
- Aligned protein sequences in each protein domain with MEGA-X
- Created and analyzed phylogenetic trees

## We thank all sponsors of iCons

# **Colicin Evolution: Lessons from Billions of Years of Bacterial Warfare** Allison Brookhart, Sofia Blomqvist, and Margaret Riley\* Department of Biology, UMass Amherst

## **Results**

0.7

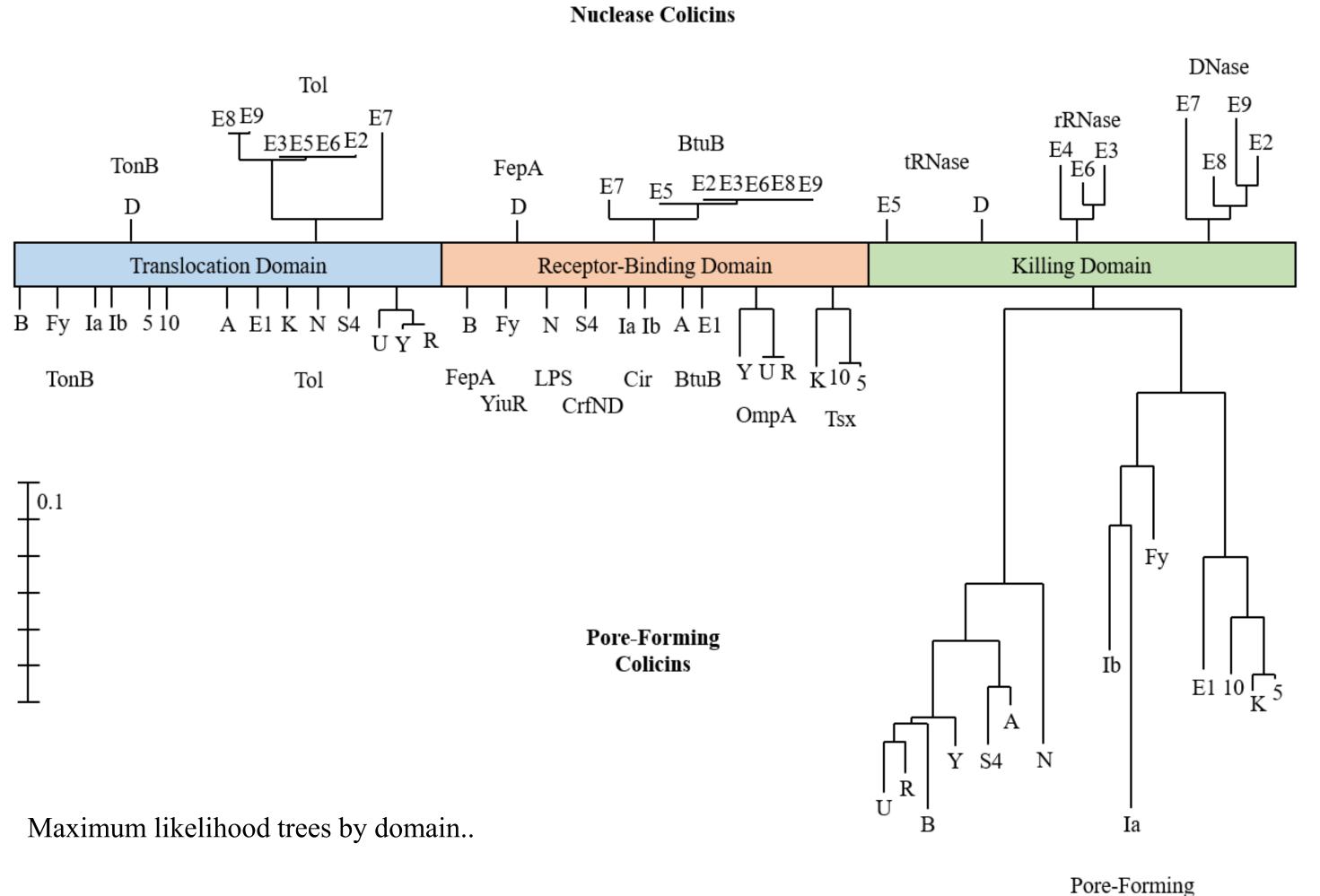
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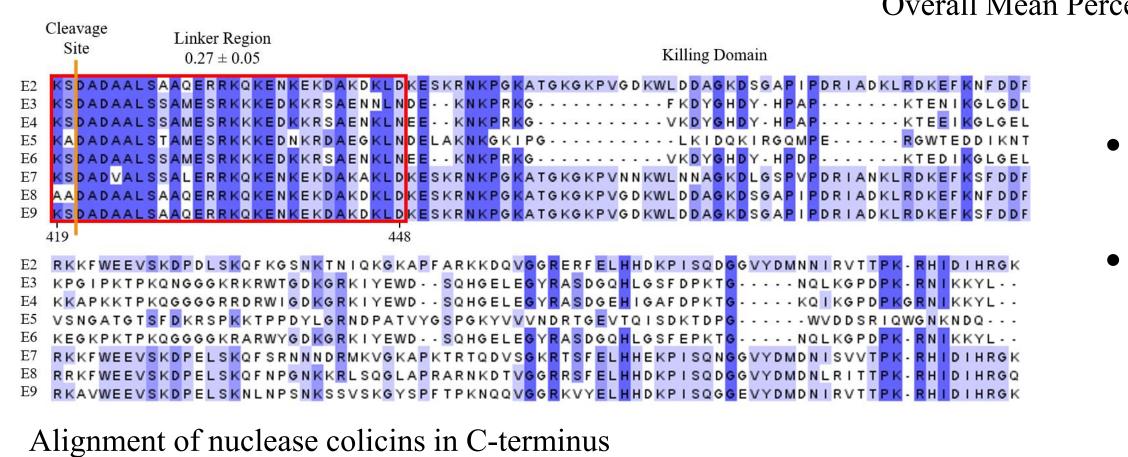
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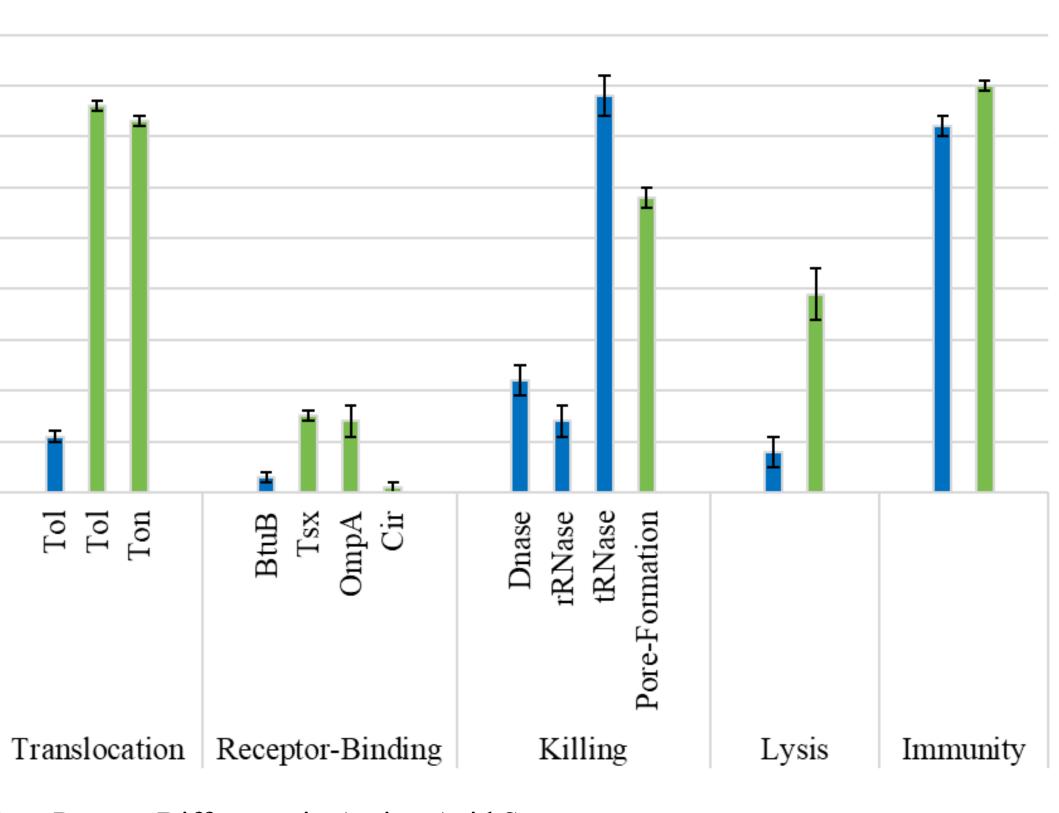


- Differences in killing mechanism can explain this variation.
- Nucleases must cross both outer and inner membranes to access nucleic acids in the cytoplasm and kill the cell, so nuclease cell entry options may be more limited.
- Pore-formers kill by inserting themselves in the inner membrane, so this killing mechanism can stay the same but cell entry methods must change to avoid resistance.
- The least costly way to evolve resistance is receptor mutations, which drives poreformers to use novel receptors.



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- Pore-forming colicins can use more receptors and translocation systems and have more sequence dissimilarity, suggesting that **pore-formers are** older than nuclease colicins.
- Pore-formers and nucleases experience different selective pressures.
- Pore-formers have most differences in translocation and receptor-binding domains, whereas nucleases are most different in the killing domain.



Pore-Formers

Overall Mean Percent Difference in Amino Acid Sequence

Nucleases

- Nuclease colicins must be cleaved in order to enter the cytoplasm.
  - **Proteolytic cleavage occurs at the region** where high sequence dissimilarity starts to occur for nuclease colicins.

# Acknowledgements

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Feldgarden, M., & Riley, M. A. (1999). The phenotypic and fitness effects of colicin resistance in Escherichia coli K-12. Evolution; International Journal of Organic *Evolution*, *53*(4), 1019–1027. https://doi.org/10.1111/j.1558-5646.1999.tb04517.x

Riley, M. A. (1993). Molecular mechanisms of colicin evolution. Molecular Biology and Evolution, 10(6), 1380–1395. https://doi.org/10.1093/oxfordjournals.molbev.a040081







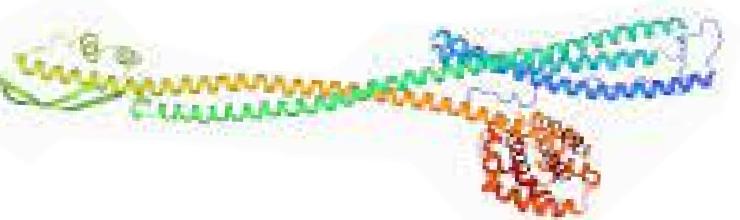


### Discussion

### **Conclusions:**

• Understanding where selective pressure encourages evolution will help us best use colicins in the future

• Ex: we can predict potential receptors • Resistance to any antimicrobial will evolve, but understanding how and when this resistance arise helps us better plan for the future.



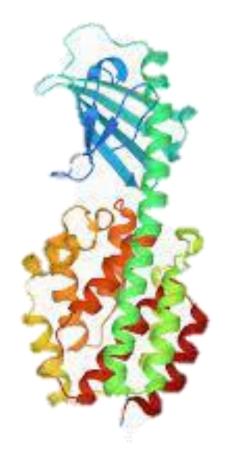
### **Future Directions:**

• Does this model of evolution apply to other bacteriocins, such as pyocins?

• How does colicin evolution affect killing ability?

• Using this information when genetically engineering bacteriocins or designing

bacteriocin-based technologies



## References