The Role of *nhaP1, nhaP2,* and *nhaP3* Antiporters in the Acid Tolerance of Vibrio cholerae.

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Oregon State

Cholera

- Extreme diarrhea
- Rehydration therapy
- Vaccines



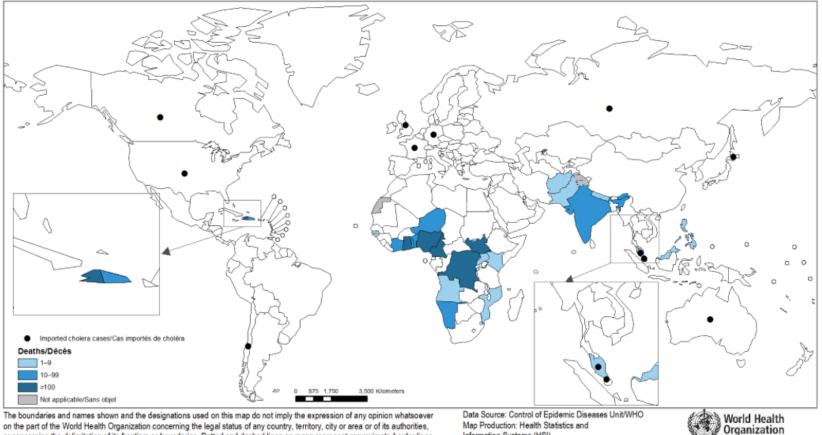




(Knobil, 2015)

Cholera

• 1.4 – 4.3 million cases per year – 28,000 to 142,000 deaths



on the part of the World Health Organization concerning the legal status of any country, territory, city or area or of its authorities, or concerning the delimitation of its frontiers or boundaries. Dotted and dashed lines on maps represent approximate border lines for which there may not yet be full agreement.

Map Production: Health Statistics and Information Systems (HSI) World Health Organization

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Cholera in Haiti

- 2010 Earthquake
 - >6% of Haitians have been infected
 - First two years: >8, 231 deaths
 - Up until 2015, >700,000 cases



Vibrio cholerae

Vibrio cholerae 0395N1 (Vc)

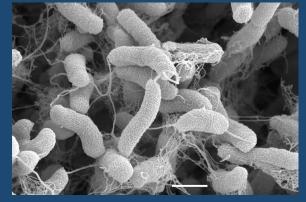
 Gram negative rod
 Marine/Brackish waters





(Munoz, 2014)

(WHO, 2014)



(Robert Koch Institute, 2014)

The Traveling Organism



(J. Silva/Reuters, 2011)

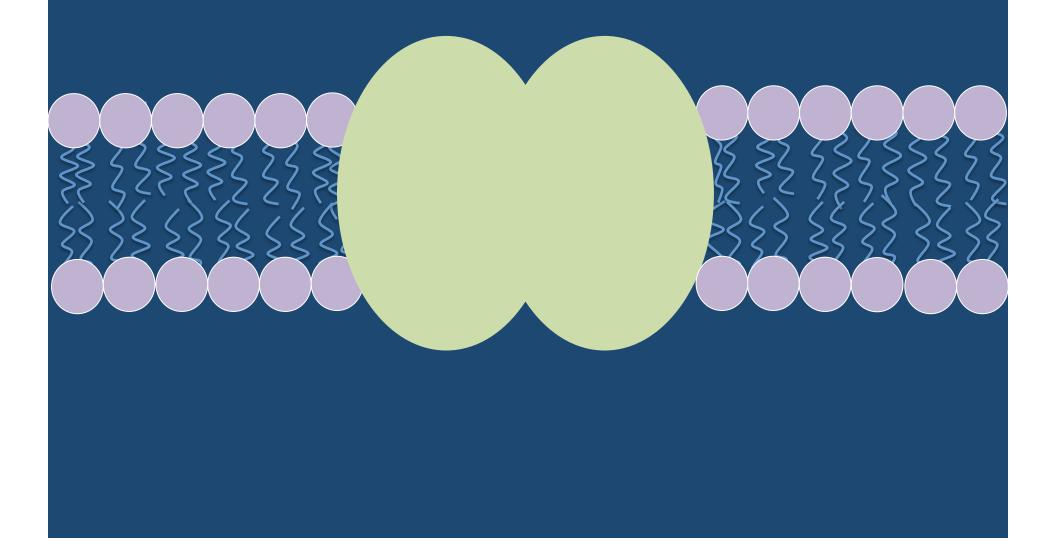
The Traveling Organism

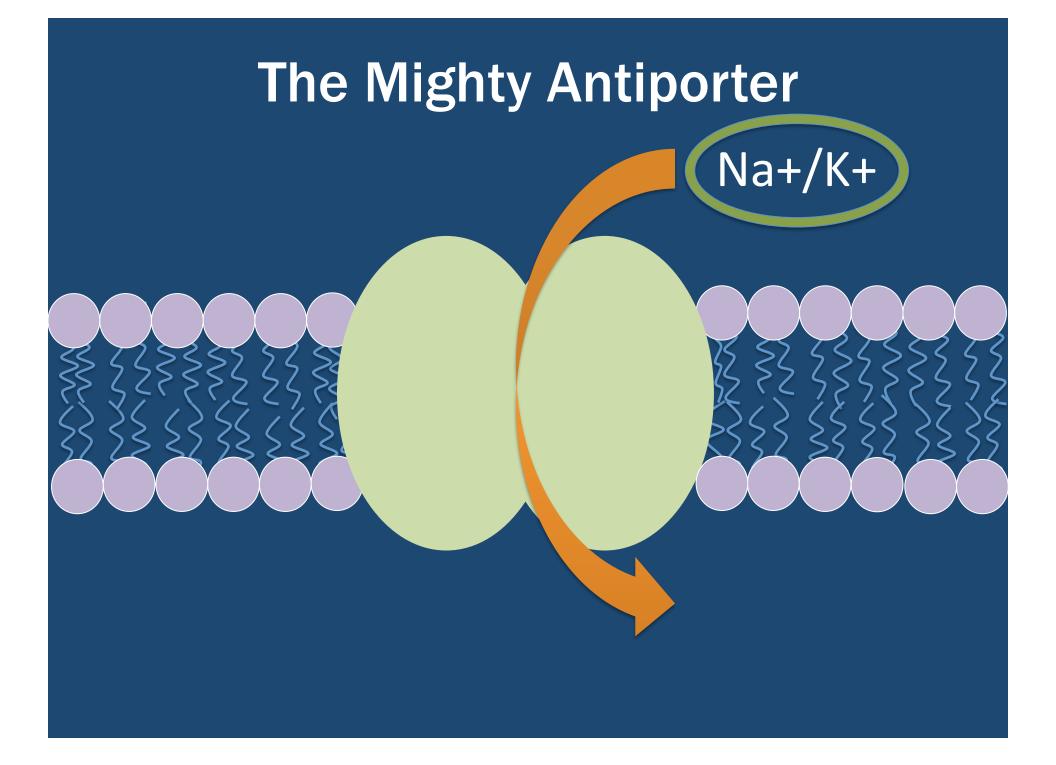


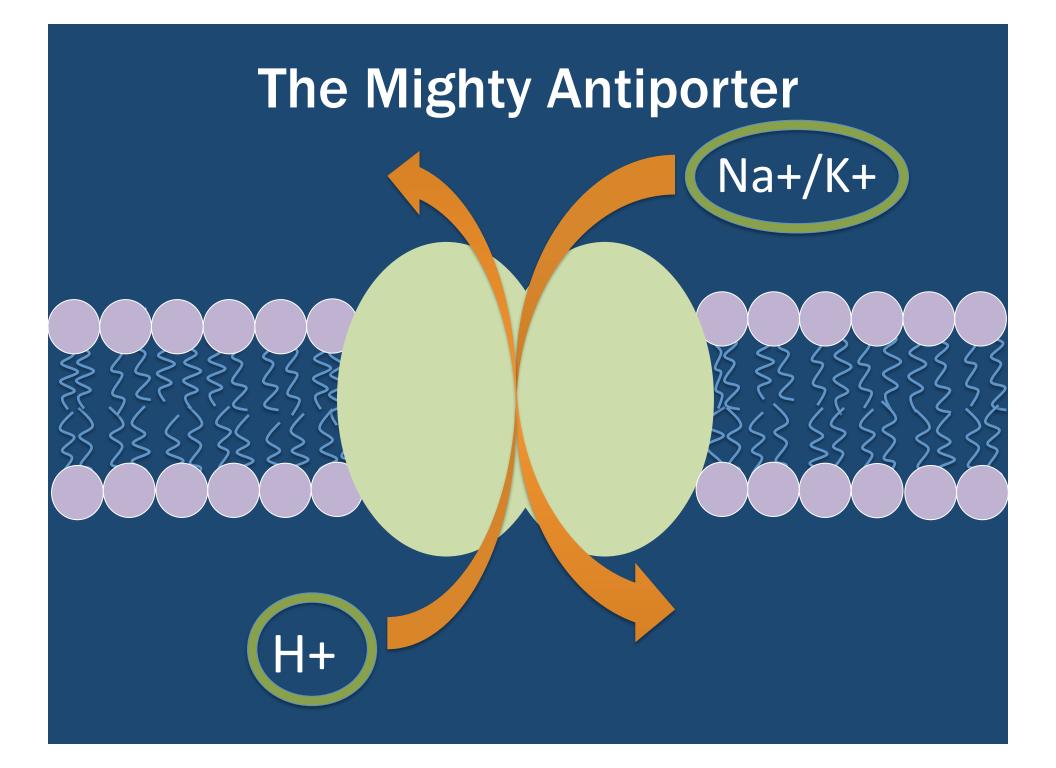
The Traveling Organism

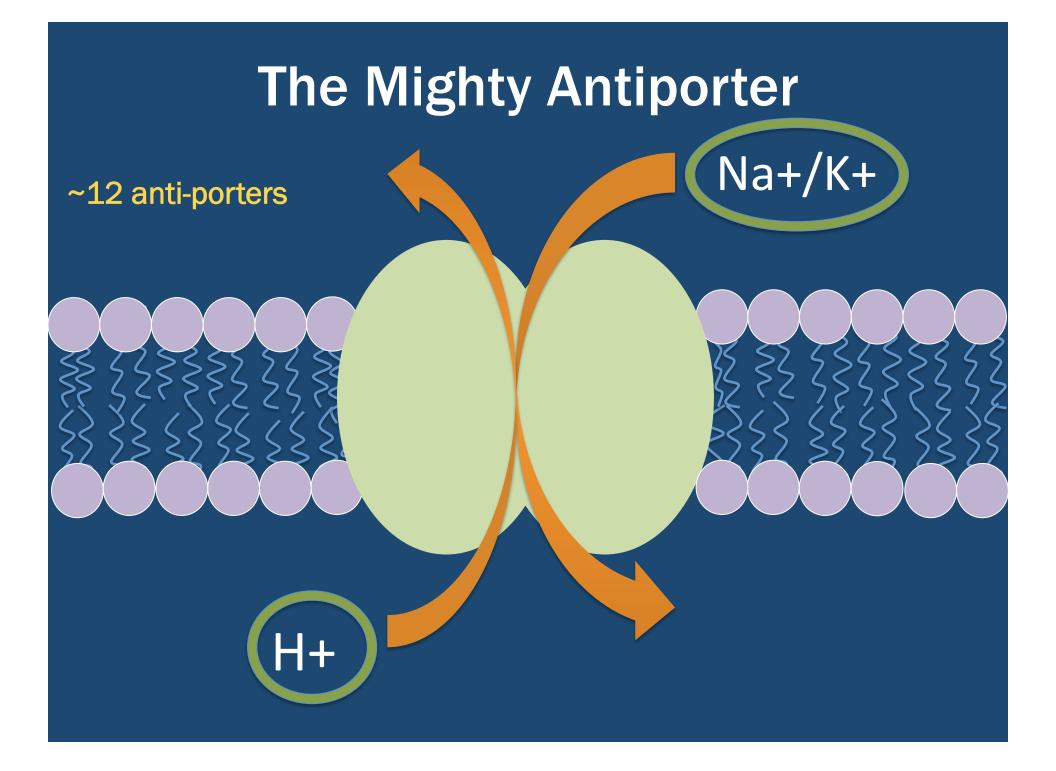


Ion Homeostasis & Antiporters









The Mighty Antiporter

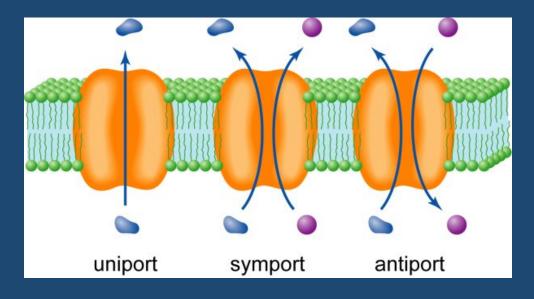
Na+/K+

~12 anti-porters -nhaP1, nhaP2, nhaP3

H+

Expected Outcomes and Impact

- Understanding
- Found in Yersinia pestis and Pseudomonas aeruginosa
- Novel intervention therapies



Question & Hypothesis

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- We hypothesized that the three genes, nhaP1, nhaP2, and nhaP3, in combination, are required for the survival of Vibrio cholerae through the extreme acidity of the human stomach.



• Wild type

nhap1

- Vibrio cholerae 0395 N1

nhap3

nhaP2

- Wild type
 Vibrio cholerae 0395 N1
 - nhapt nhapt
- Triple mutant
 - Vibrio cholerae 0395 N1 $\Delta nhaP1\Delta nhaP2\Delta nhaP3$

Wild type
 Vibrio cholerae 0395 N1

nhaP2

nhaloo

nhap1

Triple mutant

phaP1

- Vibrio cholerae 0395 N1 $\Delta nhaP1\Delta nhaP2\Delta nhaP3$

Phapo 3

nhaP2

Wild type
Vibrio cholerae 0395 N1

nhaP2

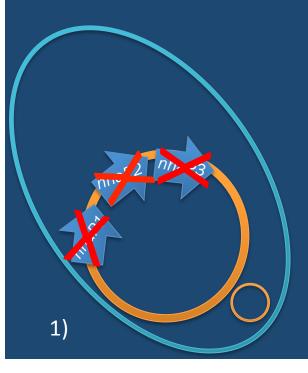
nhapos

nhaP1

- Triple mutant
 - Vibrio cholerae 0395 N1 $\Delta nhaP1\Delta nhaP2\Delta nhaP3$

- Complement P1, Complement P2, Complement P3 & Wild type with empty plasmid
 - 1) Vc Δ*nhaP1*Δ*nhaP2*Δ*nhaP3* <pBAD24::nhaP1>
 - 2) Vc $\Delta nhaP1\Delta nhaP2\Delta nhaP3 < pBAD24::nhaP2>$
 - 3) Vc Δ*nhaP1*Δ*nhaP2*Δ*nhaP3* <pBAD24::nhaP3>
 - 4) Vc pBAD24

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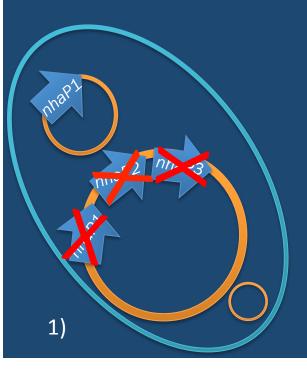
 Complement P1, Complement P2, Complement P3 & Wild type with empty plasmid

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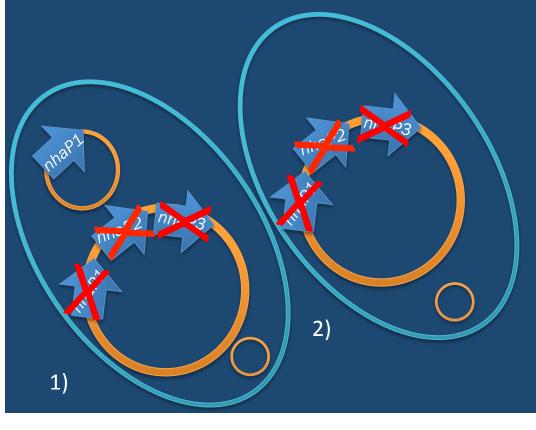
2) Vc $\Delta nhaP1\Delta nhaP2\Delta nhaP3 < pBAD24::nhaP2>$

3) Vc Δ*nhaP1*Δ*nhaP2*Δ*nhaP3* <pBAD24::nhaP3>

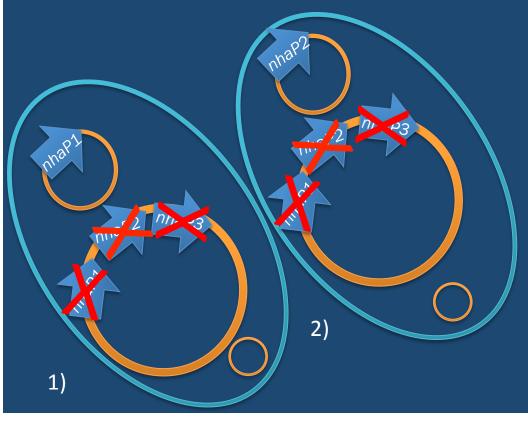
4) Vc pBAD24



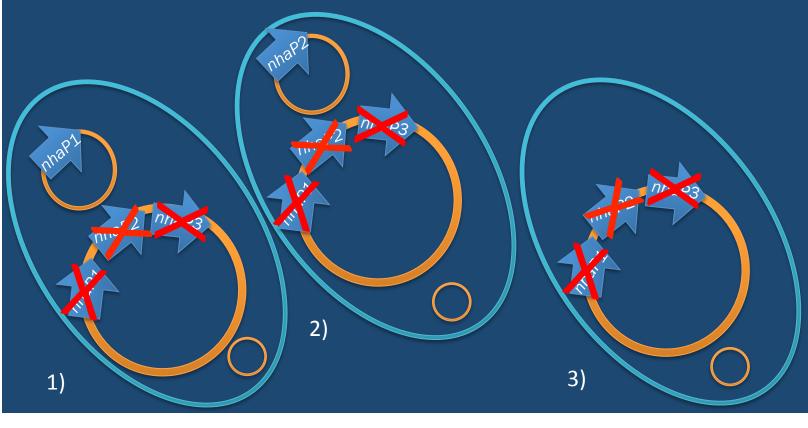
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- 4) Vc pBAD24



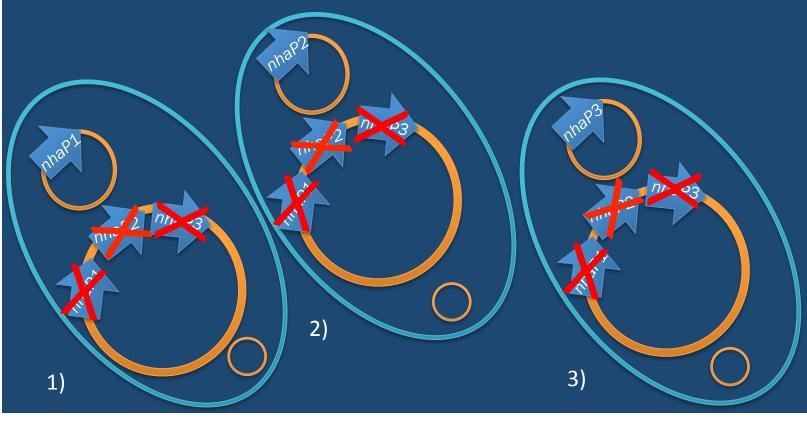
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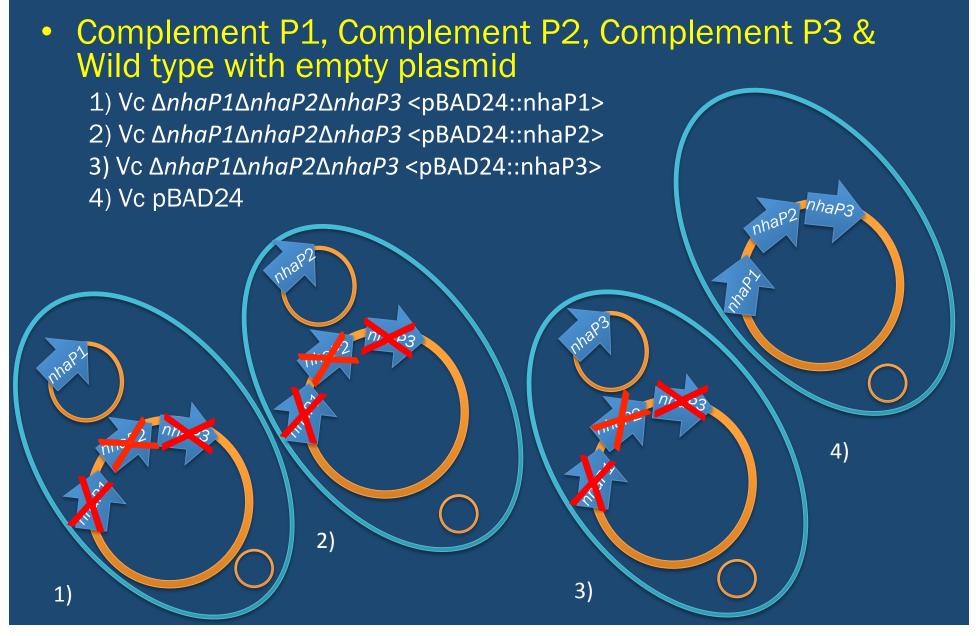


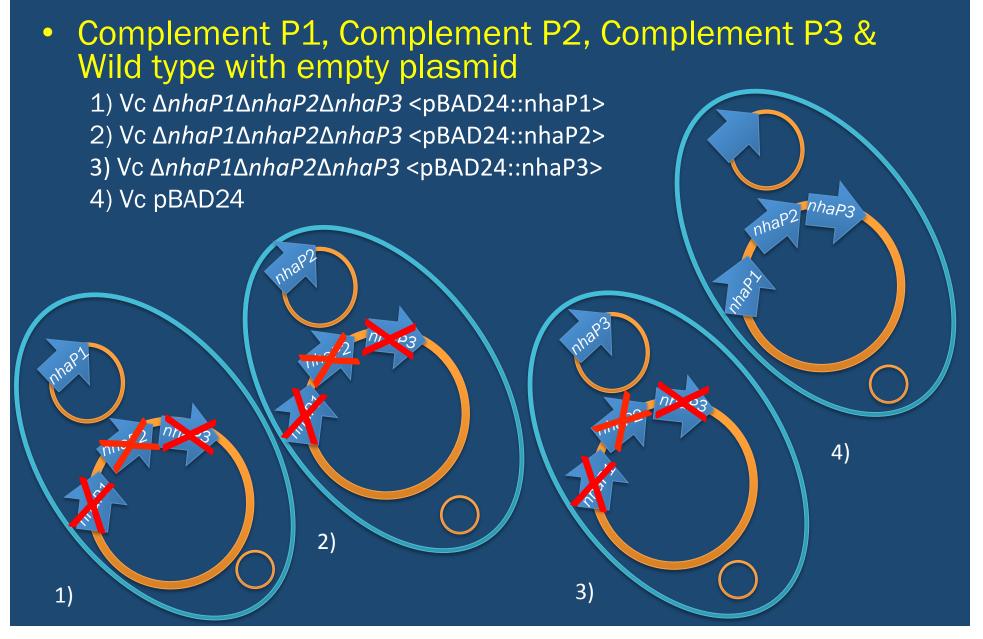
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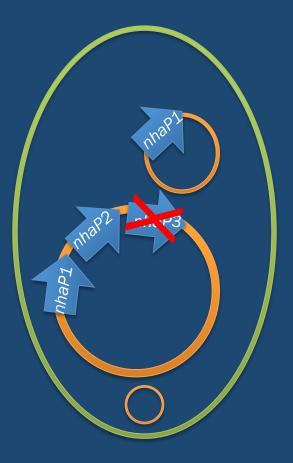


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- 3) Vc Δ*nhaP1*Δ*nhaP2*Δ*nhaP3* <pBAD24::nhaP3>
- 4) Vc pBAD24









Vibrio cholerae 0395 N1 ΔnhaP3<pBAD24::nhaP1>

Extraction with the Qiagen Plasmid Kit

Vibrio cholerae 0395 N1 ΔnhaP3<pBAD24::nhaP1>

hap

Extraction with the Qiagen Plasmid Kit

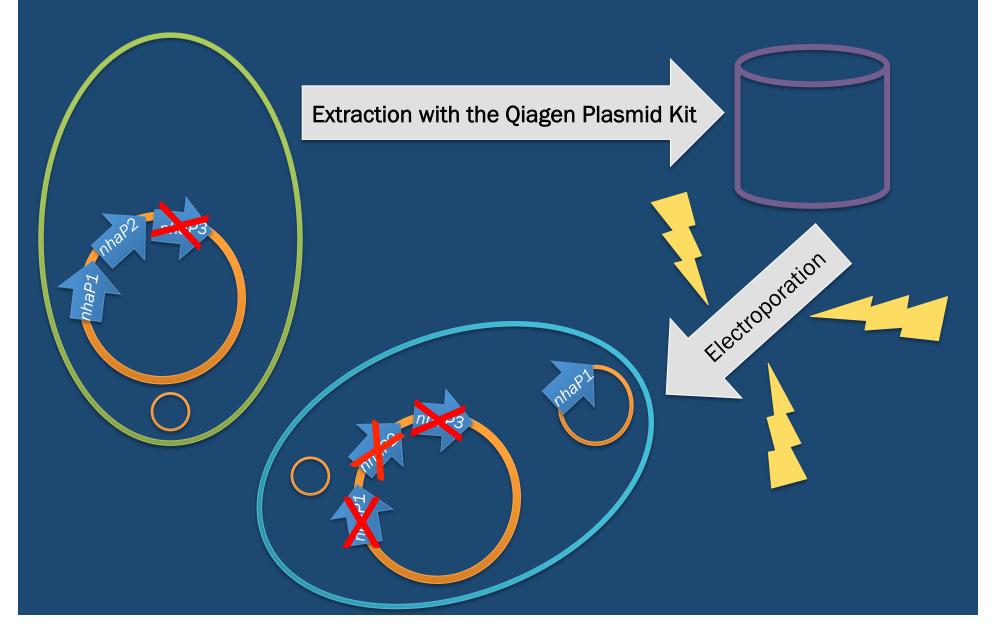
hap



Extraction with the Qiagen Plasmid Kit

hap





Methods

- Acid Tolerance Assays
 - Conditions:
 - pH 3.0, 3.5, 4.0 & 4.5 LBK
 - 0, 15, 60, 90 minutes





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Note

A 6×6 drop plate method for simultaneous colony counting and MPN enumeration of *Campylobacter jejuni*, *Listeria monocytogenes*, and *Escherichia coli*

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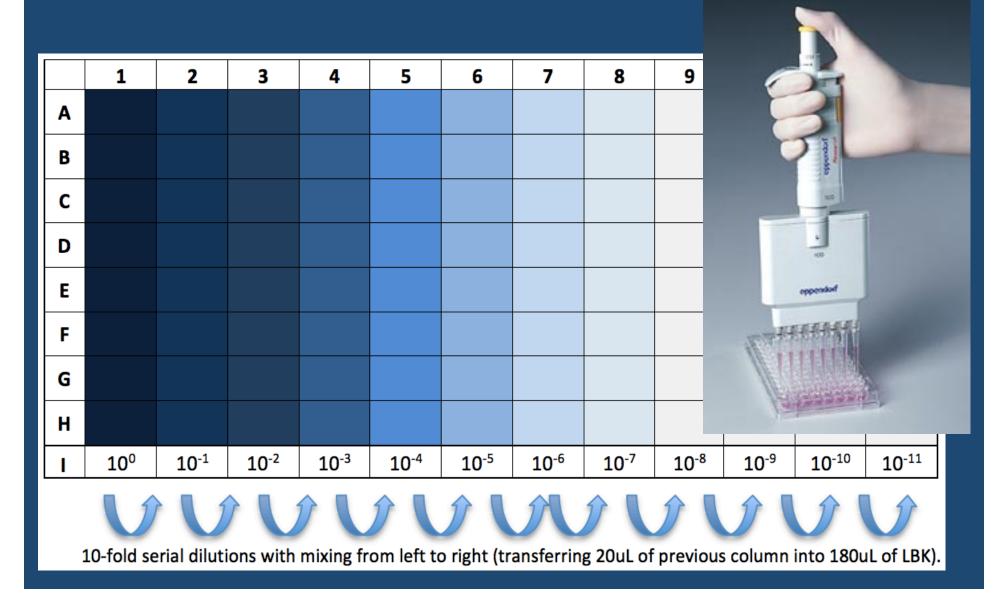
Received 28 January 2003; received in revised form 23 April 2003; accepted 9 June 2003

Serial Dilutions

	1	2	3	4	5	6	7	8	9	10	11	12	
A													
В													
С													
D													
E													
F													
G													
н													
I	10 ⁰	10-1	10-2	10 ⁻³	10-4	10 ⁻⁵	10 ⁻⁶	10 ⁻⁷	10 ⁻⁸	10 ⁻⁹	10 ⁻¹⁰	10-11	
いいいいいいいいいい													
	10-fold serial dilutions with mixing from left to right (transferring 20uL of previous column into 180uL of LBK).												

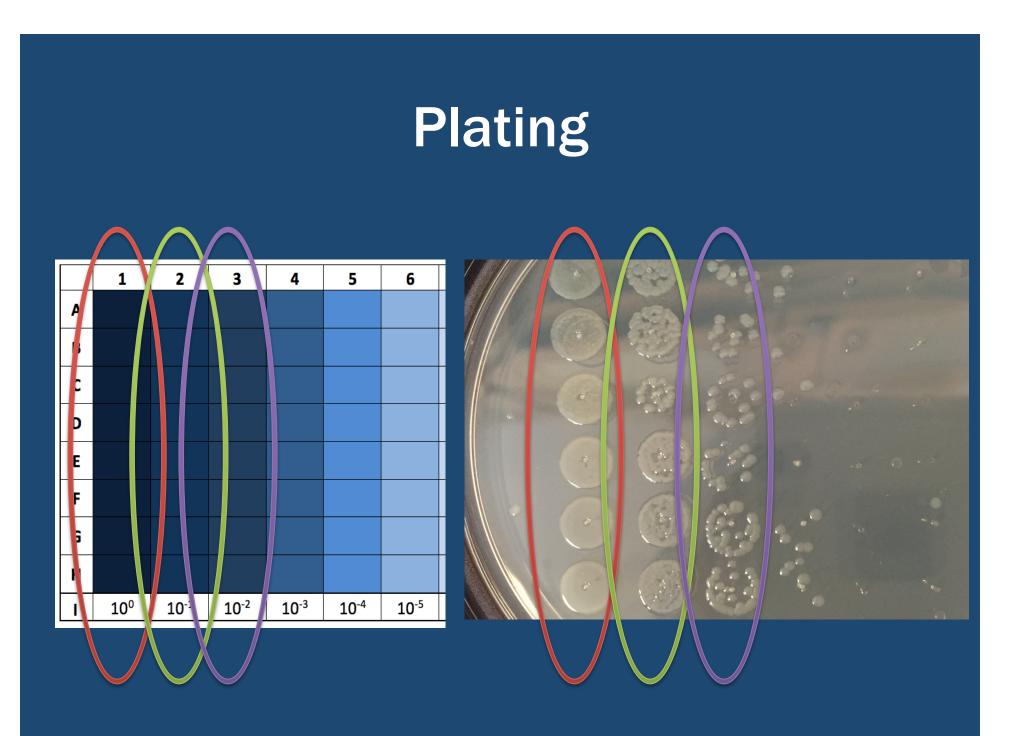
Serial Dilutions

(Biomart, 2007)

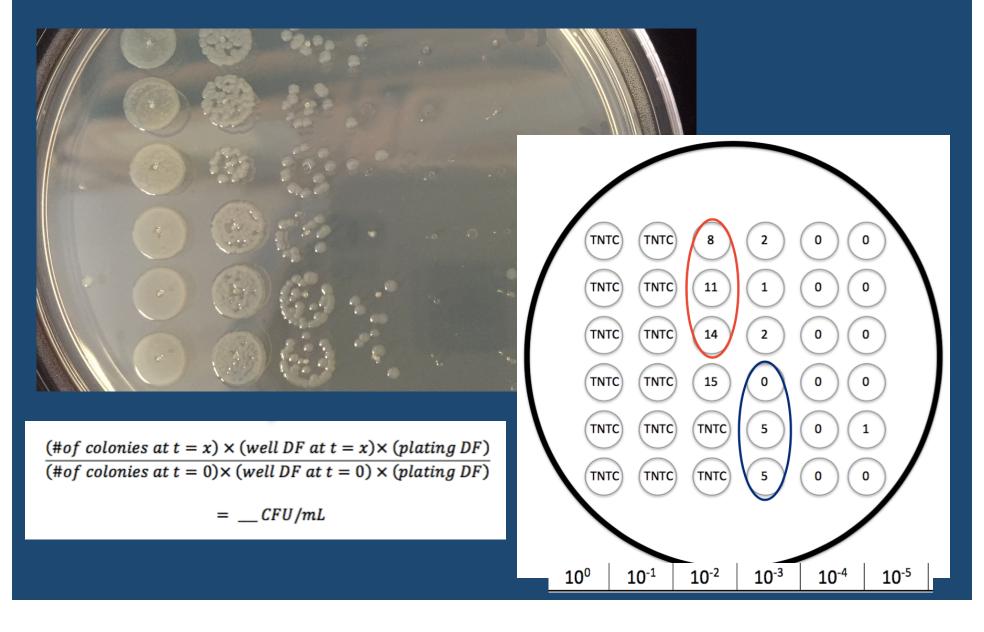


Plating 2 3 4 5 1 6 D Ε 10⁰ **10**⁻¹ **10**⁻² **10**-3 10-4 **10**-5

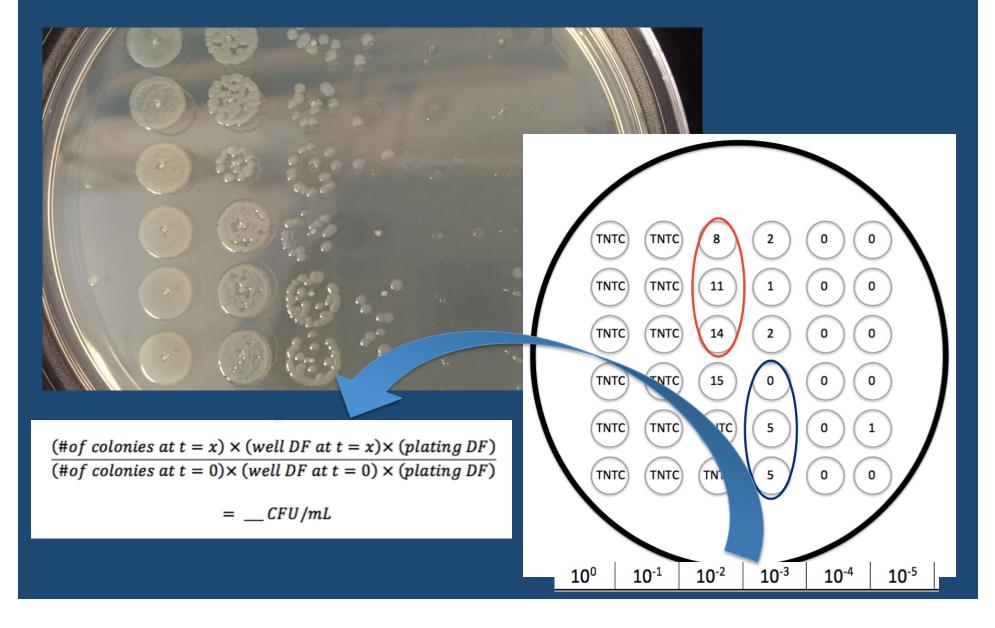
Plating 2 3 4 5 1 6 D Ε 10⁻³ 10⁰ **10**⁻¹ 10⁻² **10**⁻⁴ **10**-5



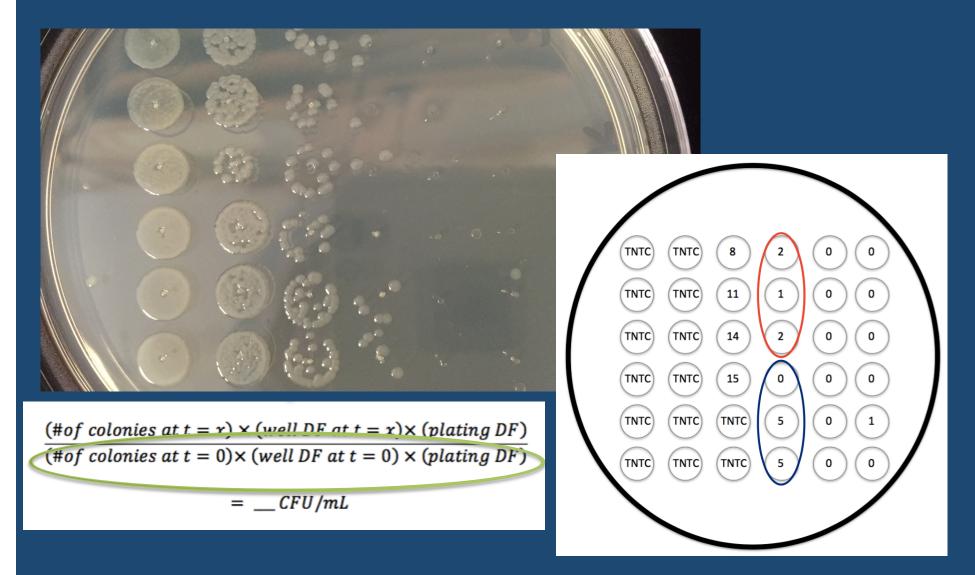
Calculation of Percentage of Cell Survival



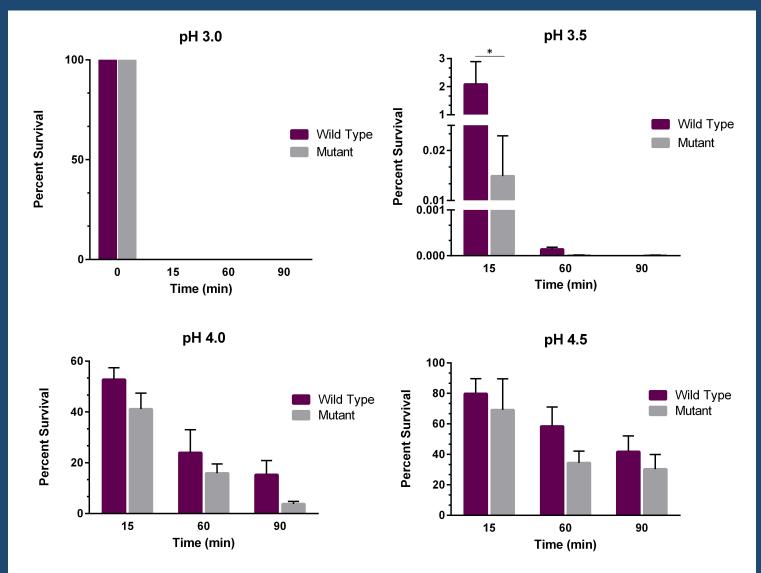
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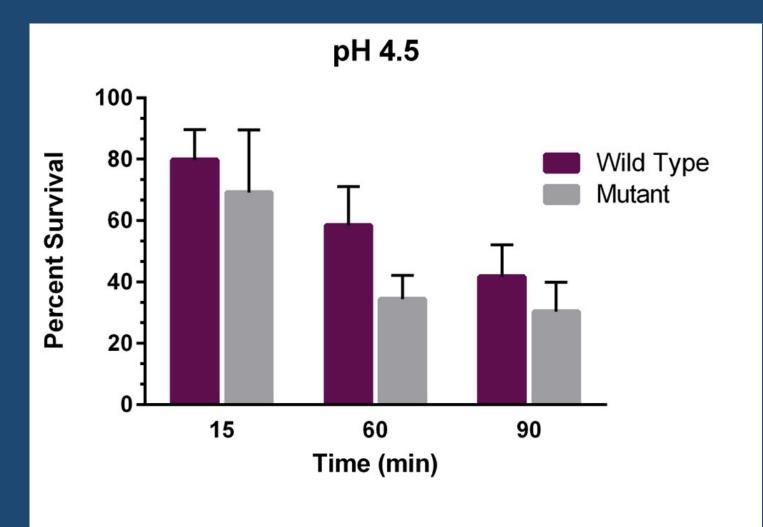


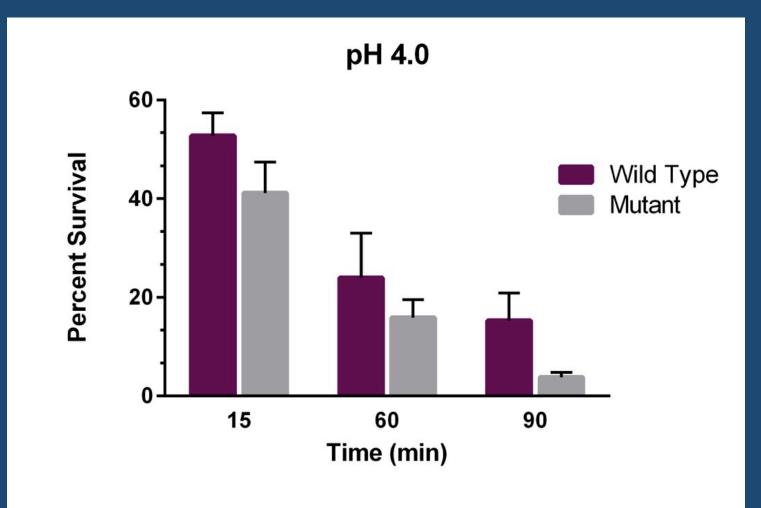
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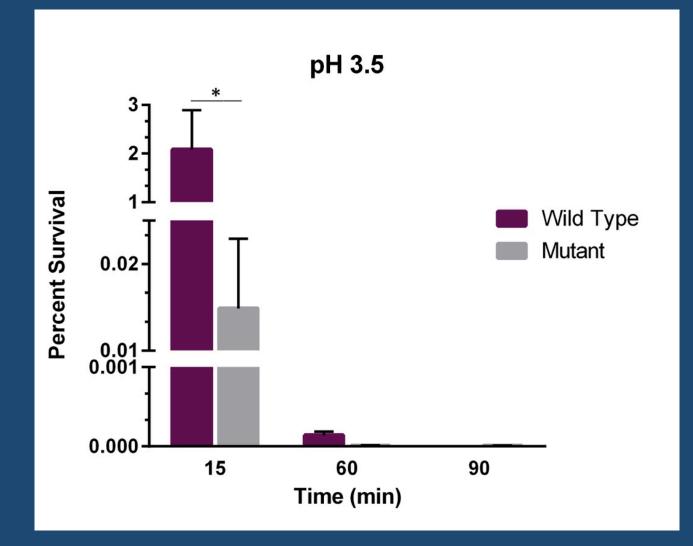


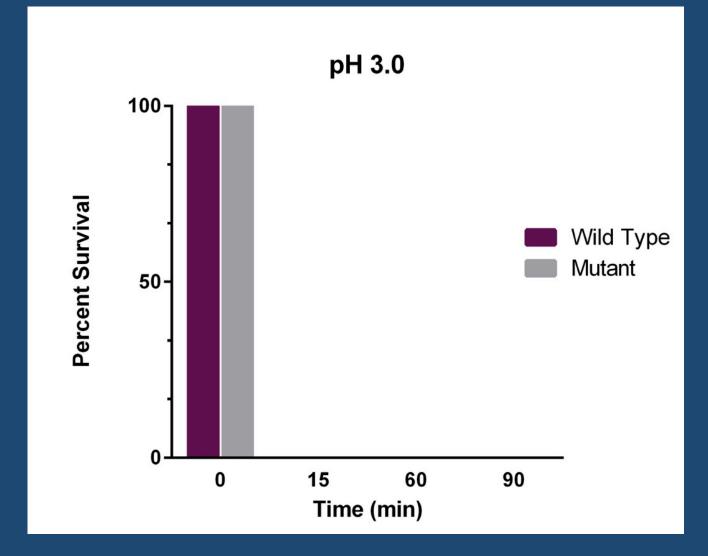
Results



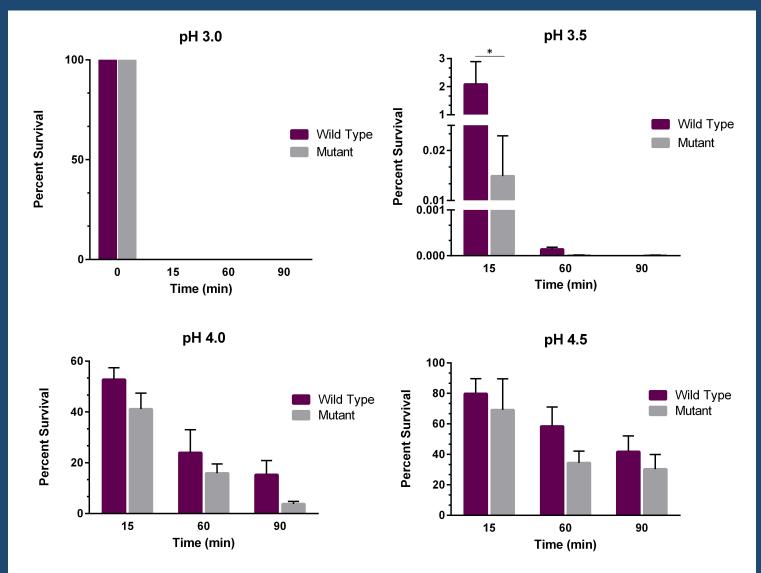




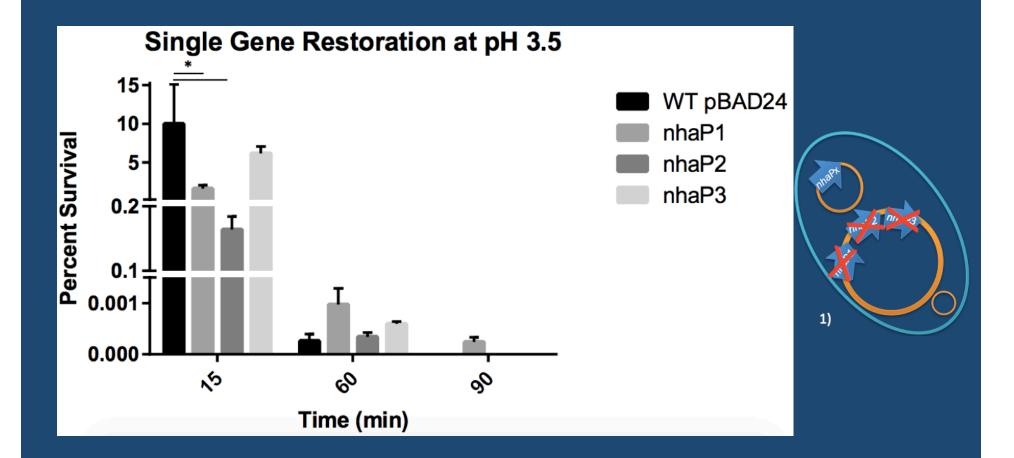




Results



Average Percentage of Survival: Compliment Testing

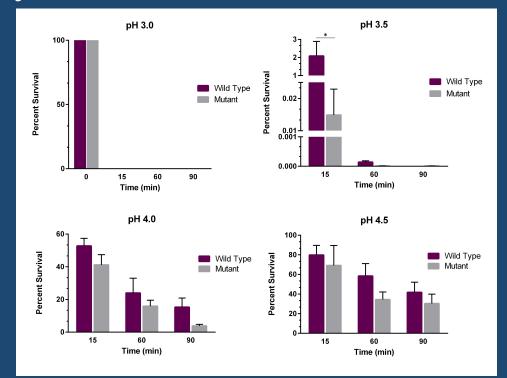


Question & Hypothesis Revisited

- Does the nhaP genes have an effect on the ability of Vibrio cholerae to survive in acidic broths for an extended period of time?
- We hypothesize that the three genes, *nhaP1*, *nhaP2* and *nhaP3*, in combination, are required for the survival of Vibrio cholerae through the extreme acidity of the human gut.

Wild Type versus Triple mutant Conclusion

Having *nhaP1*, *nhaP2* and *nhaP3* intact is beneficial for the organism under extremely acidic conditions but is not necessary for survival.



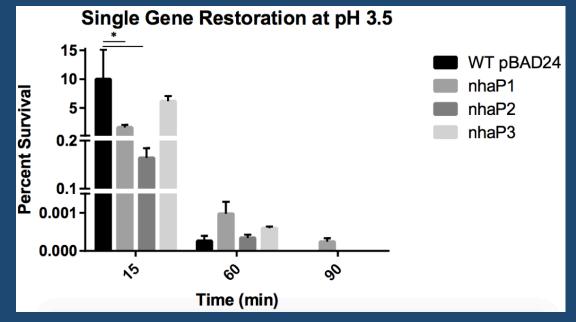
Discussion

- 6x6 Drop Plate Method
 - Saves materials & time
 - Sweet spot
 - Potential error



Complement Strain Conclusions

- Lacking two of the antiporters lead to reduced survival compared to the wild type.
- The strain having *nhaP3* restored survived at the highest rate compared to *nhaP1* and *nhaP2* at 15 minutes and acted very similar to the wild type.



Future Research

- Comparison of results w/ Standard Plate Counting
- Increasing replicates
- Combo of *nhaPx* antiporters
- Studying other antiporters in Vc to see if one has a larger effect or whether it's a combo of all antiporters that allow for survival

Acknowledgements

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- Funding for this project was supported by an NIH grant (1R21AI109435-01A).
- Additionally, special thanks to Wanda Crannell, Dr. Kate Field and other students within the Bioresource Research program at Oregon State University.



Questions?