# The Effects Of CRISPR-Cas Systems On The Rate Of Horizontal Gene Transfer

A Network Theoretic Approach



#### Siddharth Reed Undergraduate Thesis

Golding Lab, Biology Department, McMaster University

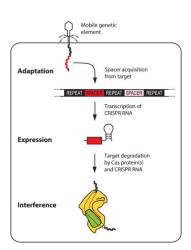
January 22, 2019

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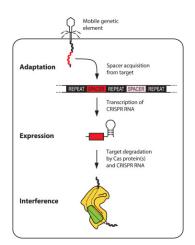
- 1. CRISPR-Cas systems
- 2. Horizontal Gene Transfer
- 3. Phylogenomic Networks
- 4. Do CRRISPR Systems Affect Horizontal Gene Transfer?
- 5. My Project

# CRISPR-Cas systems

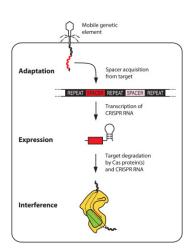
 Adaptive Bacterial Immune System



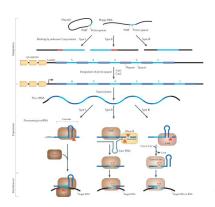
- Adaptive Bacterial Immune System
- Protects against foreign DNA



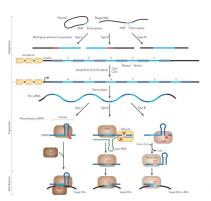
- Adaptive Bacterial Immune System
- Protects against foreign DNA
- Requires Cas proteins and CRISPR loci



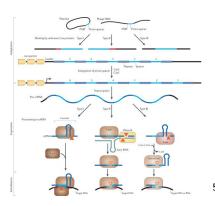
• 45% of bacteria have CRISPR loci  $(n = 6782)^2$ 



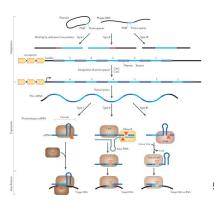
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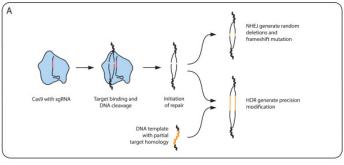


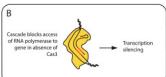
- 45% of bacteria have CRISPR loci  $(n = 6782)^2$
- 3 Main Types, multiple subtypes<sup>3</sup>
- CRISPR arrays represent unique life history of an organism
- 11% 28% are false or orphaned CRISPR loci<sup>4</sup>

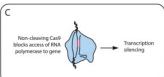


## Biotech Application

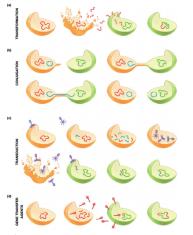
## **Biotech Application**





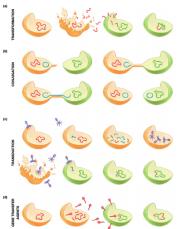


## Horizontal Gene Transfer

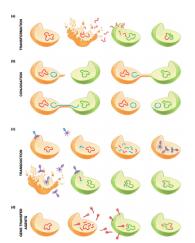


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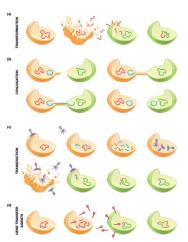


 Conjugation: Transfer of DNA through cell-cell connections<sup>6</sup>



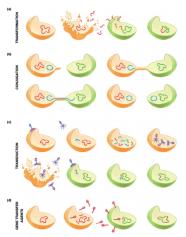
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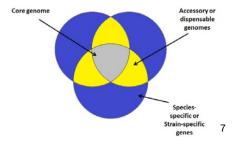
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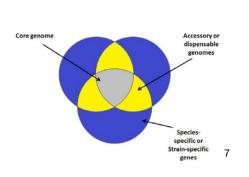
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- CRISPR-Cas directly affects Transduction and Transformation<sup>6</sup>

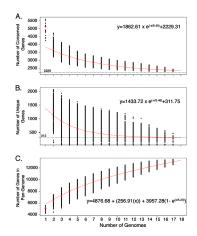
## Pan-Genomes

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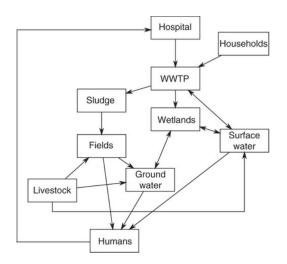
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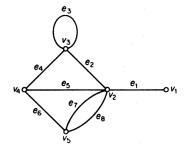
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- Sequence compatibility

## Applications

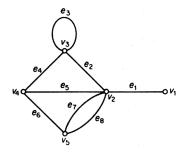
## **Applications**



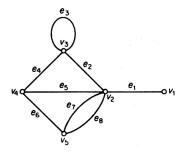
## Phylogenomic Networks



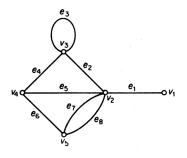
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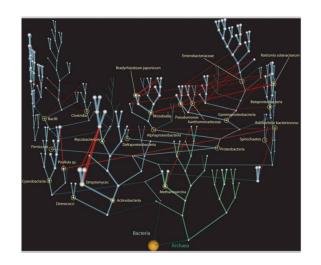
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- Directed or Undirected Edges



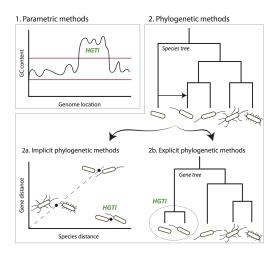
- Useful mathematical abstraction of real world system
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- Directed or Undirected Edges
- Weighted or Unweighted Edges

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### Construction



# Do CRRISPR Systems Affect Horizontal Gene Transfer?

# Yes

• Cost tradeoff factors:

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  - Prophage abundance<sup>15</sup>

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- CRISPR can enhance transduction-mediated HGT<sup>15</sup>

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  - Can see inhibitory effects of CRIPSR on HGT over short evolutionary time scales
  - Higher gene indel rates for CRISPR containing genera than non-CRISPR containing outgroups

# My Project

### **Hypothesis**

### **Null Hypothesis**

Bacterial strains or genera with known CRISPR systems will show no significant differences in network statistics compared to those strains or genera without known CRISPR systems.

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### Alternative Hypothesis

Bacterial strains or genera with known CRISPR systems will show a significant difference in at least 1 network statistic compared to those strains or genera without known CRISPR systems.

### Within Network Comparisons

For genera with CRISPR conatining strains, compare the node statistics of CRIPSR-containing strain to non-CRISPR-containing strains.

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### Between Network Comparisons

For genera with no CRISPR conatining strains, compare the network statistics of mixed to non-CRISPR-containing networks.

### Within Network Comparisons

For genera with CRISPR conatining strains, compare the node statistics of CRIPSR-containing strain to non-CRISPR-containing strains.

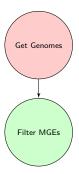
### Between Network Comparisons

For genera with no CRISPR conatining strains, compare the network statistics of mixed to non-CRISPR-containing networks.

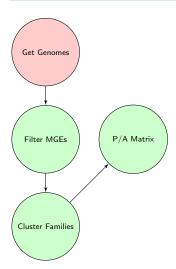
#### Gene Indel Rates vs. Network Statistics

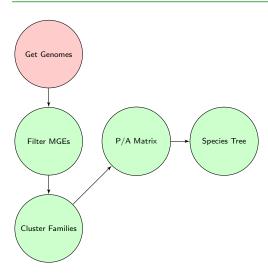
Compare gene InDel rates to node/network statistics for CRISPR-containing and non-CRISPR-containing strains/genera.

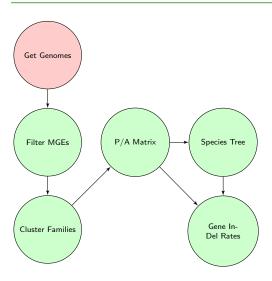


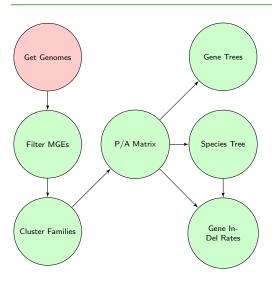


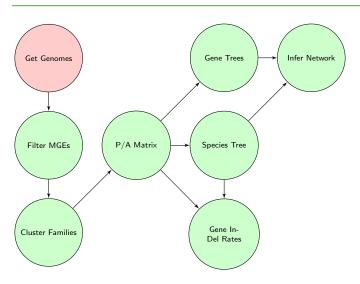


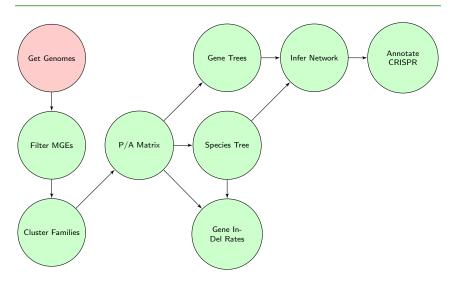


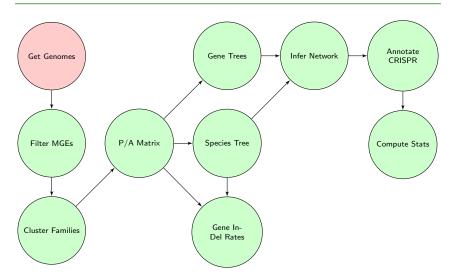


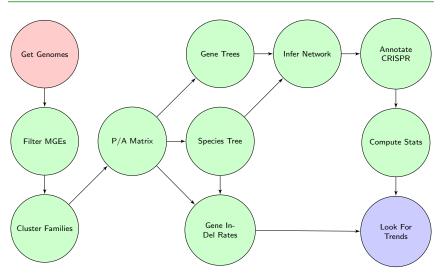












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• Total/Average edge weight:  $\sum_i w_i, \frac{\sum_i w_i}{N}$ 

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- Node Associativity:  $\frac{j(j+1)(\overline{k}-\mu_q)}{2E\sigma_q^2}$  where j is the excess degree of the node and  $\overline{k}$  is the average excess degree of the node's neighbors and  $\mu_q$  and  $\sigma_q$  are the mean and standard variation of the excess degree distribution.

#### **Thanks**

#### Thank you to

- Dr. G. Brian Golding
- The Golding lab
  - Caitlin Simopoulos
  - o Daniella Lato
  - Zachery Dickson
  - Sam Long
  - Lucy Zhang
  - Brianne Laverty
  - Nicole Zhang
- · Everyone here for listening

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