## How Quorum Sensing Interactions Affect Population Structure 02-712 Final Project

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## Background

## **Quorum-Sensing Systems**

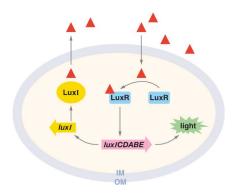


Figure 1: Waters and Bassler (2005)

 Signal-Receptor molecule pairs that modulate gene expression

## **Quorum-Sensing Systems**

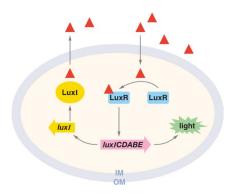


Figure 1: Waters and Bassler (2005)

- Signal-Receptor molecule pairs that modulate gene expression
- Once threshold density is reached, enough signal is received to upregulated target genes

## **Quorum-Sensing Systems**

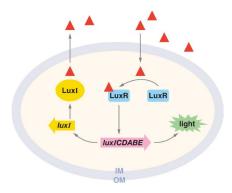


Figure 1: Waters and Bassler (2005)

- Signal-Receptor molecule pairs that modulate gene expression
- Once threshold density is reached, enough signal is received to upregulated target genes
- Can lead to biofilms, antibiotic production etc.

### Public Goods and Cheating

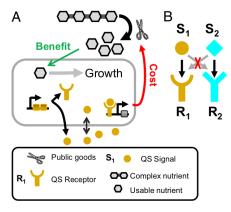


Figure 2: Eldar (2011)

When quorum is reached, bacteria produce a "public good"

### Public Goods and Cheating

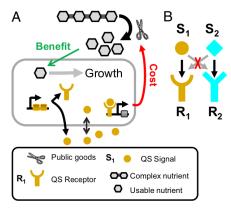


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- Everyone benefits from this even if they don't contribute

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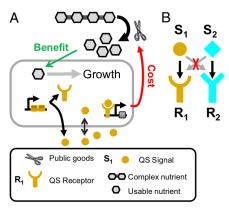


Figure 2: Eldar (2011)

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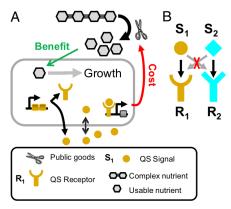


Figure 2: Eldar (2011)

- When quorum is reached, bacteria produce a "public good"
- Everyone benefits from this even if they don't contribute
- Must produce the receptor, signal molecule and good to contribute
- Cheaters DO prosper (if you are a bacterium)

#### Who Cares?

▶ check the discussion from Eldar (2011) for references

Maintaining Freeloaders as a Diversity Reservoir

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Kin Recognition for Strains

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Kin Recognition for Strains

Designing Cheaters to Disrupt Pathogen Growth

#### Methods

#### Basic ODE Model

## Social conflict drives the evolutionary divergence of quorum sensing

Avigdor Eldar<sup>1</sup>

Department of Molecular Microbiology and Biotechnology, Faculty of Life Sciences, Tel Aviv University, Tel Aviv 69978, Israel

▶ Equations taken from Eldar (2011)

#### Basic ODE Model

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- Equations taken from Eldar (2011)
- Deterministic ODE model of bacteria
- ▶ We focused on QS interactions over Evolution

## Signal-Receptor Activation Matrix $K_{ac}$

Represents all receptors-signal pairs  $(R_iS_i)$  present in at least 1 OTU in the population

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## Signal-Receptor Activation Matrix $K_{ac}$

- ▶ Represents all receptors-signal pairs  $(R_iS_i)$  present in at least 1 OTU in the population
- ▶ Different sets of receptor-signal combinations can produce the same  $K_{ac}$
- $K_{ac}$  is of dimension  $|R| \times |S| = |N| \times |N|$

#### Facultative Cheaters

Matrix for 2 strains  $R_1S_1$  and  $R_2S_2$ 

$$\begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix}$$

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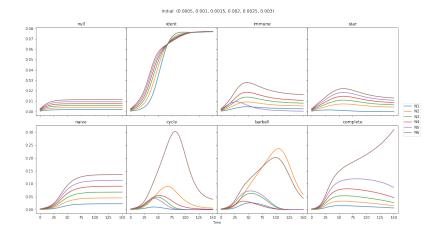
#### **Custom Matrix**

Matrix for 2 strains  $R_1R_2S_1$  and  $R_2S_2$ or 3 strains  $R_1S_1$ ,  $R_2S_1$  and  $R_2S_2$ 

$$egin{bmatrix} 1 & 0 \ 1 & 1 \end{bmatrix}$$

#### Results

## Comparing Different $K_{ac}$ Matrices



## How $K_{ac}$ Sparsity Affects Population Structure

## Simulating With Human Gut Microbiome Data

#### Discussion

Chaining  $K_{ac}$  has strong effects on model dynamics

▶ changes can occur independent of sparsity

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#### Chaining $K_{ac}$ has strong effects on model dynamics

- changes can occur independent of sparsity
- decreasing sparsity does . . .

## Moral of the Study

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# Cheating may help but cooperating is best!

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# Cheating may help but cooperating is best!

(for bacteria)

## Bibliography I

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