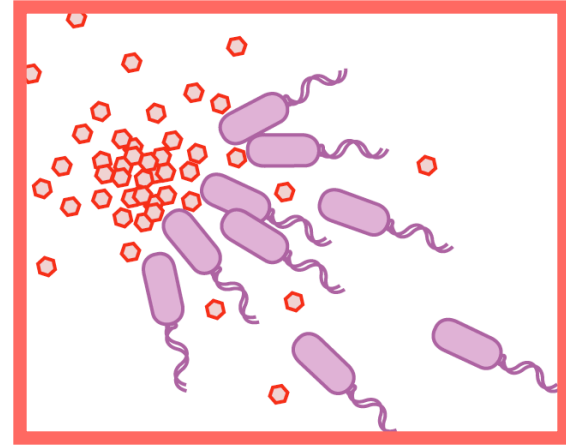
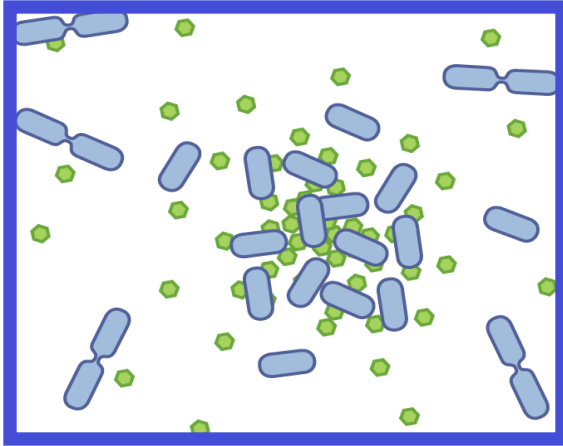




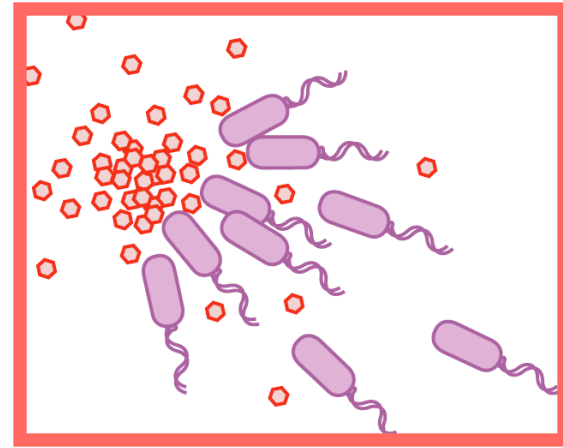
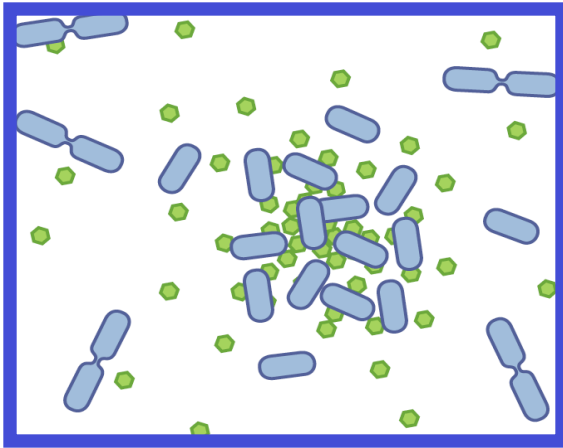
Quorumtaxis:

Programming *E. coli* to Eavesdrop,
Stalk, and Kill *B. subtilis*

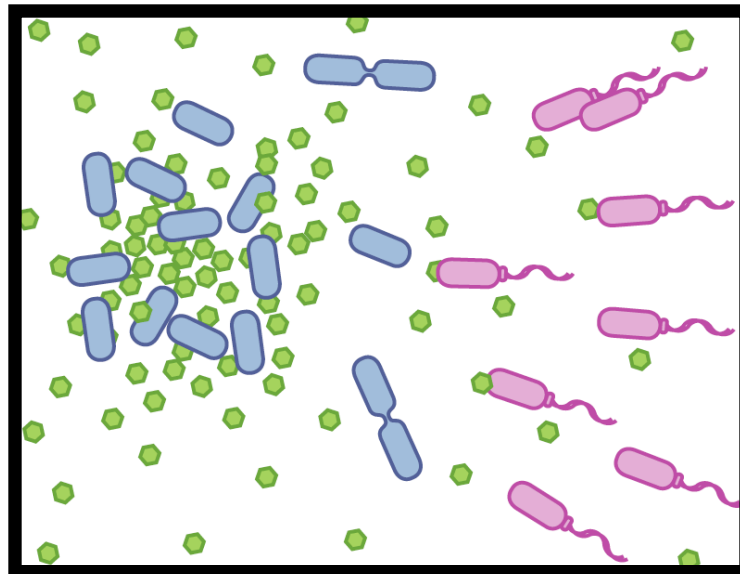
Quorum-sensing + Chemotaxis



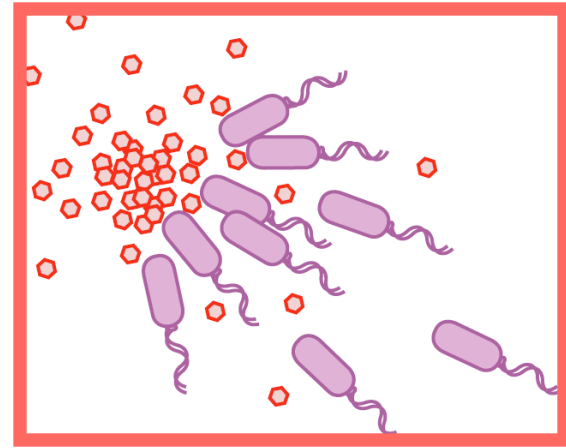
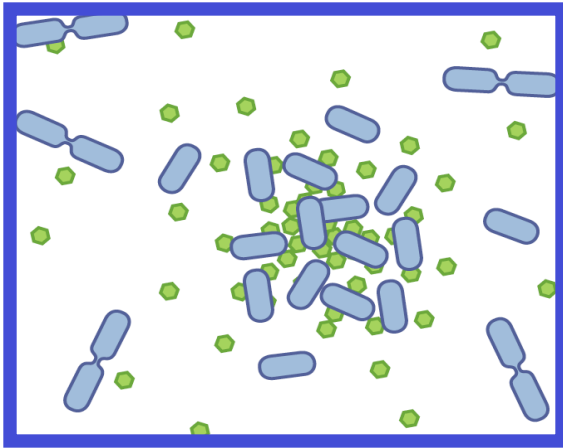
Quorum-sensing + Chemotaxis



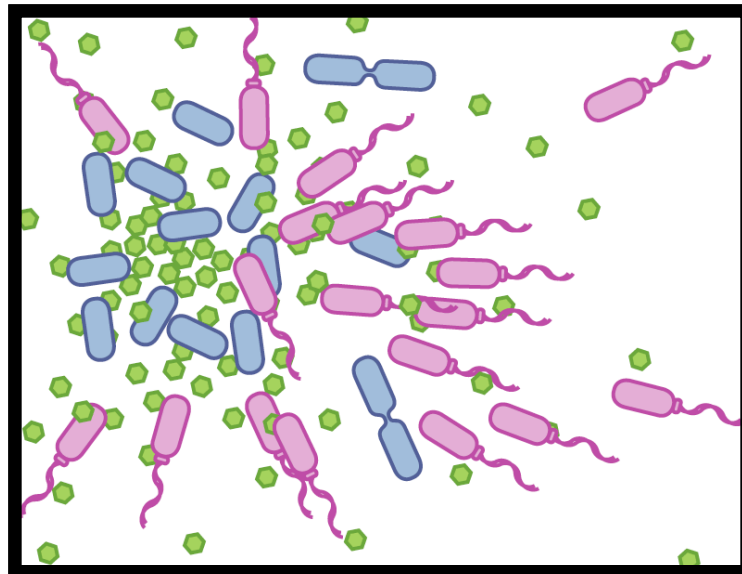
= Quorumtaxis



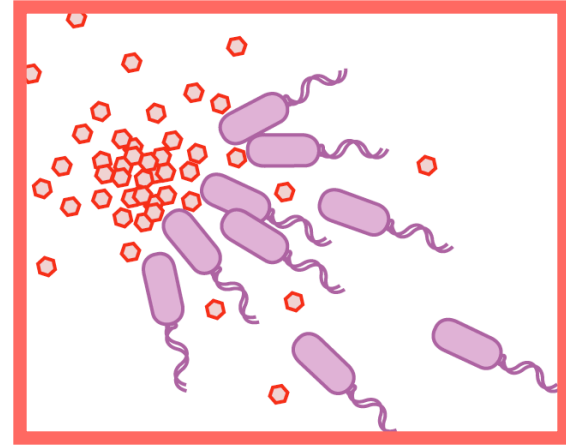
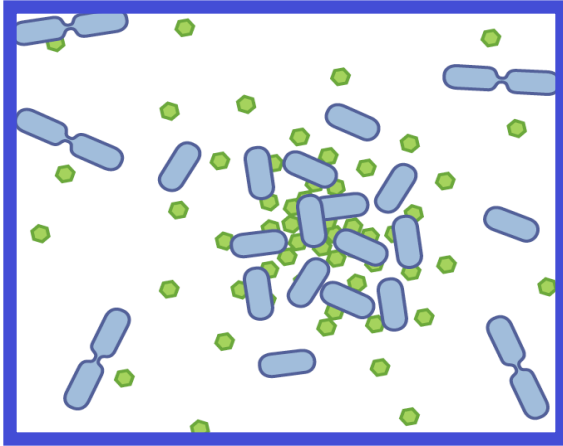
Quorum-sensing + Chemotaxis



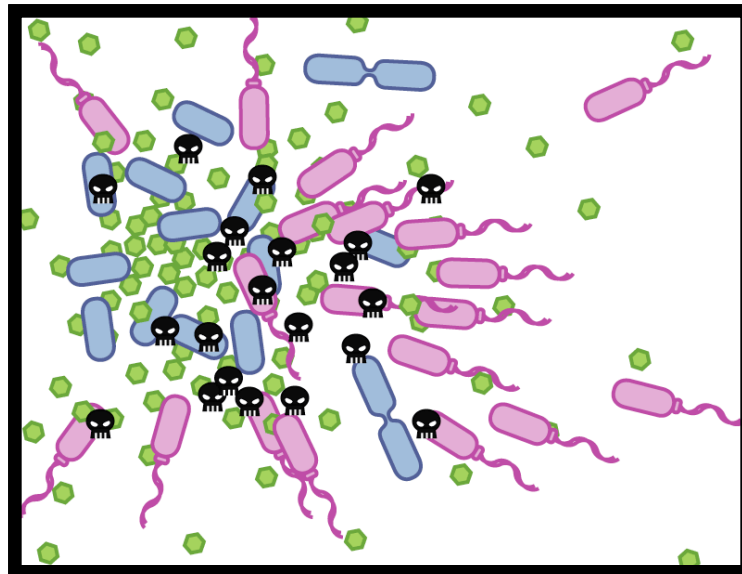
= Quorumtaxis



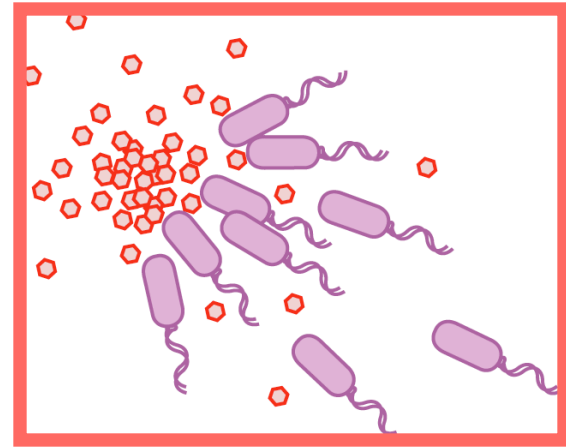
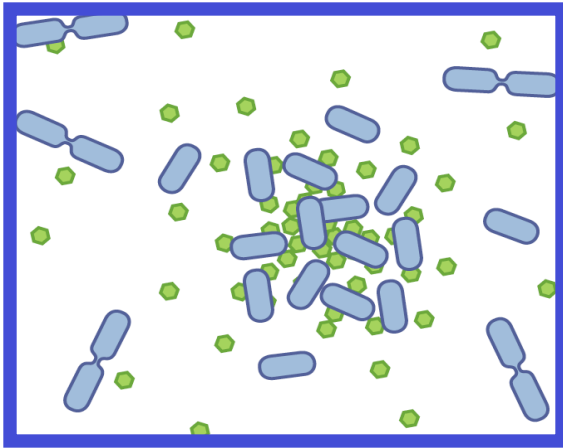
Quorum-sensing + Chemotaxis



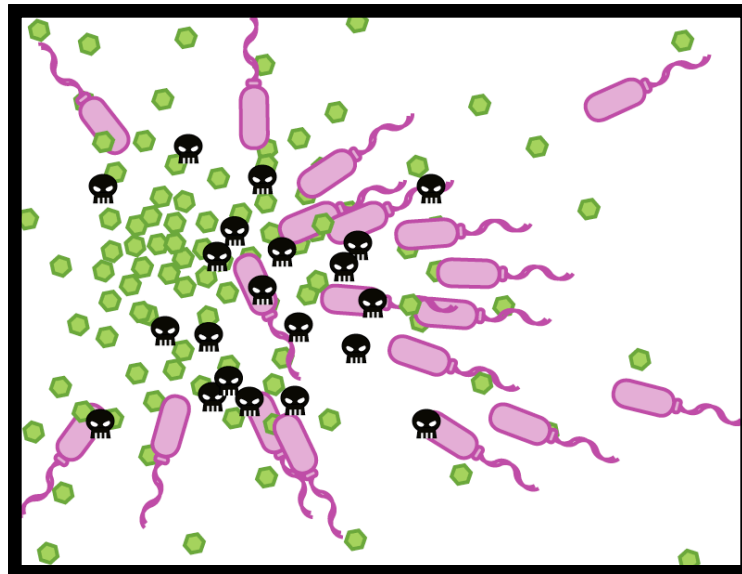
= Quorumtaxis



Quorum-sensing + Chemotaxis



= Quorumtaxis

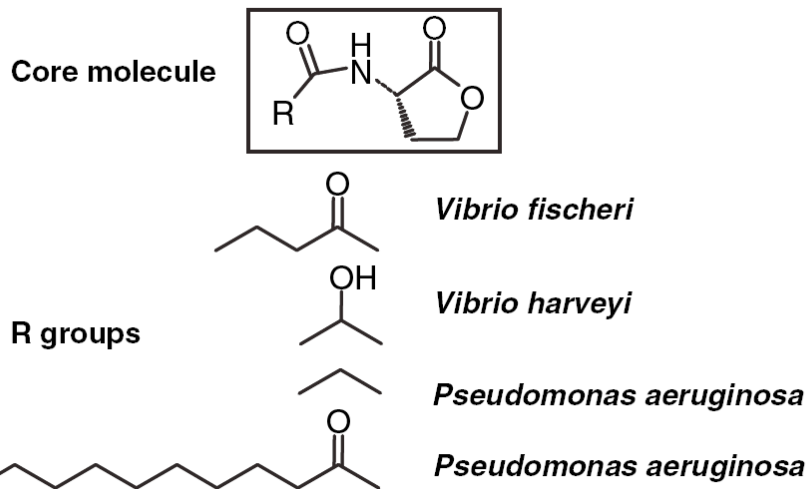


What is a Good Pheromone to Use for Quorumtaxis?

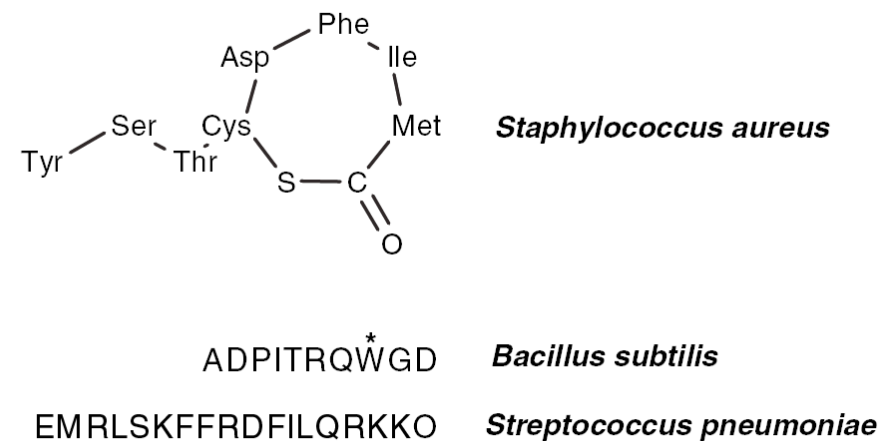
Gram-negative bacteria

Gram-positive bacteria

Acyl homoserine lactone autoinducers



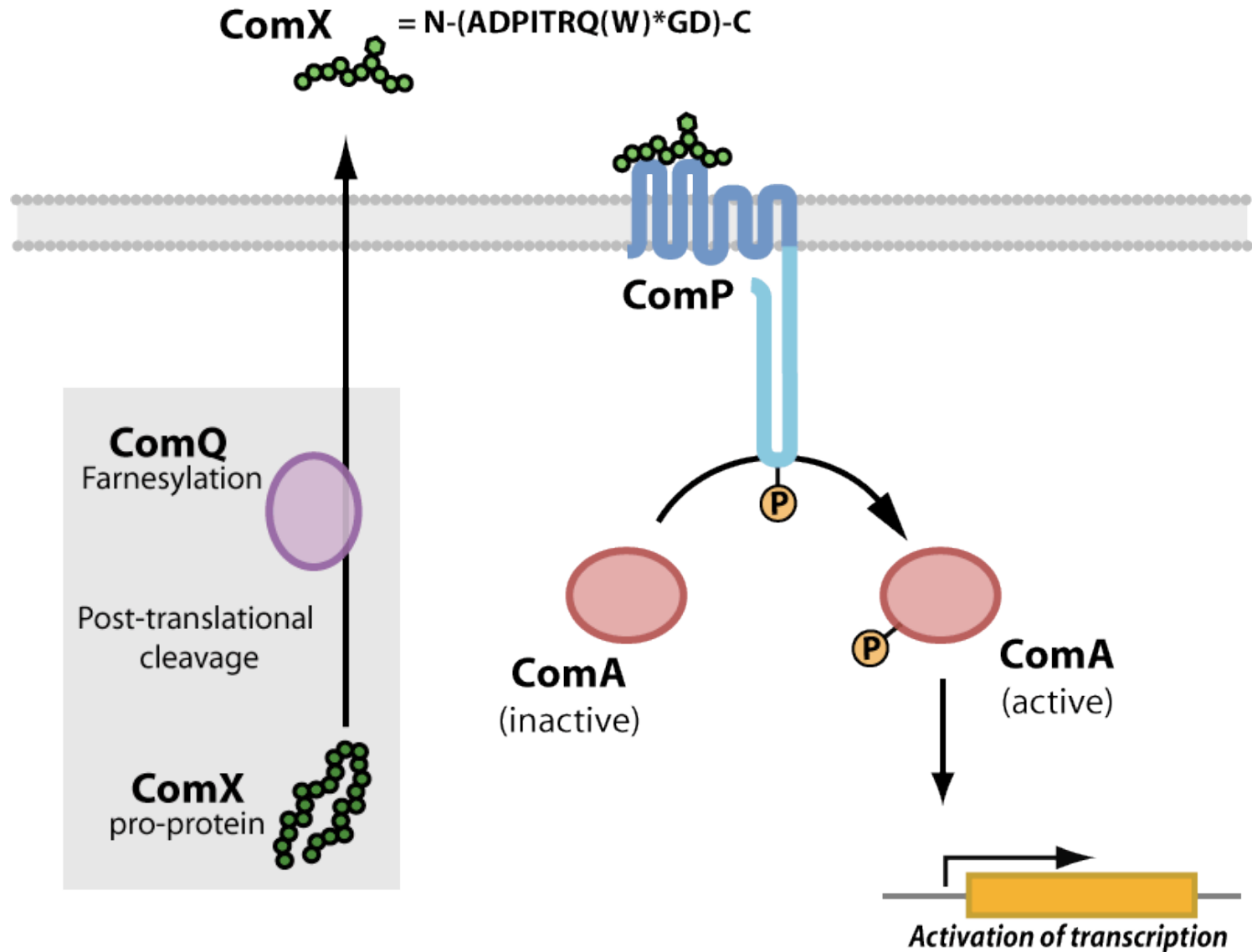
Oligopeptide autoinducers



AHL-based pheromones are metabolic products with limited diversity.

Oligopeptides are genetically encoded with a tremendous potential for diversity.

The *B. subtilis* Peptide-based Quorum Sensing System has 4 Protein Components



The ComXP Systems Could be Used to Create Diverse Quorumtaxis Systems

ComX precursors exhibit tremendous sequence diversity,

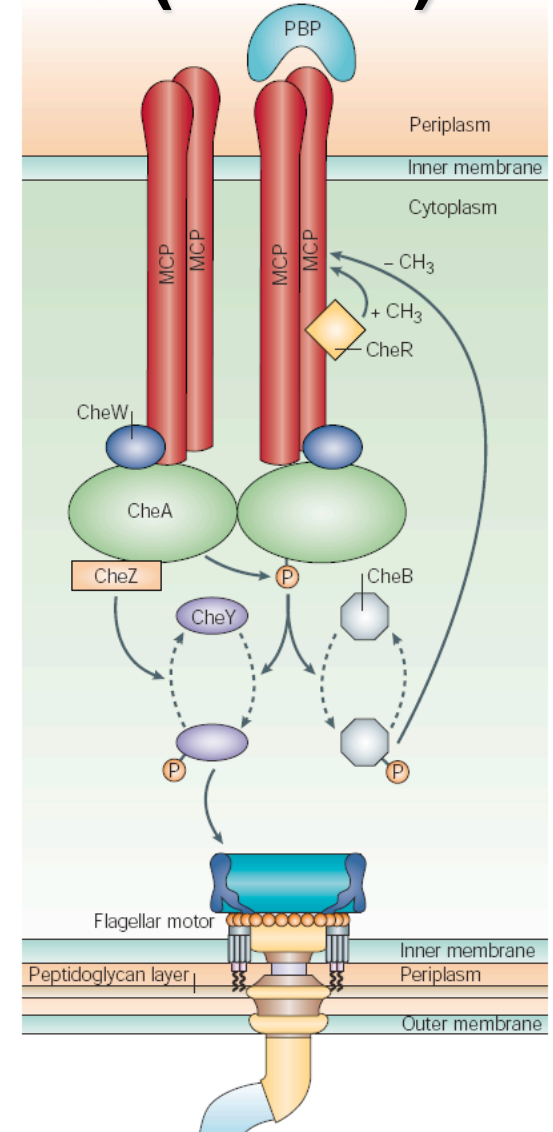
<u><i>Bs</i> strain</u>	<u>ComX precursor (C-terminus)</u>	
RO-E-2DKLIPSIVDIFNKKMTLSKKCKGIF W EQ	58
RO-H-1QSECIINGFK---GLEIYSMLD W KY	53
RO-B-2QLKSIVDAFG---GLQIYTNGN W VPS	53
RO-OO-2QLKSIVDAFR---GMQIYTNGN W VPS	53
RO-C-2EKKAILYAFEQ---GEVPRTSK W PPI	53
RO-FF-1ETQLIIEGFEGIEEVKRGNAGK W GP	53

...and orthogonal ComX systems have evolved that do not communicate, eliminating cross-talk between species and reducing background.

ComP Receptor is Related to Methyl-Chemotaxis Proteins (MCPs)

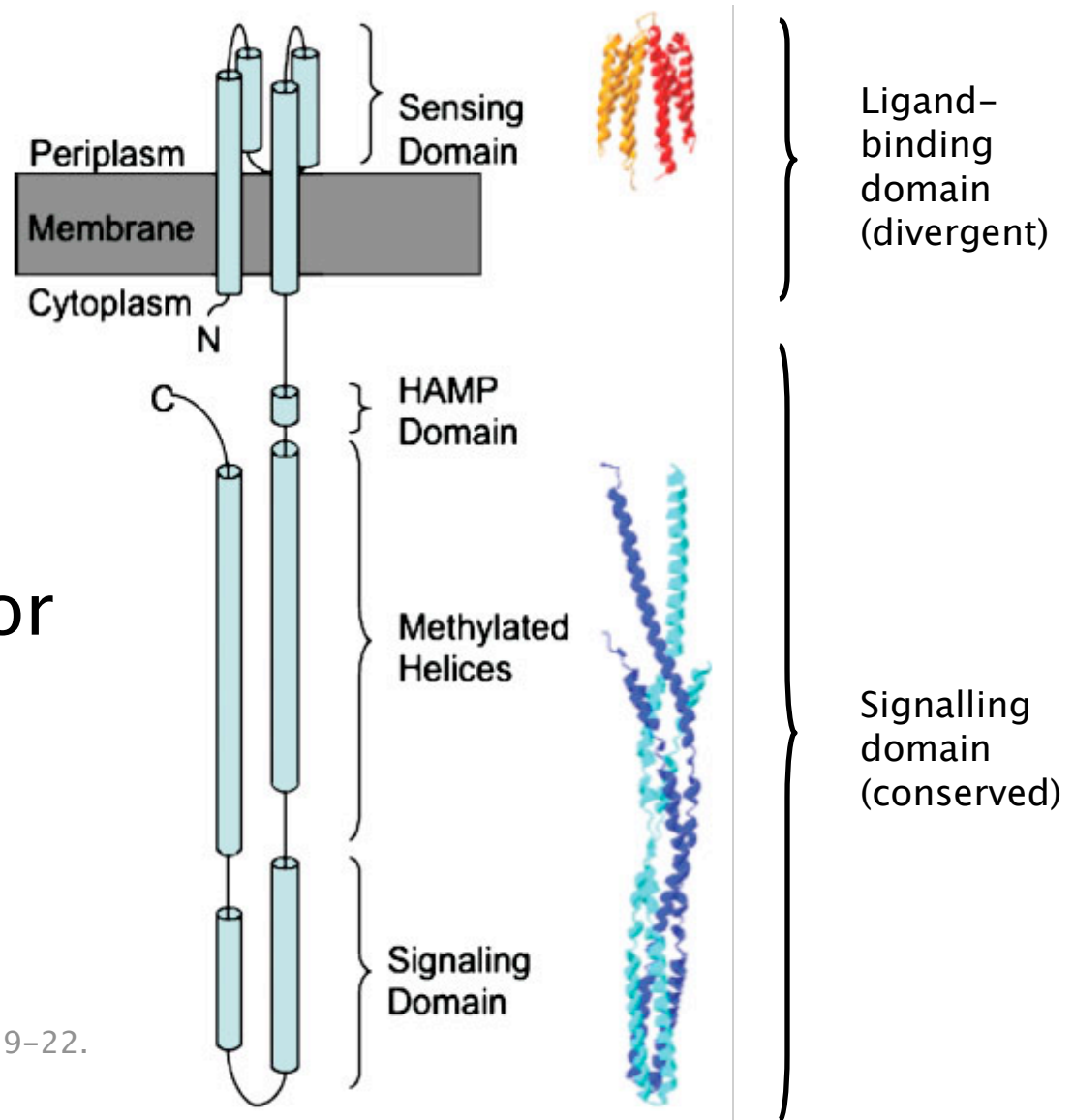
Gene	Protein	Response
tsr	Tsr	serine
tar	Tar	aspartate
trg	Trg	ribose
tap	Tap	peptides
aer	Aer	oxygen

§ ComP and Tsr exhibit ~11% identity.

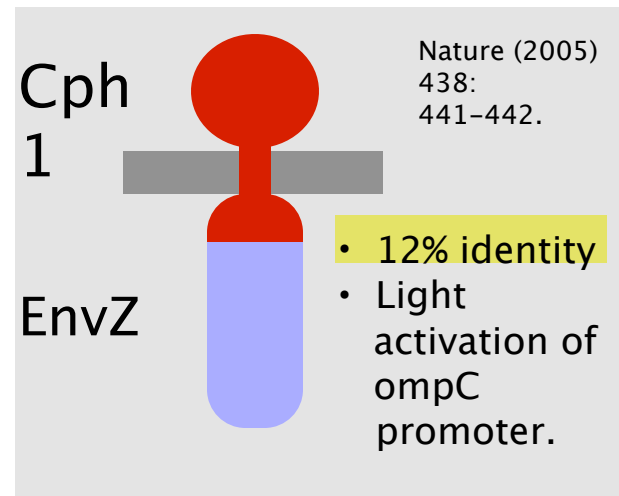
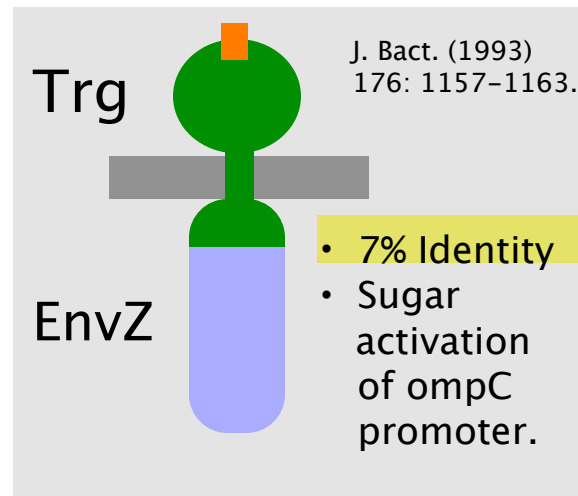
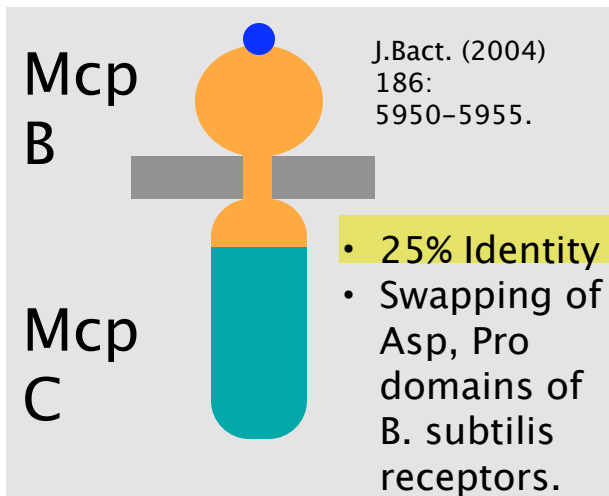
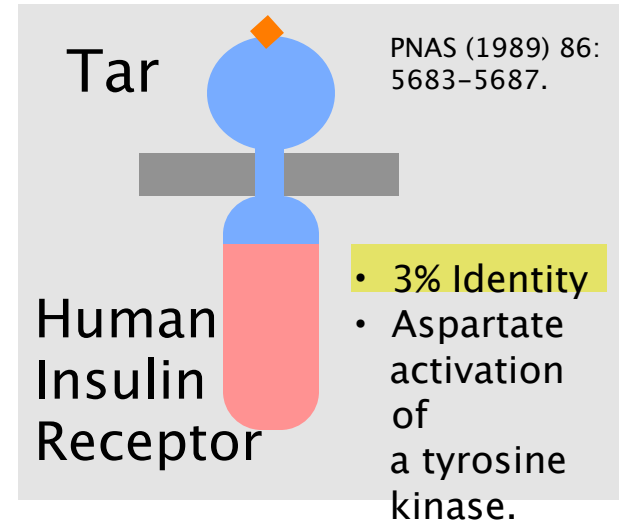
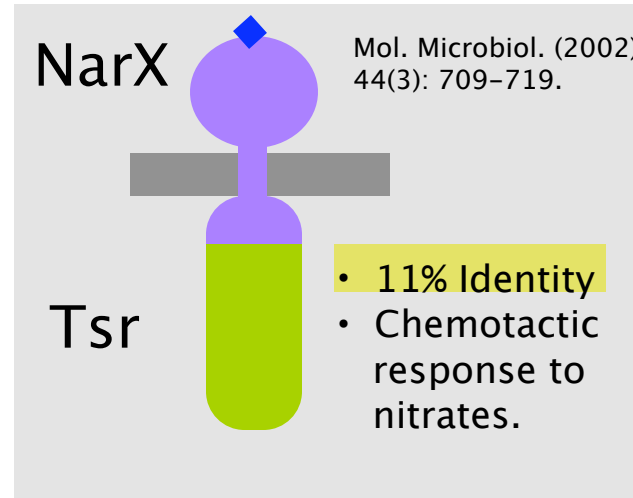
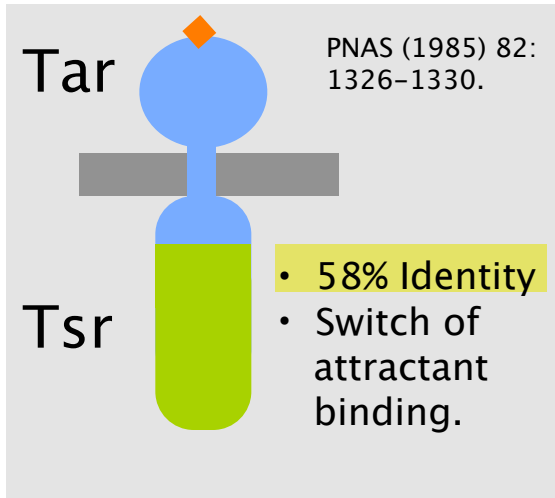


MCPs Have a Modular Architecture

Tsr Receptor



MCP-like Proteins Have Been Recombined to Create Novel Receptors



Strategy

Phase I

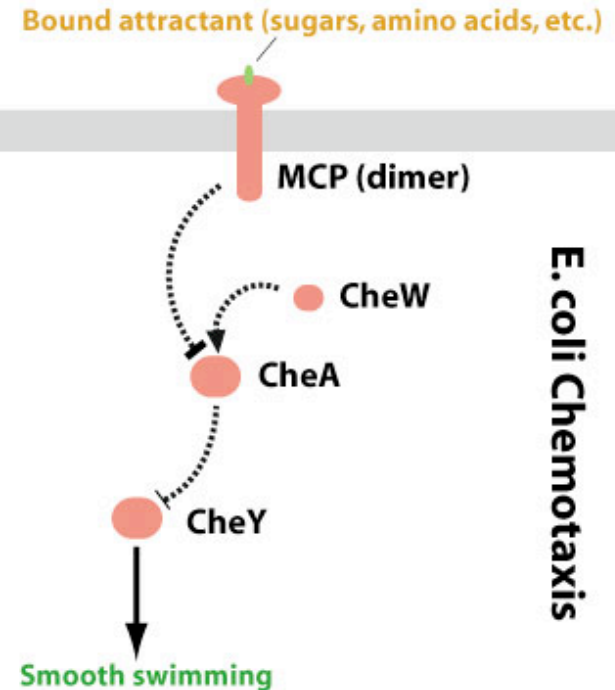
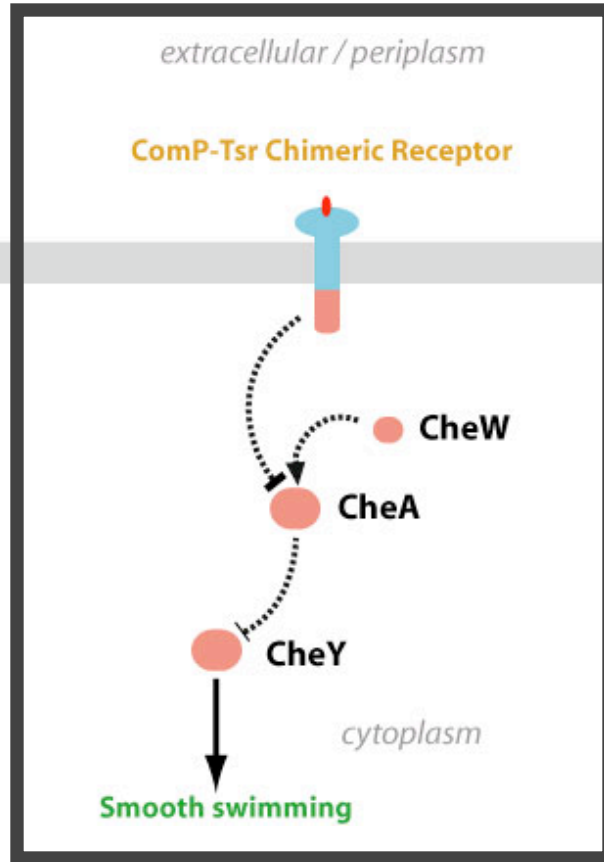
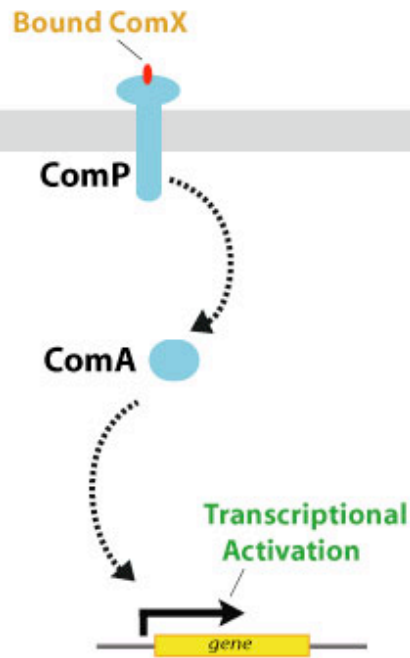
- § Use BioBricks and ComP–Tsr chimeras to construct quorumtaxis circuit.
- § Use chemotaxis models to design and optimize quorumtaxis circuit.

Phase II

- § Extend circuit to include destroy output, e.g., SdpC toxin production.

Module Integration by a Chimeric Receptor

Bacillus Quorum Sensing

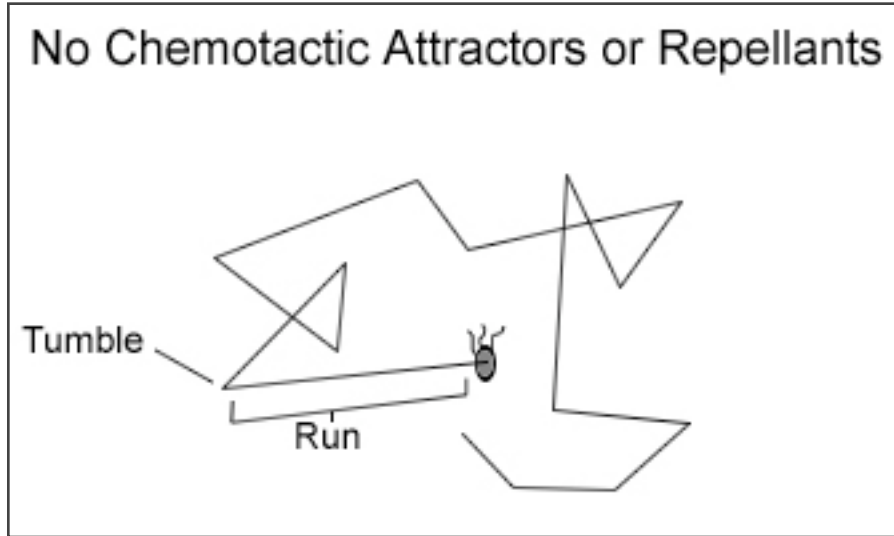


Issues That May Arise With Chimeric Receptors

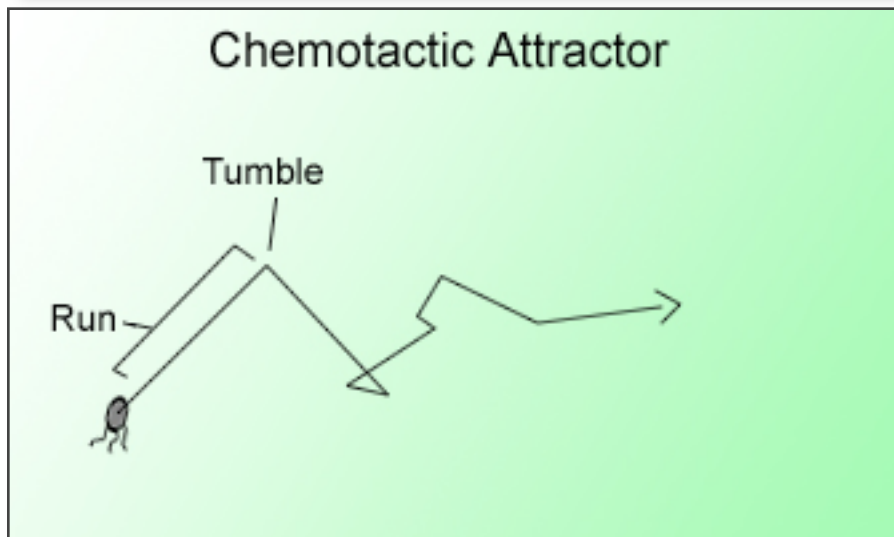
§ ComP–Tsr chimeras may be weakly functional because they:

1. Weakly bind ComX.
2. Bind ComX, but transmit the signal poorly to the cytoplasmic domain.
3. Bind ComX, transmit the initial signal to the flagella, but are unable to adapt to gradients of ComX.

Bacterial Chemotaxis Works by Modulating Run Lengths

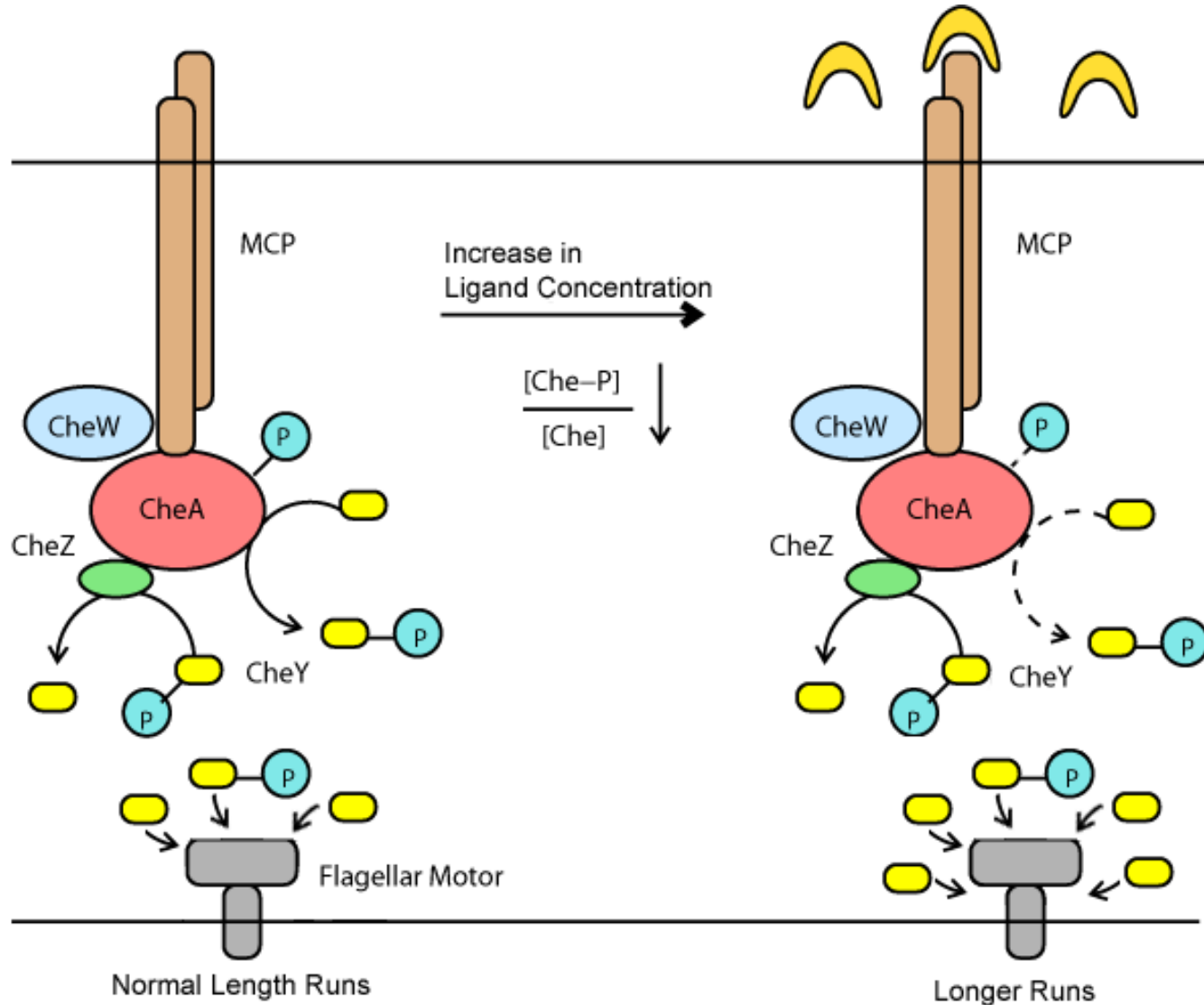


§ Runs with an average length of 1 second between tumbles.

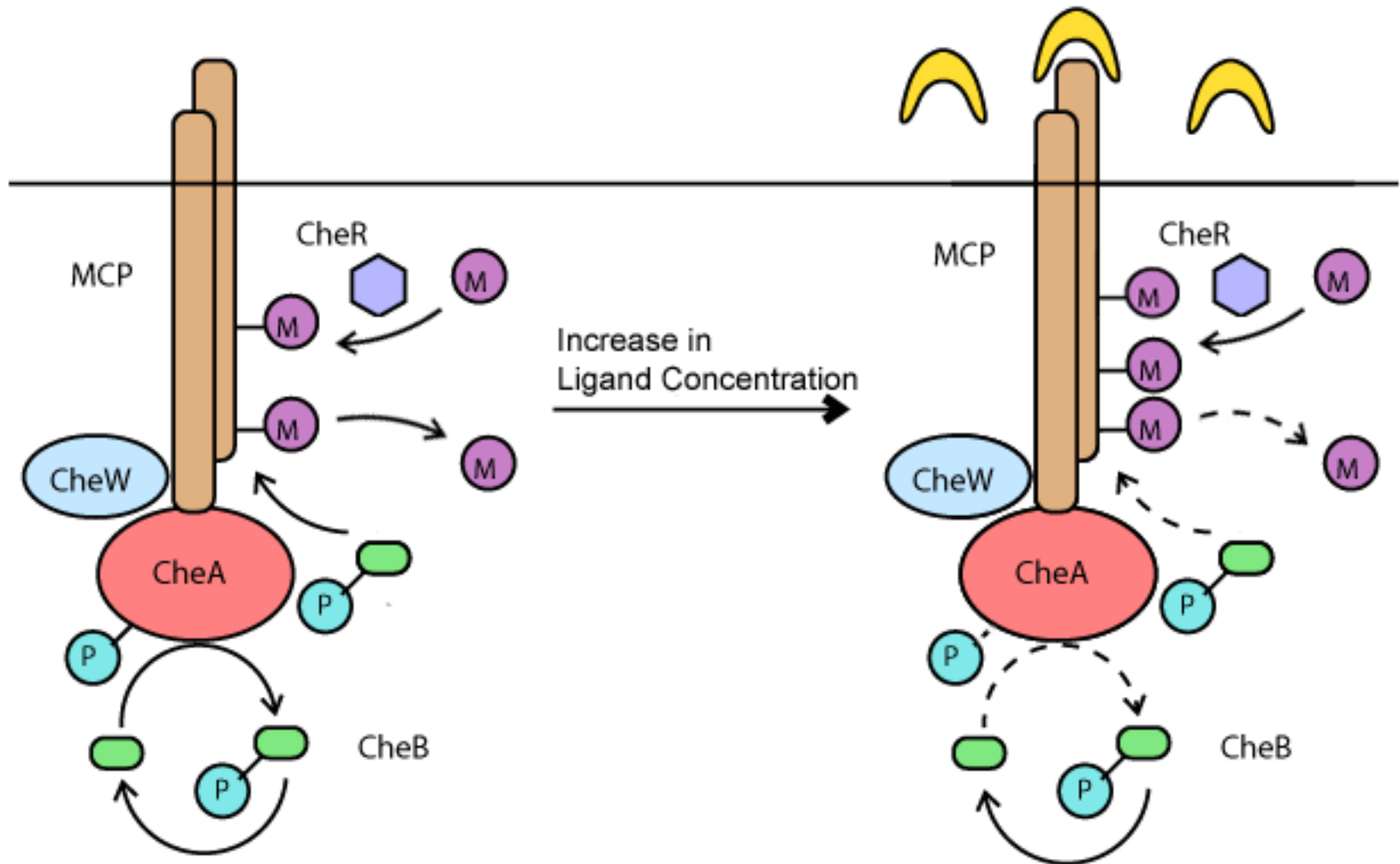


§ Run lengths are increased when moving up the gradient, and decreased when moving down.

Flagellar Motion is Controlled by Fast Reactions Involving CheY

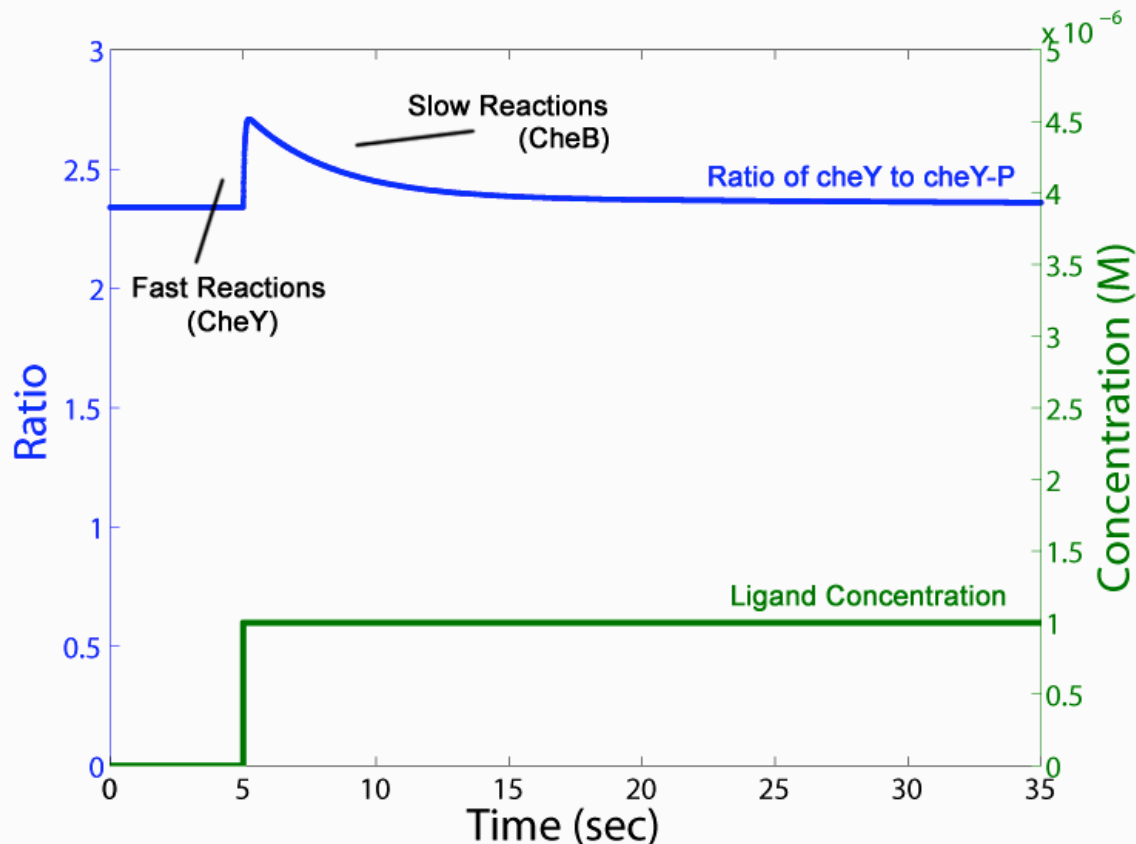


Adaptation is Facilitated by a Relatively Slow Chain of Reactions Involving CheB



Chemotaxis Can be Modeled by Combining the CheY and CheB Reaction Sequences

- § Our model uses 17 differential equations and takes into account over 20 reactions.
- § Reactions are modeled using the Law of Mass Action and Michaelis–Menten enzyme kinetics.



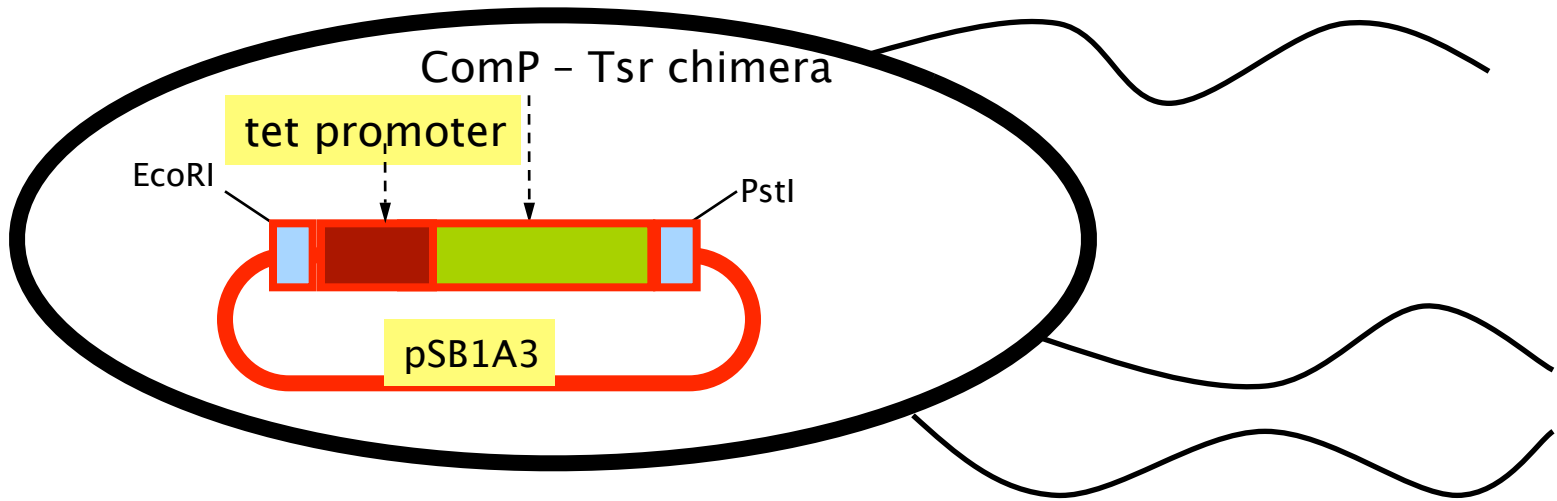
We Will Use the Model to...

- § Identify ways in which our chimeric receptors differ from natural MCPs based on experimental data.
- § Determine which of the working chimeras best suits our project goals.
- § Characterize the specifics of our chimeras in order to make them easier to understand and use.

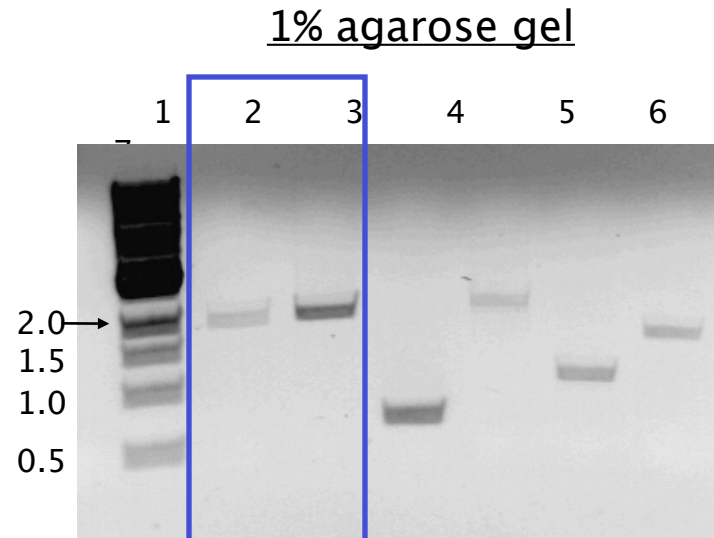
Outline of Experimental Strategy

1. Amplify genetic components.
2. Create ComP/Tsr chimera library and build quorumtaxis circuit.
3. Screen for functional quorumtaxis circuit.

Step 1: Amplify BioBricks

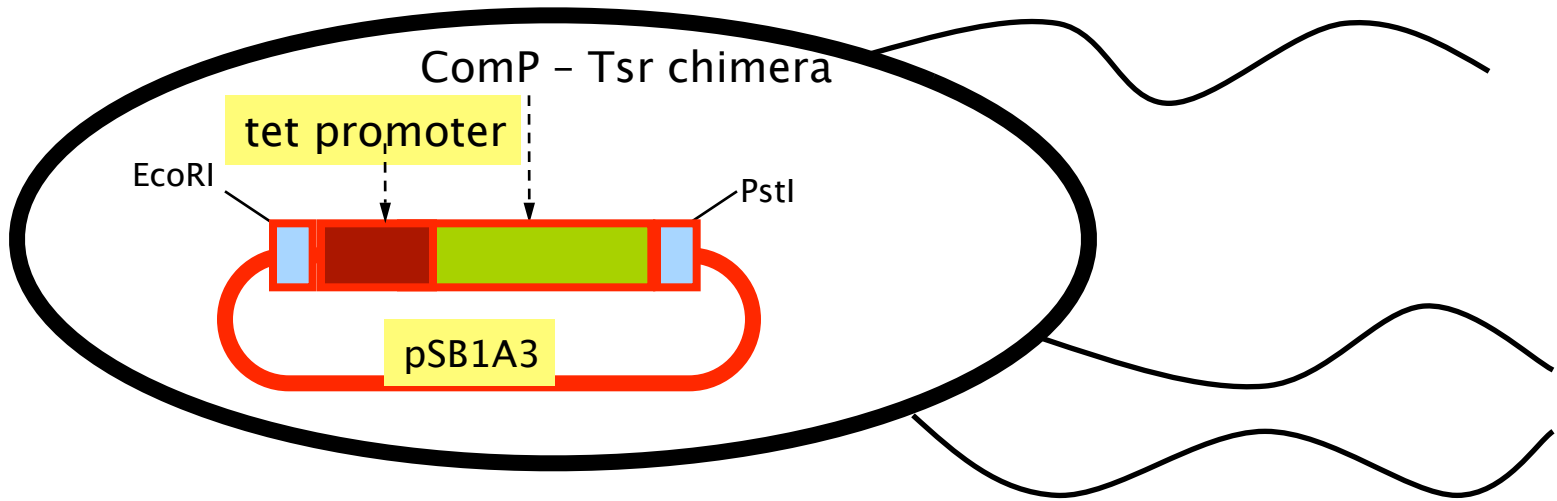


<u>Lane</u>	<u>Sample</u>	<u>Size</u>
1	1 kb std	--
2	pSB1A3 plasmid	2157 bp
3	pTetRBS + pSB1A3	2231 bp
4	ComA	650 bp
5	ComP	2400
6	ComQX	1100 bp
7	Tsr	1600 bp



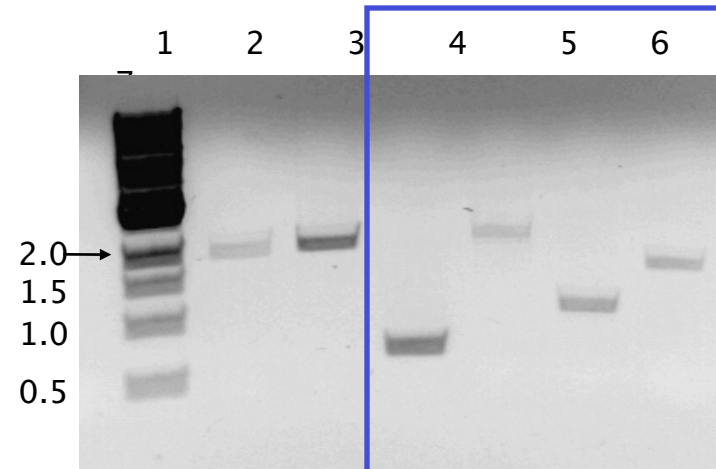
Step 2: Create Com and Tsr BioBricks

(from *B. subtilis* 168 and *E. coli*)

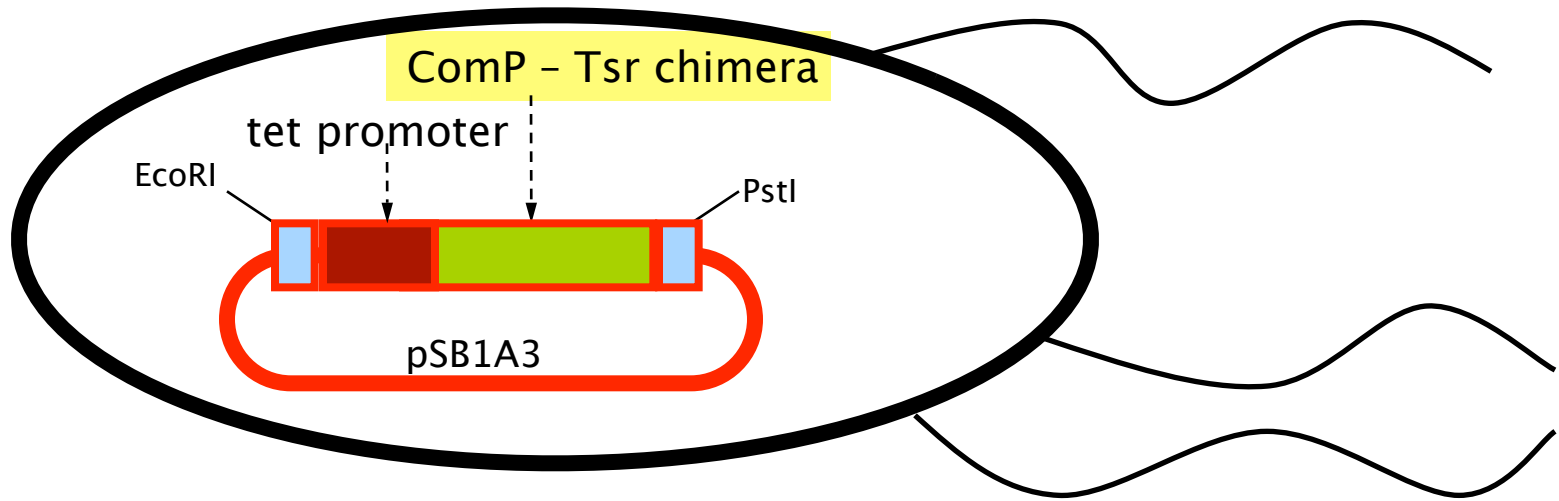


1% agarose gel

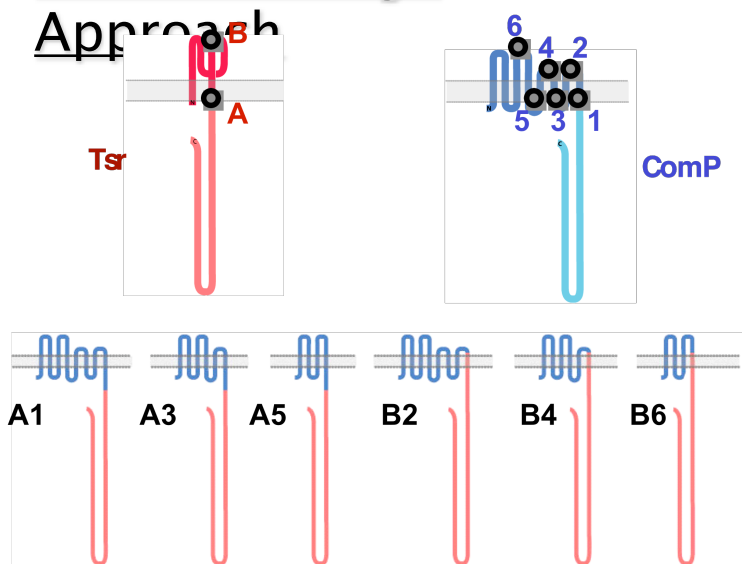
<u>Lane</u>	<u>Sample</u>	<u>Size</u>
1	1 kb std	--
2	pSB1A3 plasmid	2157 bp
3	pTetRBS + pSB1A3	2231 bp
4	ComA	650 bp
5	ComP	2400
6	ComQX	1100 bp
7	Tsr	1600 bp



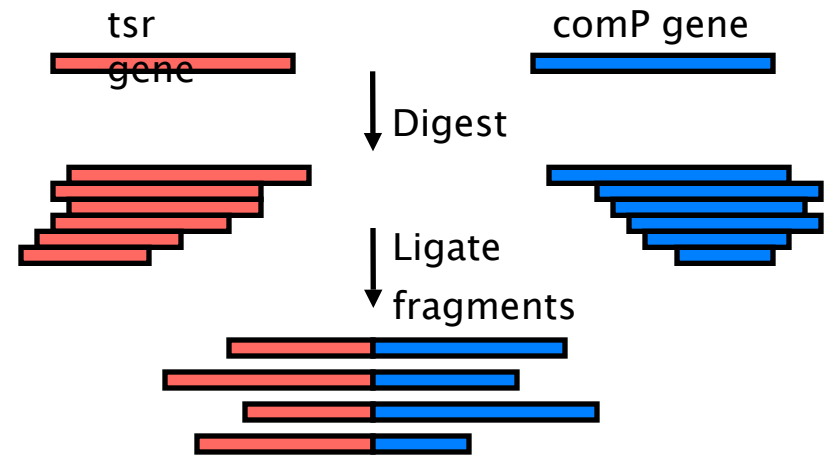
Step 3: Create ComP-Tsr Chimeras



Rational Design Approach



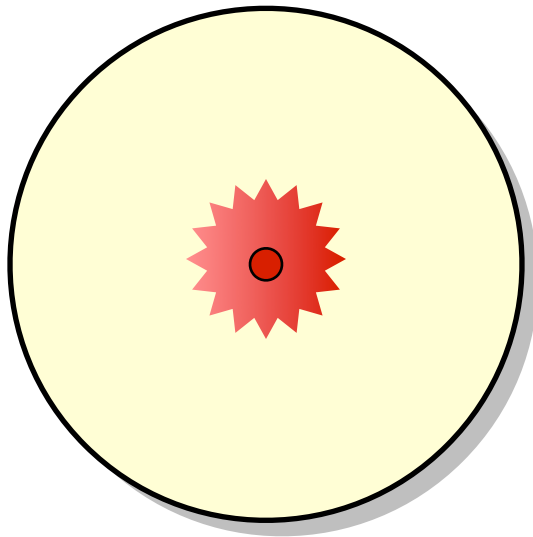
Combinatorial Library



Step 4: Test Δmcp *E. coli* Swarming

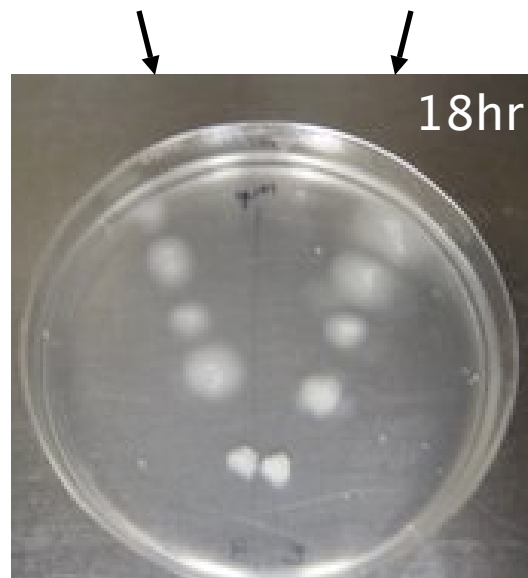
(strain that will be used to screen chimera function)

Swarming assay

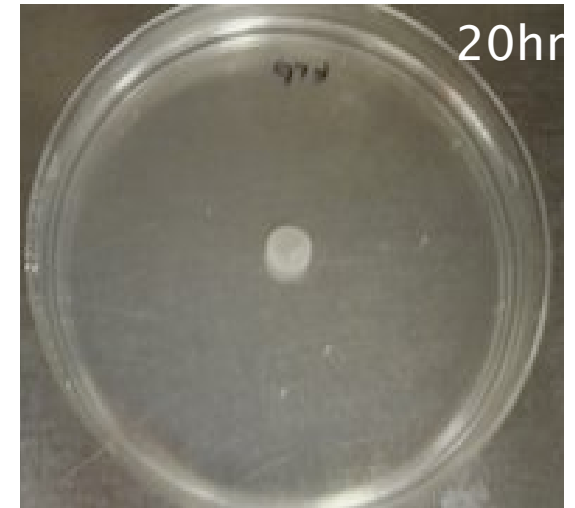


Spot cells on soft agar plate and assay outward growth.

B. subtilis & *E. coli*

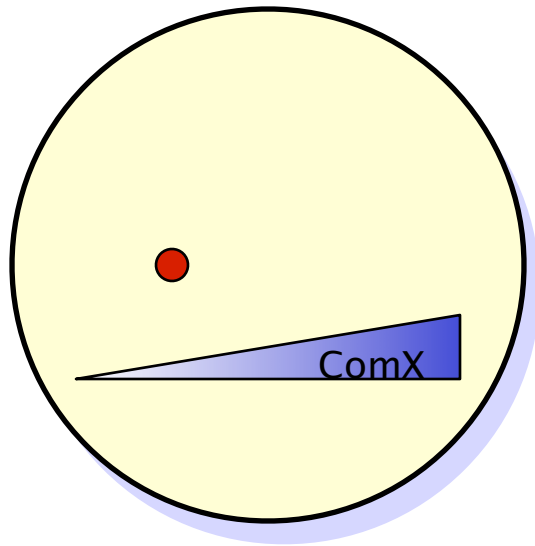


E. coli strain RP8611 (Δmcp)*

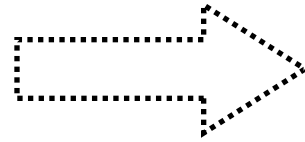


* acquired from Prof. John S. Parkinson, University of Utah

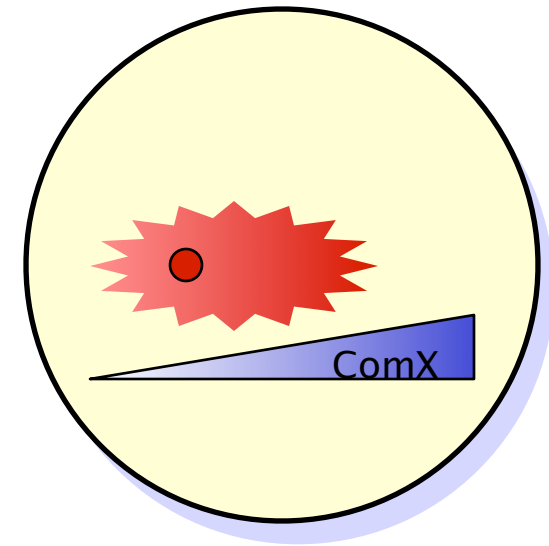
Step 5: Screen for Functional Chimeras



Spot Δmcp *E. coli* on soft agar plate containing *B. subtilis* culture extract (i.e., ComX)



Quorumtaxis



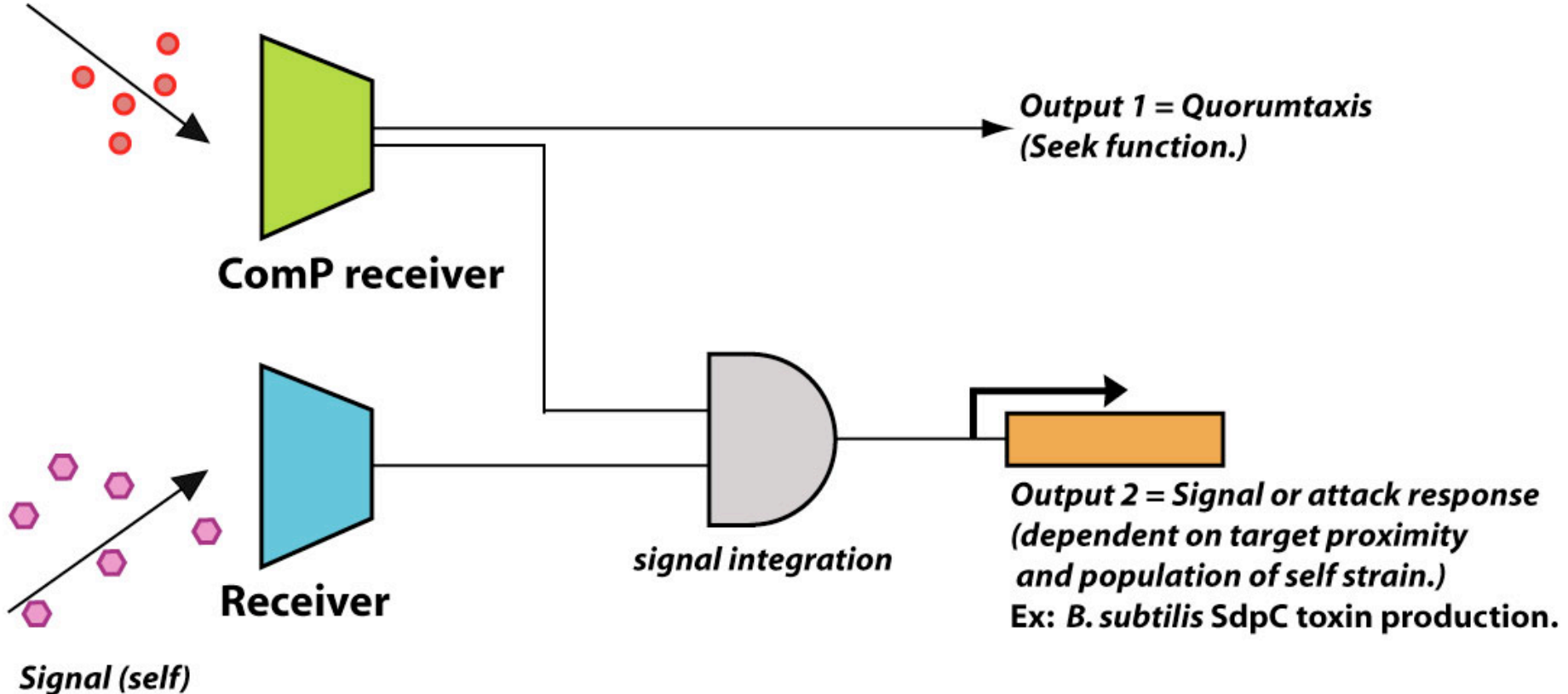
Expected swarming pattern

Progress and Future Work

1. Amplify genetic components.
2. Create ComP/Tsr chimera library and build quorumtaxis circuit.
3. Screen for functional quorumtaxis circuit.

Phase II: Develop 'Destroy' Output

ComX signal (target)



BIOCHEMISTRY AND CELL BIOLOGY

Beth Beason

George Bennett

Tina Chen

Chris Conner

Shan Gao

Leah McKay

Teresa Monkkonen

Bibhash Mukhopadhyay (BCM)

Peter Nguyen

Joff Silberg

Mary Kay Thompson

Jeremy Thompson

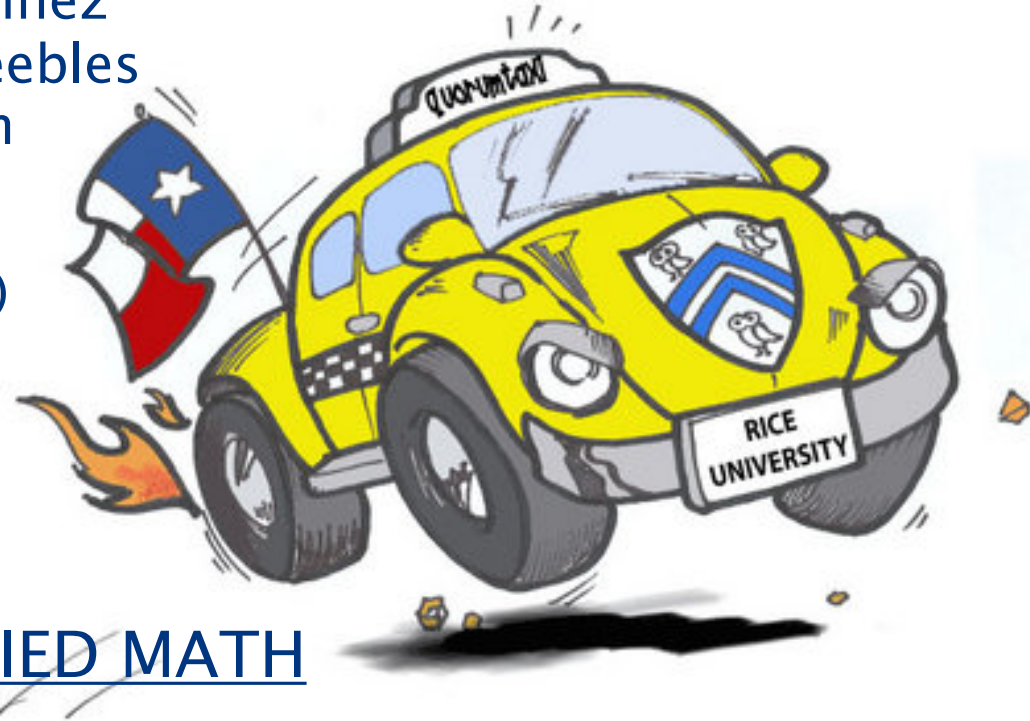
BIOENGINEERING

Irene Martinez

Christie Peebles

Ka-Yiu San

The Rice Taxis Drivers



COMPUTATION AND APPLIED MATH

Steve Cox

Jay Raol

Thomas Segall-Shapiro

CAIN PROJECT

Elizabeth McCormack

CHEMICAL AND BIOMOLECULAR ENGINEERING

Ken Cox

Dario Prieto

Miinkay Yu

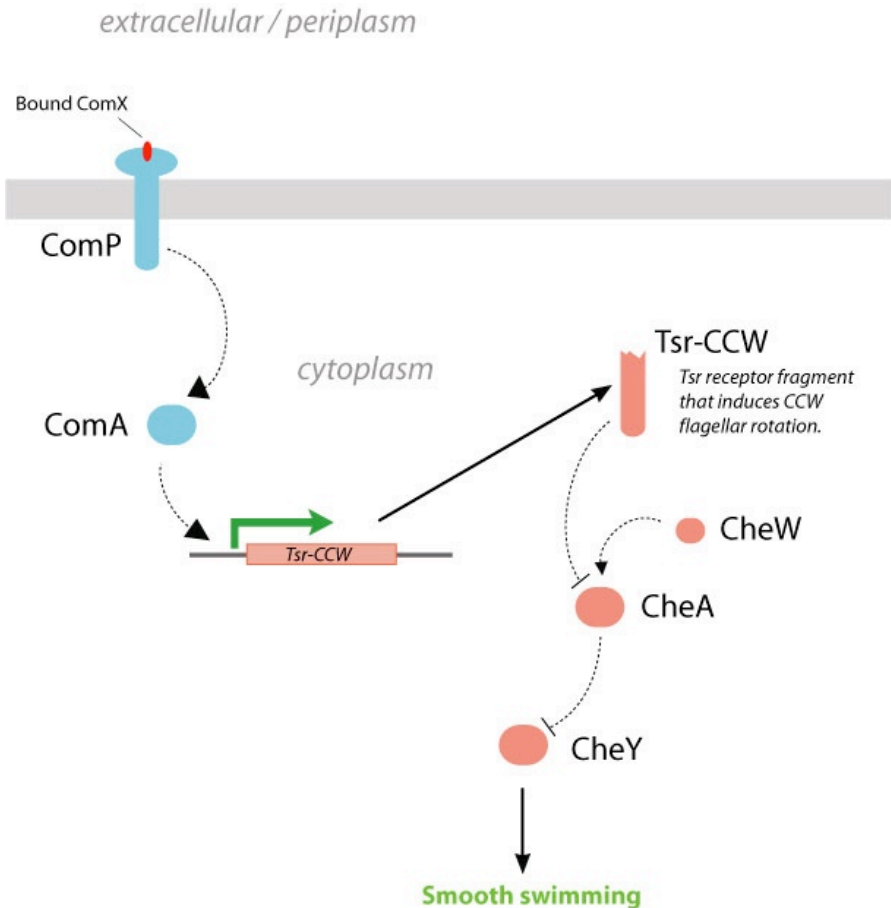
iGEM

Andrew Hessel

Extra Slides

Alternative Strategy

Transcriptional Control of Chemotaxis



§ Engineer *E. coli* cell to express ComP and ComA components of the *B. subtilis* quorum sensing pathway.

§ Reception of ComX by ComP

§ Activation of ComA, which induces transcription of a Tsr-CCW fragment leading to a smooth swimming phenotype