

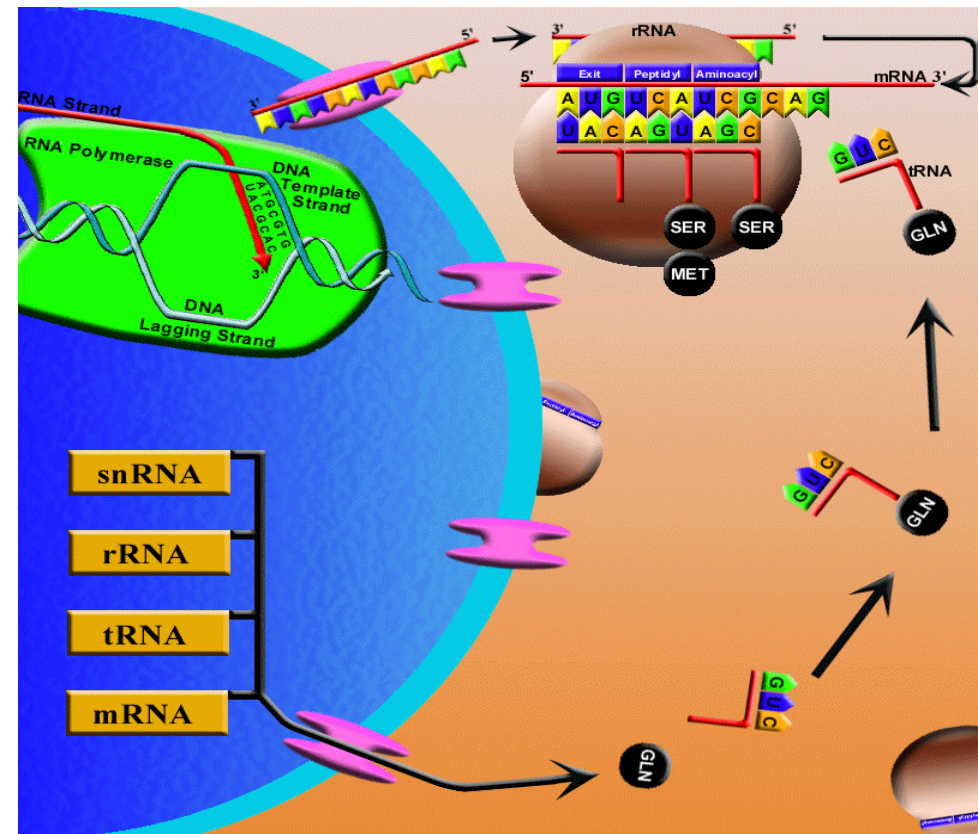


IPRO 302: Synthetic Biology

Engineering a novel organism

Nature's Engineering

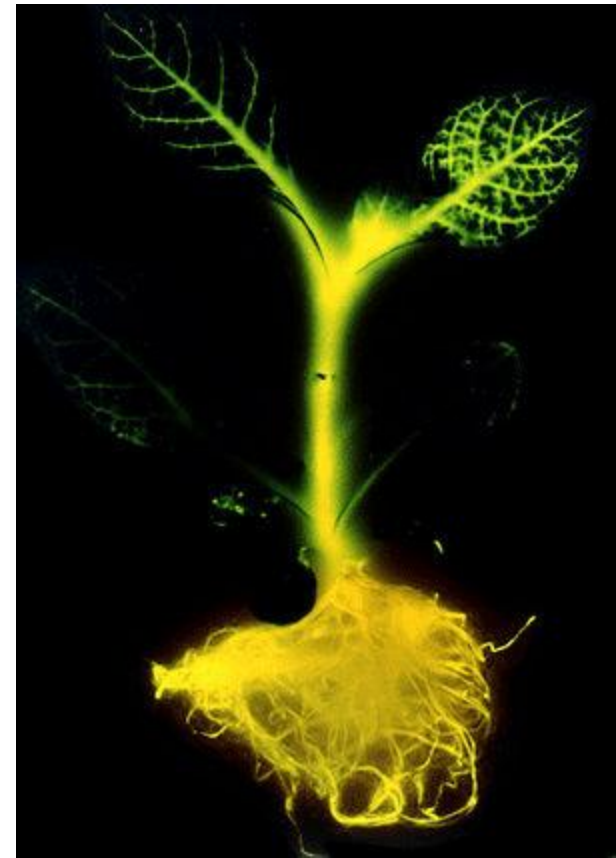
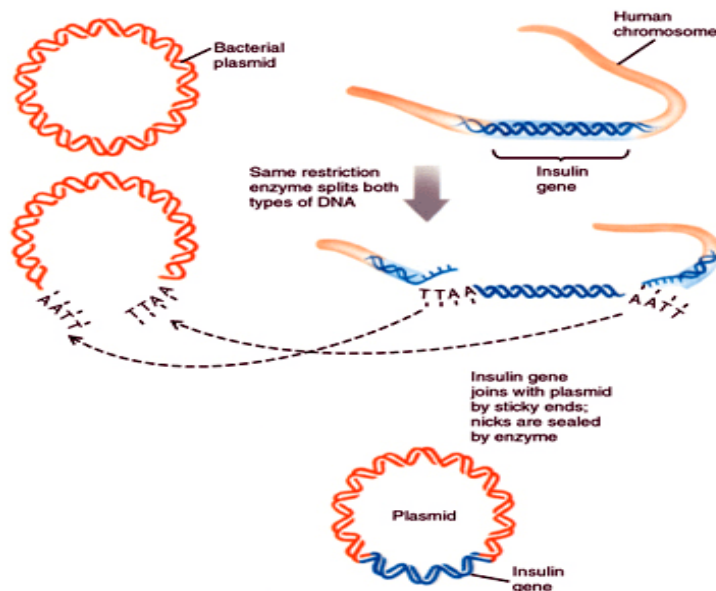
- Living things are made up of biological machines – mostly protein.
- Proteins made according to info stored in DNA
- Each organism has a master plan of DNA
- **So, can we make our own blueprints?**



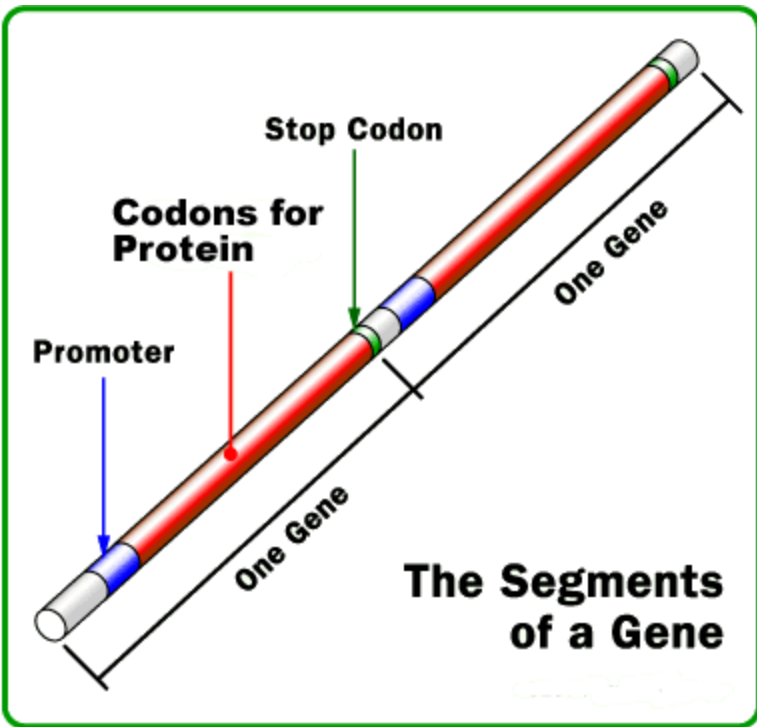
Genetic engineering

Add a DNA piece to a new organism, get a new protein

- 1000 s of examples



Elements of Control



- Not just machines, but the *right* machine in the right place at the *right* time
- **Control**
 - Promoters, terminators, localization signals
 - Syntax and punctuation

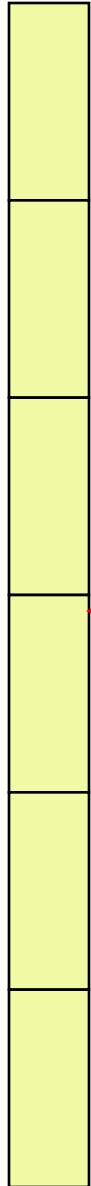


Higher Levels of Organization

- More Complex pathways with intercepting parts
 - Omega-3 pigs
 - Landmine sensing plants
 - Golden rice
 - Artemesin



Increasing Complexity



- Single gene = 1

- Gene and control = 2

- Gene and simple sensing system ~5

We are here

- Viruses ~50 genes

- Bacteria ~5000-30000

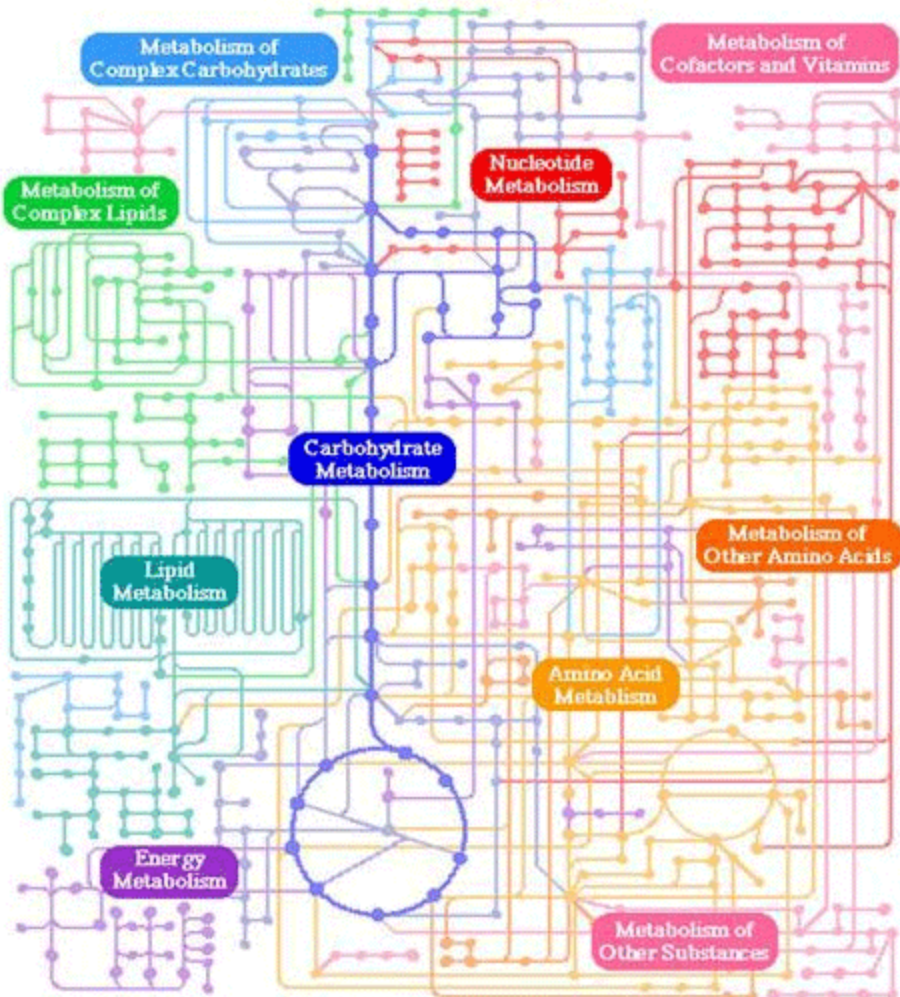
- Humans ~30,000, ~100,000 elements.

Genetic circuits

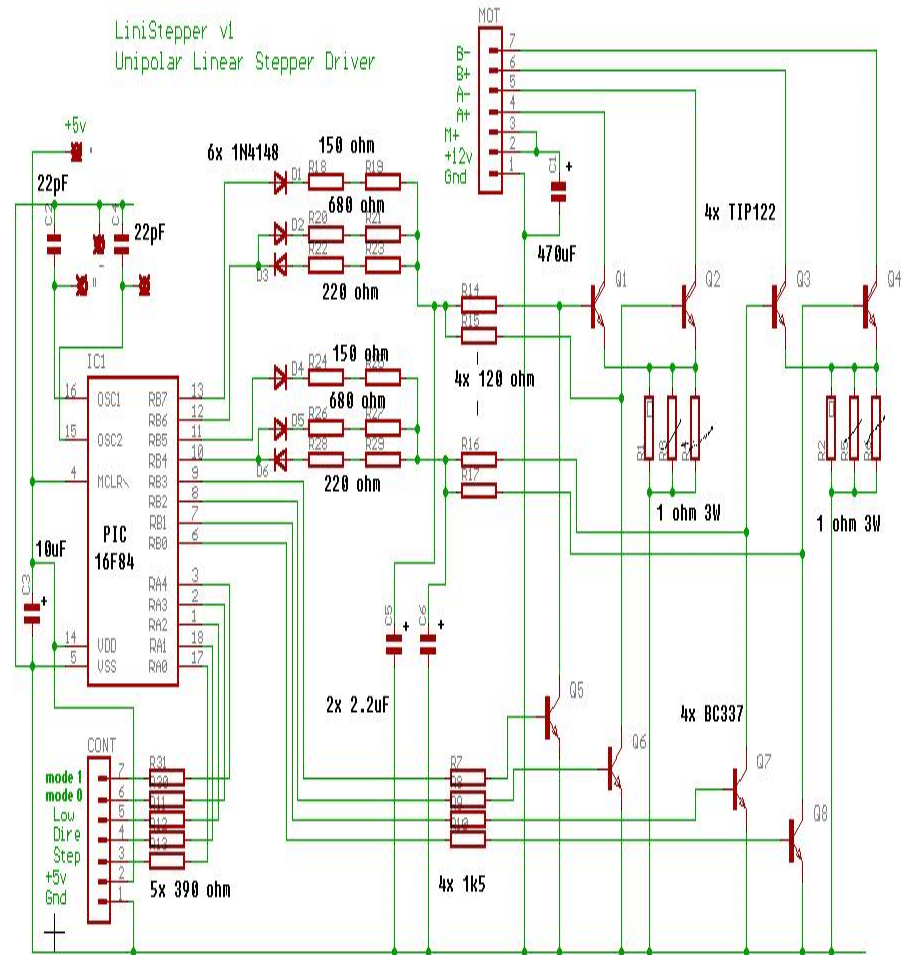
Metabolic pathways

Electronic circuits

METABOLIC PATHWAYS



LinStepper v1
Unipolar Linear Stepper Driver



Our project

The creation of dynamic system in a bacterium

- Tractable (<50-genes)
- Human readable

Implementation

- Construction through cheap lab methods

Understanding and design

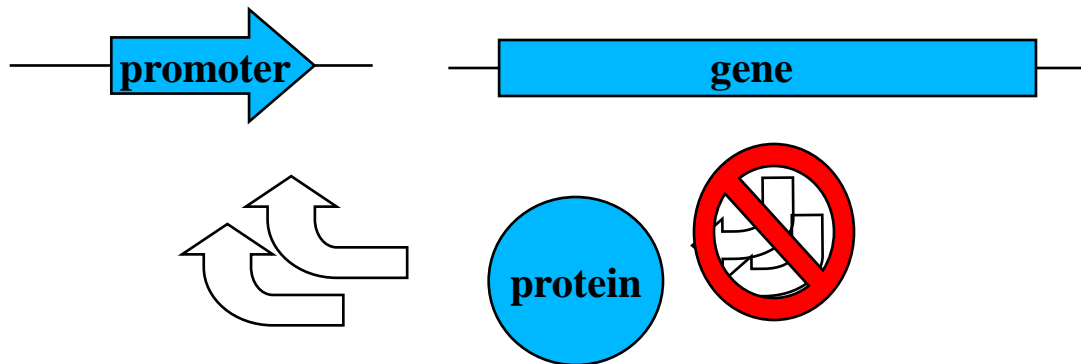
- Simulation through math modeling

Open-ended

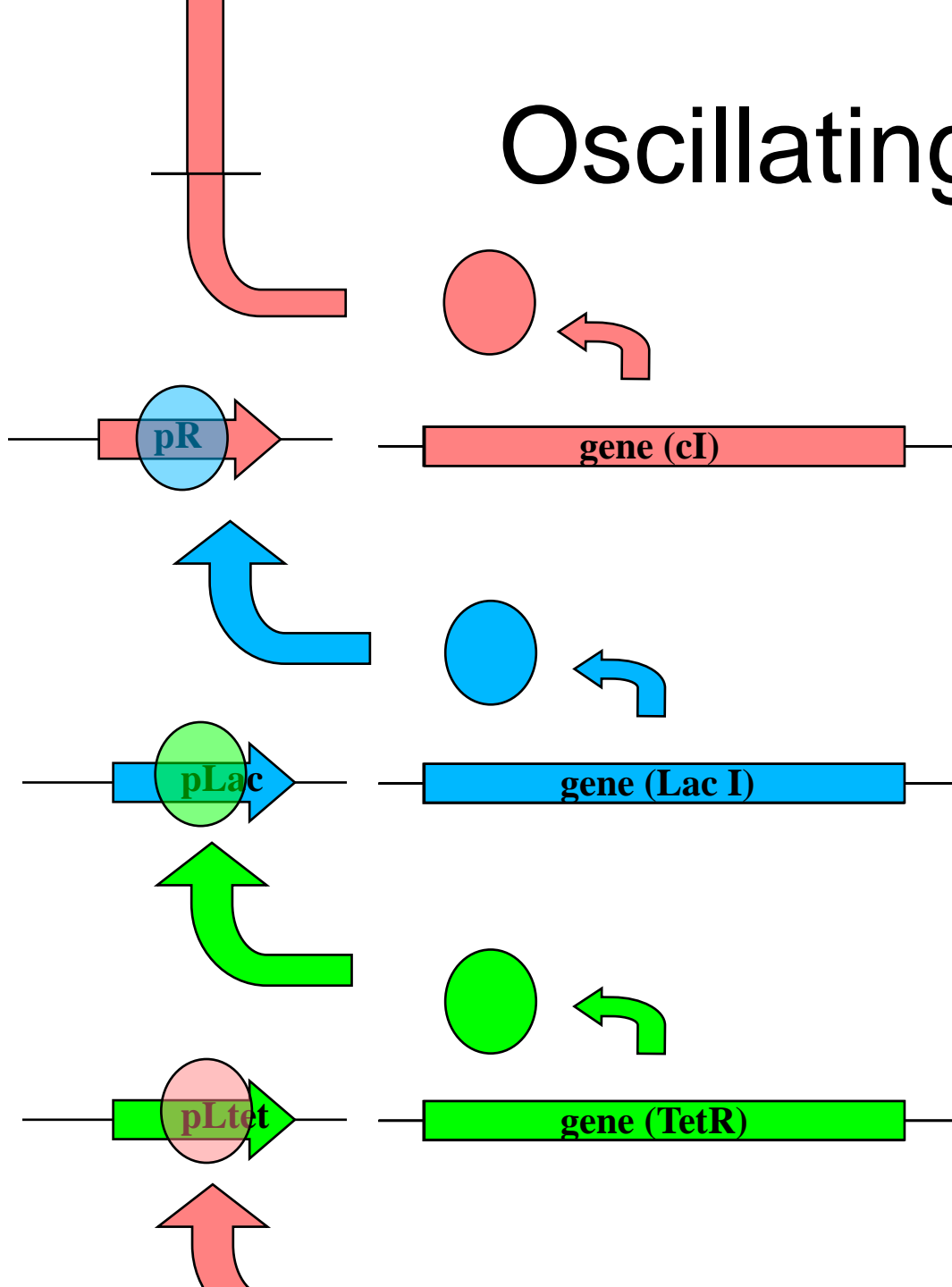
- Synchronization
- More charismatic organisms and designs

Biological System

- Feedback Systems
 - Thermostat
 - Hormone Homeostasis
- Negative Feedback
 - Product Inhibits Production



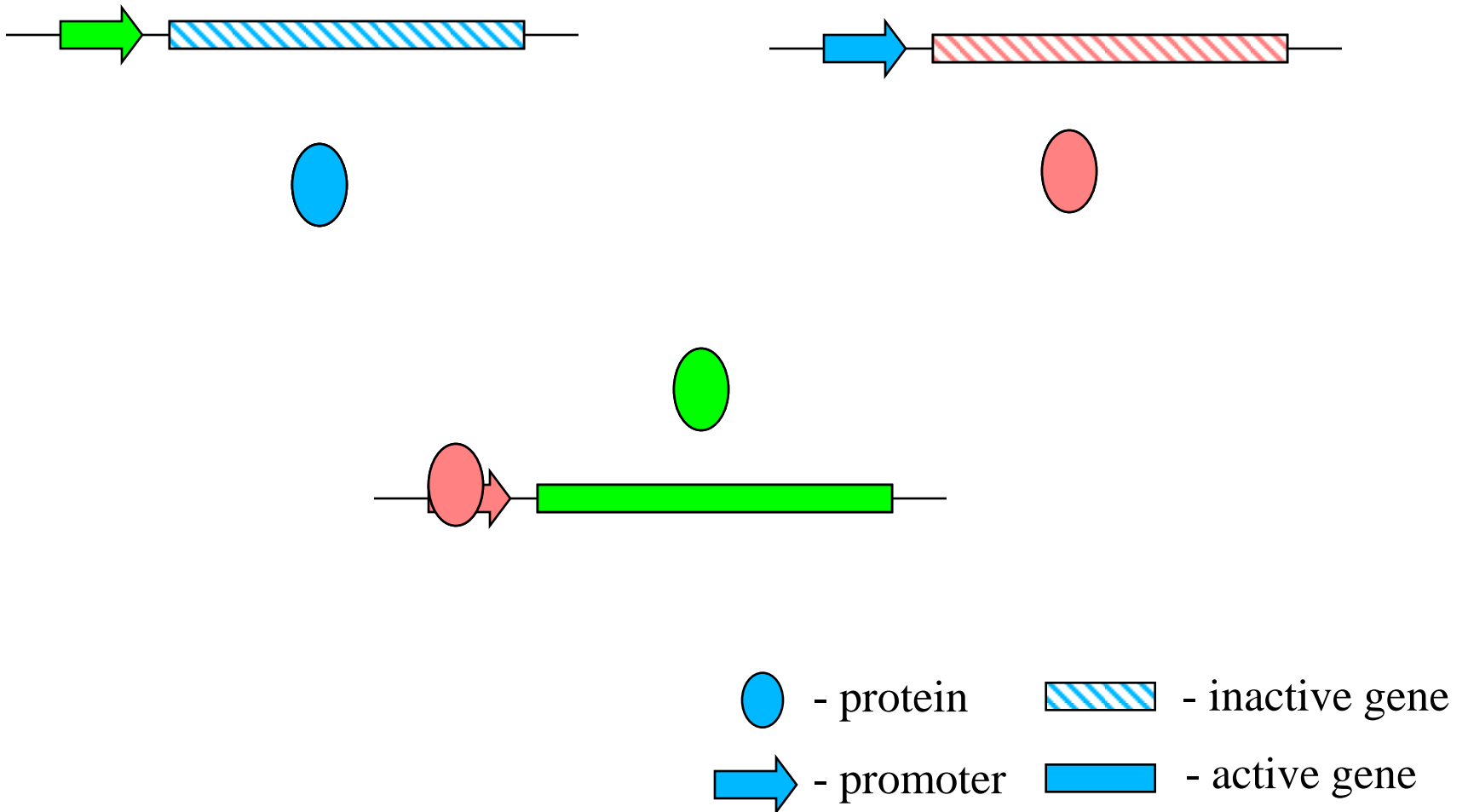
Oscillating System



Three gene system
that communicates
amongst its parts

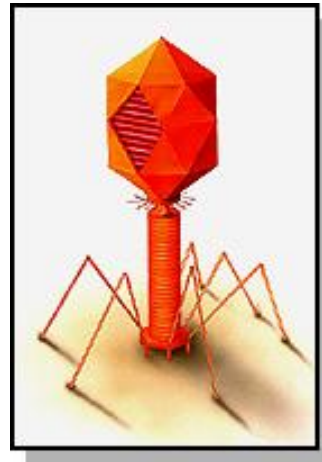
From Biology to Engineering

Rearrange gene pieces to obtain an oscillating system – how it all works together?



Obtained Genes from Many Sources

- Jelly Fish
 - XFP
- Bacteria
 - lacI
 - TetR
- Virus/Phage
 - pR – T7
 - cl
- Synthetic
 - pLTet
 - pLlac

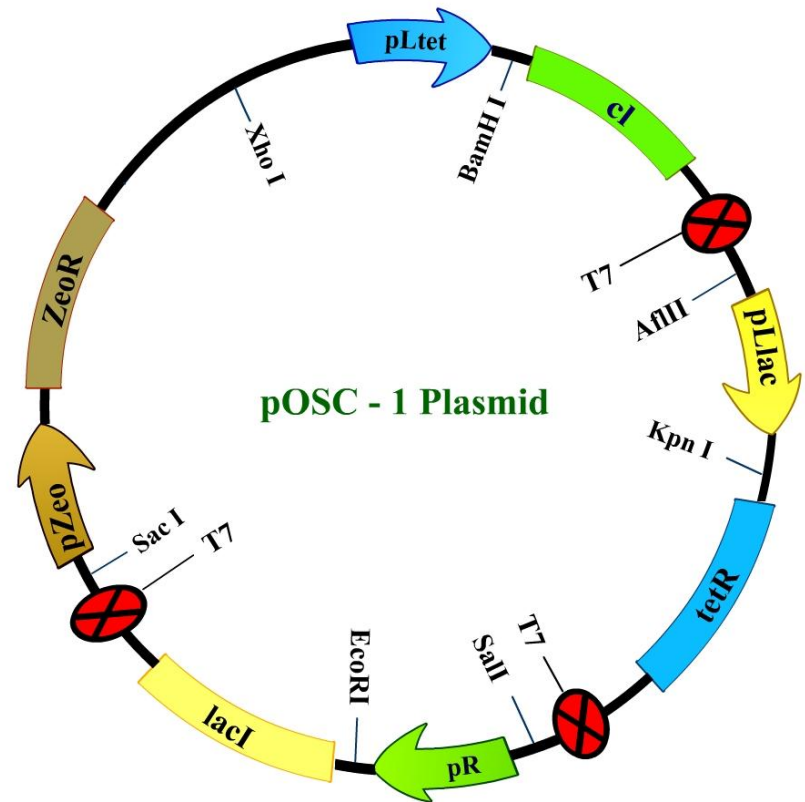
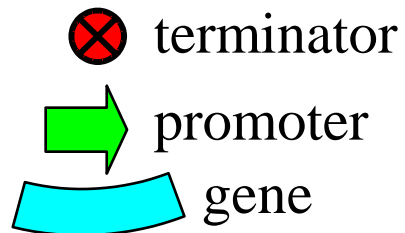


Assembly Techniques

- DNA Extraction
 - Remove DNA from Host Cells
- Amplify Target Gene
 - PCR Amplification
- Connect Genes into Plasmids
 - Fusion PCR
- Insert Plasmids into Target Cells
 - Competent Cell Transformations

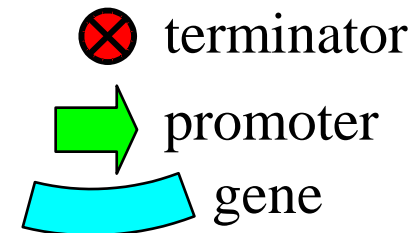
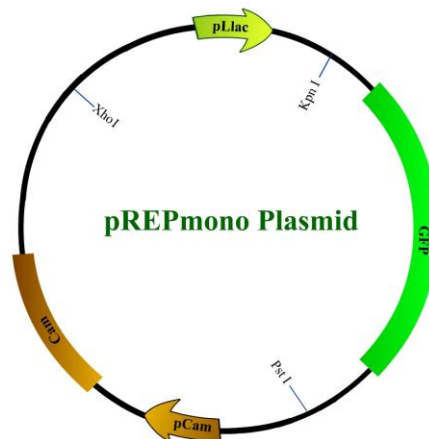
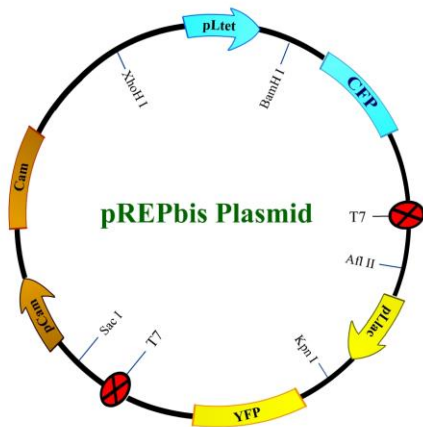
Oscillator plasmid

- All gene modules are put together in a plasmid
- The system oscillates so that only some genes are active at a certain time



Reporter Plasmid

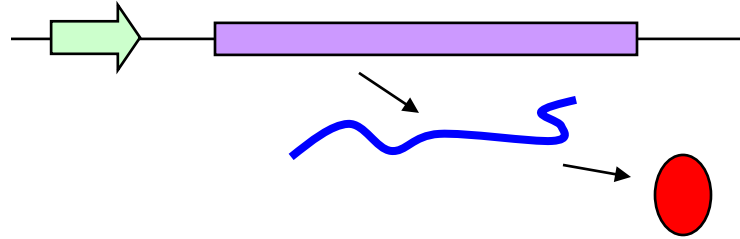
- Cannot see the Oscillations!!!
- Can See Proteins from Oscillatory Plasmid
 - Duplicate Promoter to Track Active Genes
- Fluorescent Protein Produced Depends on Gene activated in pOSC



Design Aspects

- In order to have direction in this project, we need:
 - Stochastic arithmetic modeling
 - Imaginative visions for the future

Modeling



$$\text{Behavior}(t) = \text{Model}(\alpha_0, b, m, p, k \dots)$$

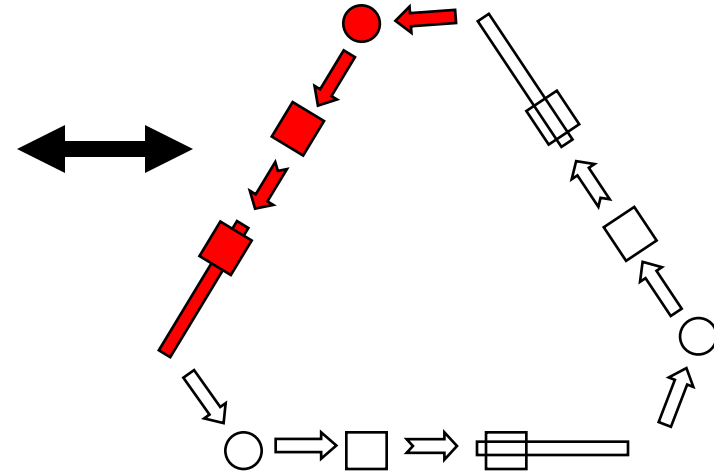
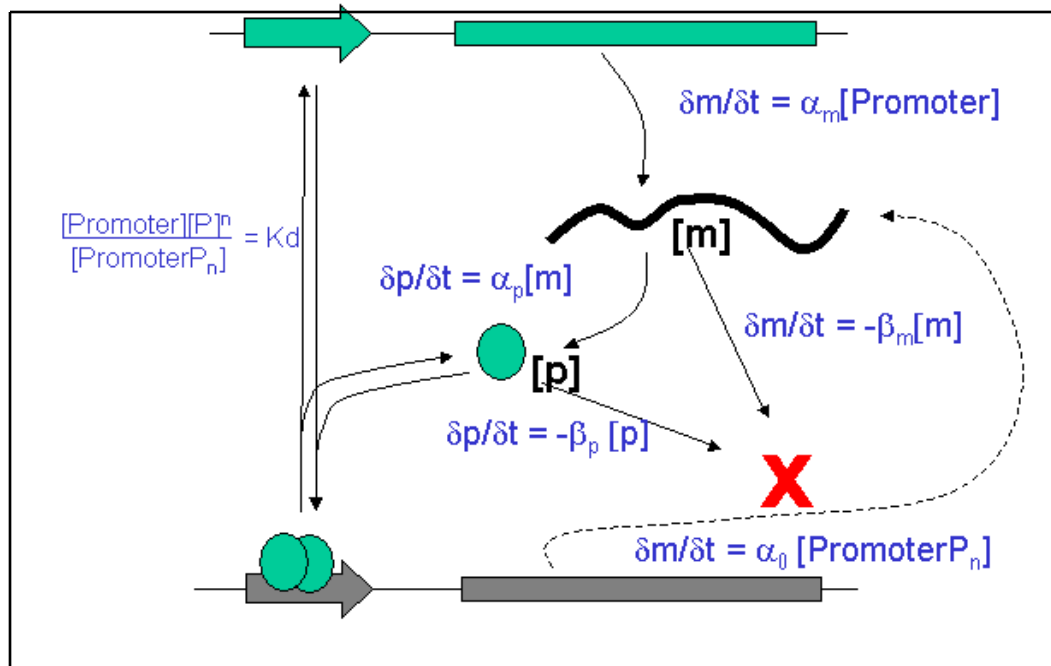
α_0 = promotor strength

b = RNA half life

m = amount of mrna

p = amount of protein

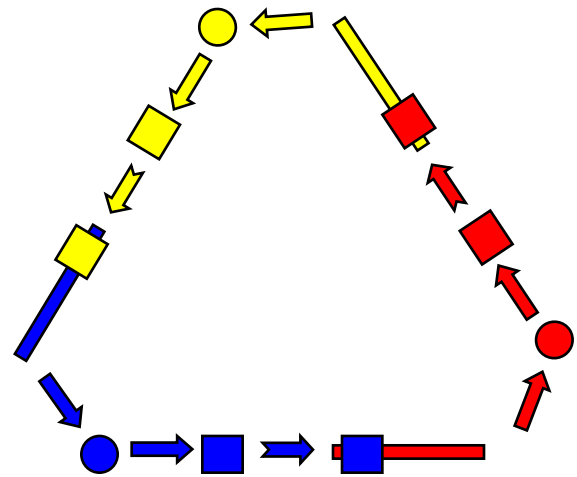
Basic system: One branch



$$\delta m_i / \delta t = \alpha 0_{i+1} + \alpha 1_{i+1} p^n_{i+1} / (K^n_{i+1} + p^n_{i+1}) - \beta_m m_i$$

$$\delta p_i / \delta t = \alpha_m m_i - b_p p_i$$

All three



mRNA Equations:

$$dm/dt = -\beta_m m + \alpha_1 p^n / (k_d + p^n) + \alpha_0$$

$$dm/dt = -\beta_m m + \alpha_1 p^n / (k_d + p^n) + \alpha_0$$

$$dm/dt = -\beta_m m + \alpha_1 p^n / (k_d + p^n) + \alpha_0$$

Protein Equations:

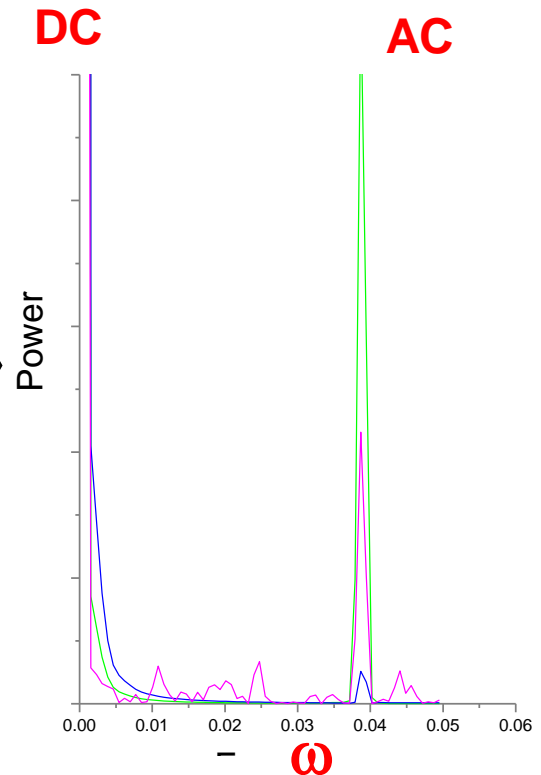
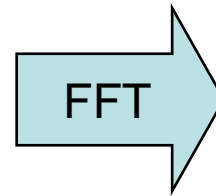
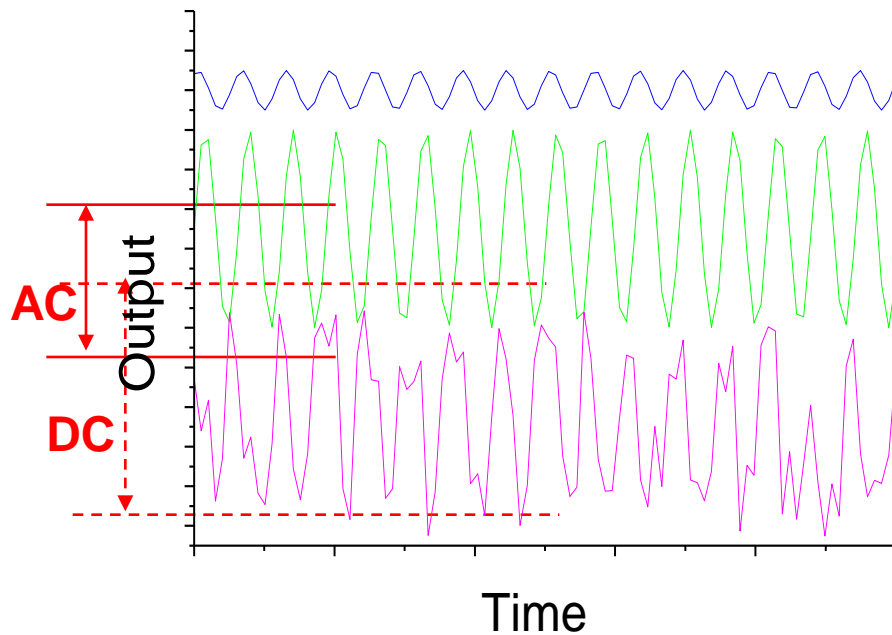
$$dp/dt = t_s * m - bp * m$$

$$dp/dt = t_s * m - bp * m$$

$$dp/dt = t_s * m - bp * m$$

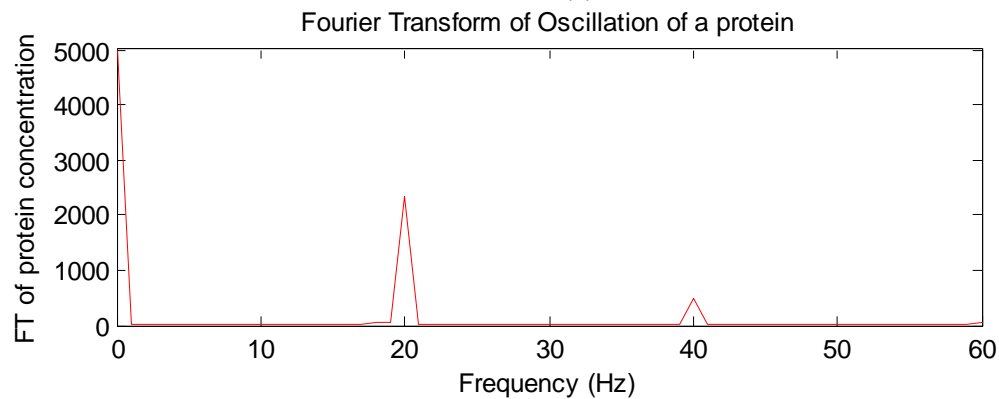
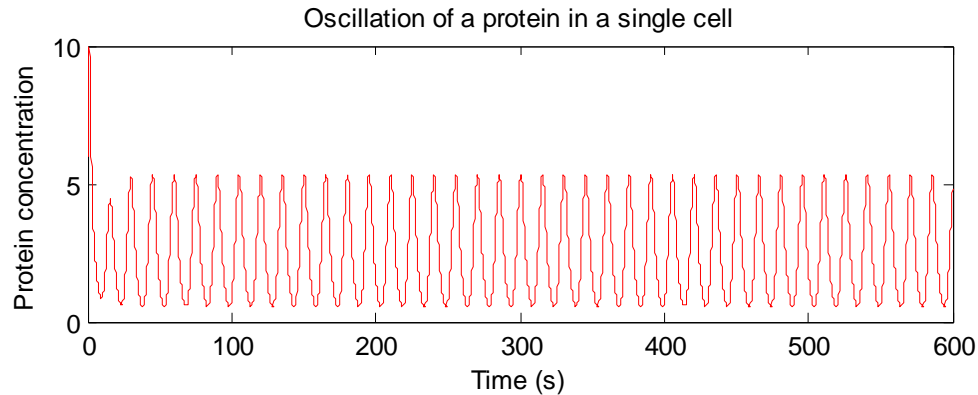
Analysis

- What do we need to know and how do we measure it?



First simulation:GUI

Deterministic



Stochastic Model

- Functional species are limited integer amounts only

– Poisson distribution

$$\lambda_m = -\beta_m m + \alpha_1 p^n / (k_d + p^n) + \alpha_0$$

$$dp/dt = t_s * m - bp * m$$

$$P_{\lambda_m}(n) = \lambda^n e^{-\lambda} / n!$$

$$\Delta m / \Delta t = \text{rand}(P_{\lambda_m}(n))$$

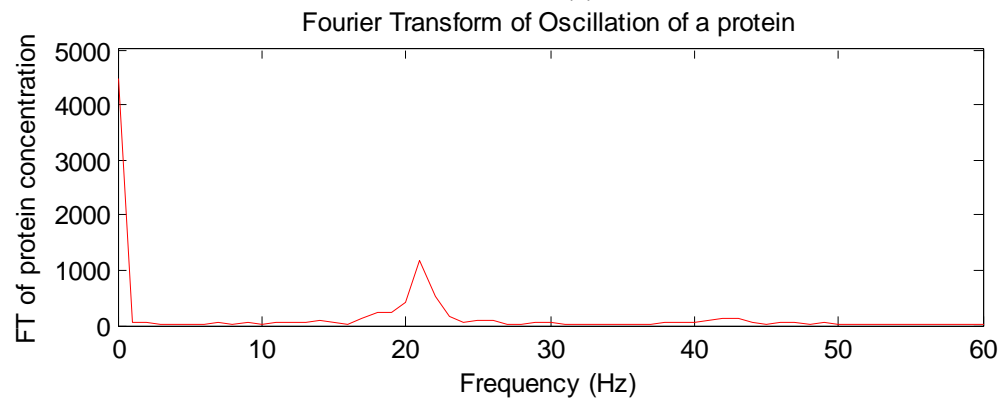
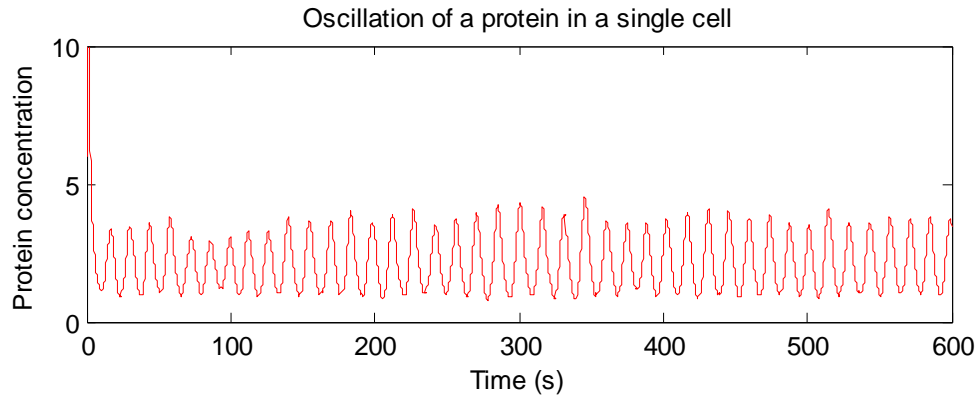
$$dm/dt = -\beta_m m + \alpha_1 p^n / (k_d + p^n) + \alpha_0$$

$$dp/dt = t_s * m - bp * m$$

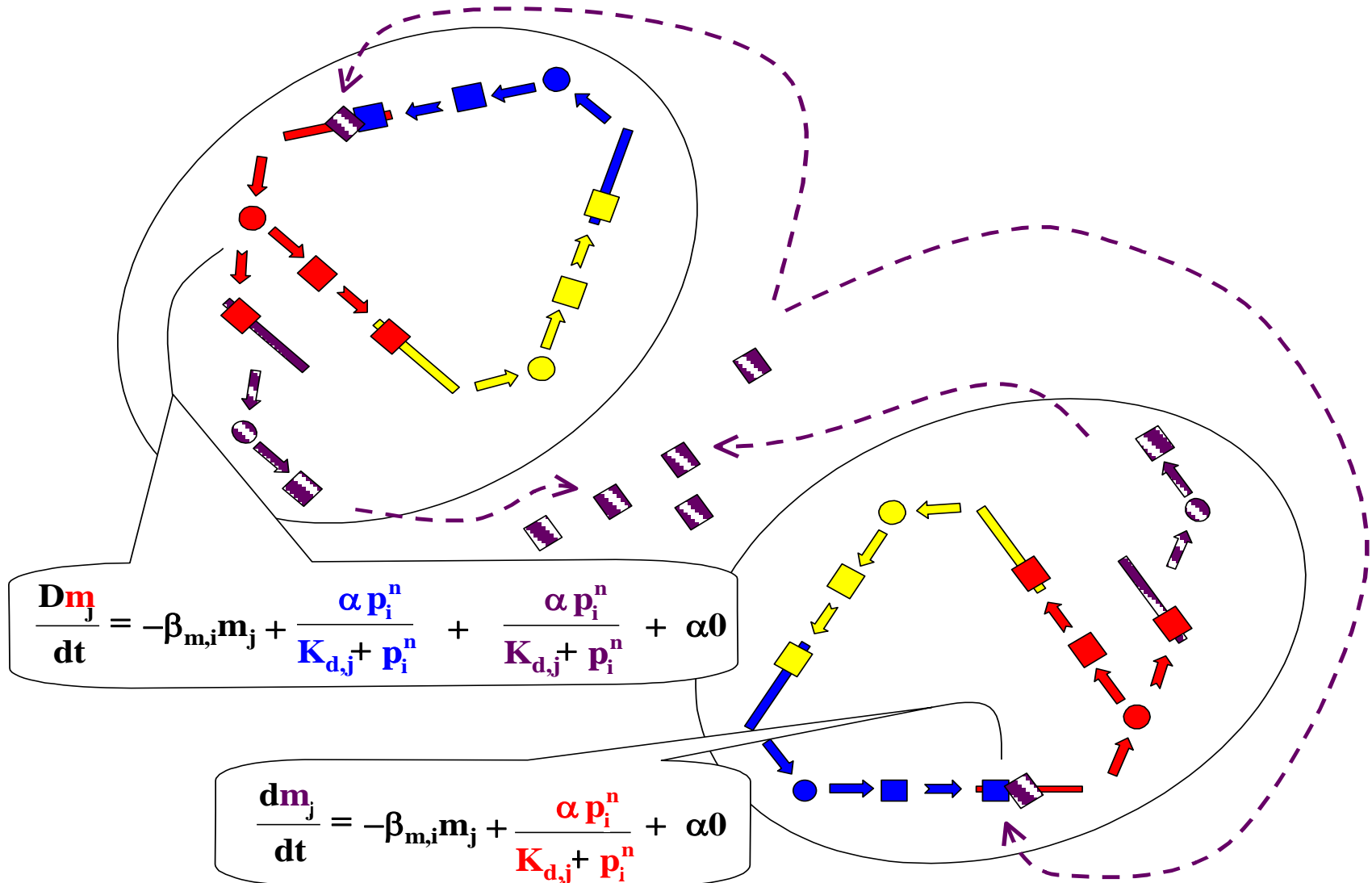
$$dm/dt = -\beta_m m + \alpha_1 p^n / (k_d + p^n) + \alpha_0$$

$$dp/dt = t_s * m - bp * m$$

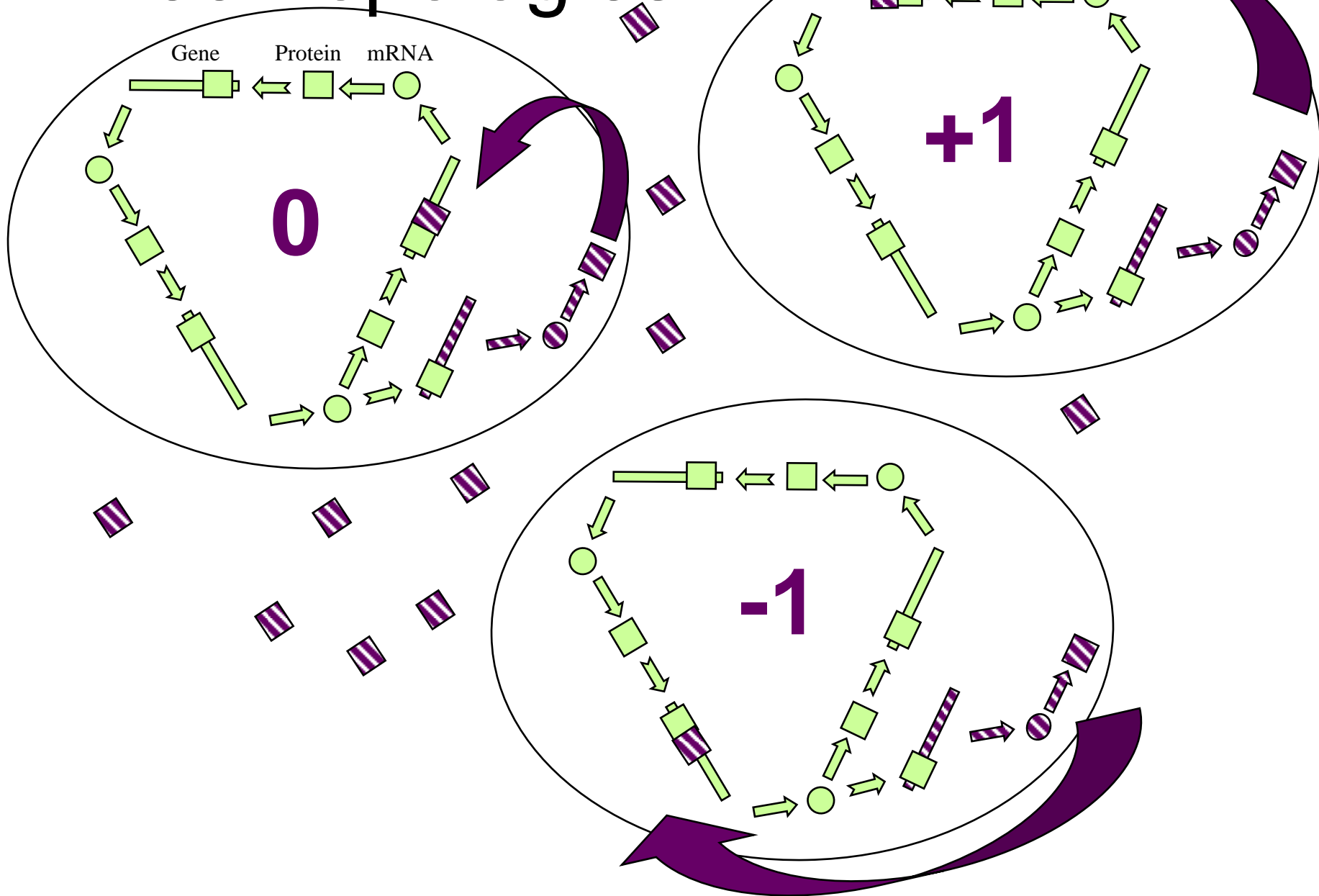
Second Simulation: GUI Stochastic



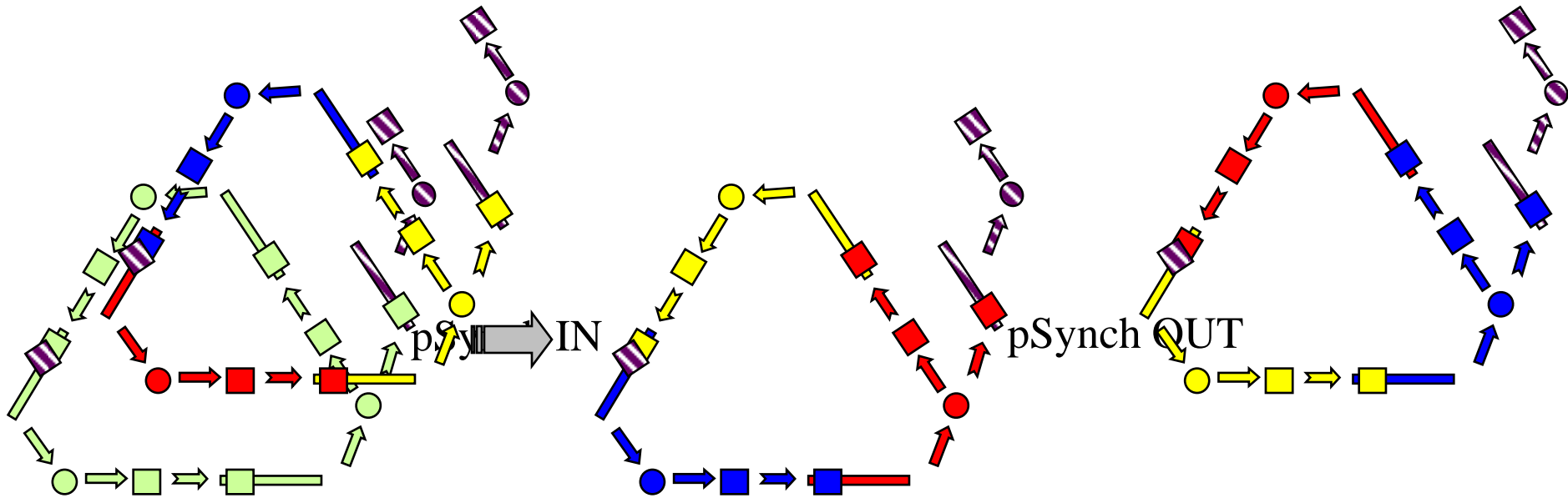
Synchronization



Three Topologies



Three Branches



$$\frac{dm_j}{dt} = -\beta_{m,i}m_j + \frac{\alpha_1}{K_{d,j} + p_i^n} + \alpha_0$$

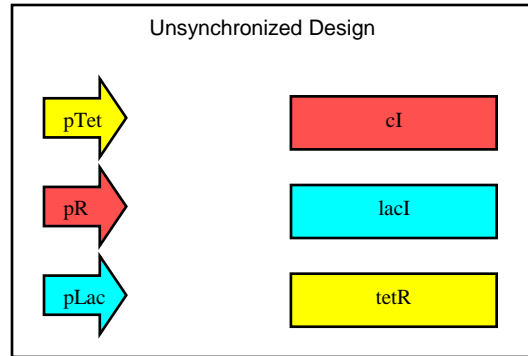
Which one?

- Strongest α_1 ?
- Least leaky α_0 ?
- Highest ON/OFF ratio α_1/α_0 ?
- $K_{d,s}$, $K_{d,out}$?
- n_h ?

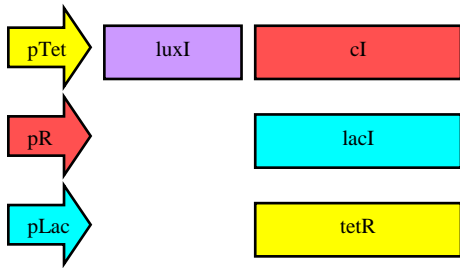
Other factors

- $\beta_{m,s}$
- $\beta_{p,s}$

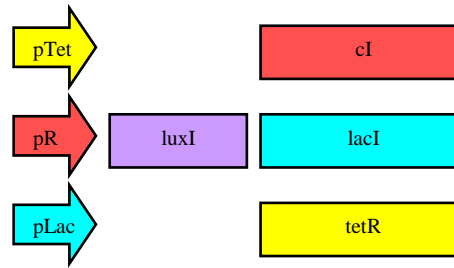
Synchronization Designs



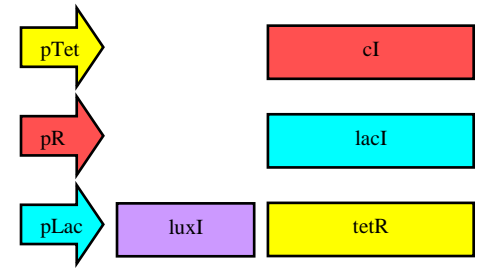
Design 1



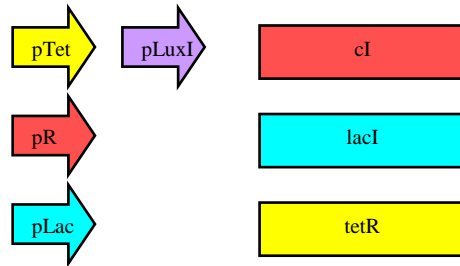
Design 2



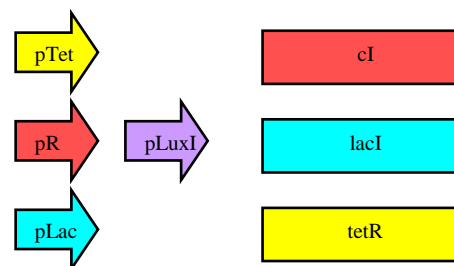
Design 3



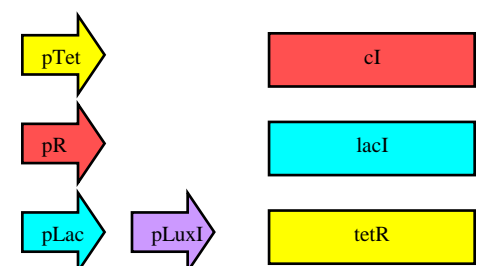
Design A



Design B



Design C



mRNA Equations:

Modeling

Protein Equations:

$$\frac{dm_i}{dt} = -m_i + \frac{\alpha_1}{1 + p_k^n} + \alpha_0 + \frac{kS_{int}}{1 + S_{int}}$$

$$P_{\lambda m}(n) = \lambda^n e^{-\lambda} / n!$$

$$\Delta m / \Delta t = \text{rand}(P_{\lambda m}(n))$$

$$\frac{dm_j}{dt} = -m_j + \frac{\alpha_1}{1 + p_i^n} + \alpha_0 + \frac{kS_{int}}{1 + S_{int}}$$

$$P_{\lambda m}(n) = \lambda^n e^{-\lambda} / n!$$

$$\Delta m / \Delta t = \text{rand}(P_{\lambda m}(n))$$

$$\frac{dm_k}{dt} = -m_k + \frac{\alpha_1}{1 + p_j^n} + \alpha_0 + \frac{kS_{int}}{1 + S_{int}}$$

$$P_{\lambda m}(n) = \lambda^n e^{-\lambda} / n!$$

$$\Delta m / \Delta t = \text{rand}(P_{\lambda m}(n))$$

$$\frac{dp_i}{dt} = -\beta (p_i - m_i)$$

$$P_{\lambda m}(n) = \lambda^n e^{-\lambda} / n!$$

$$\Delta m / \Delta t = \text{rand}(P_{\lambda m}(n))$$

$$\frac{dp_j}{dt} = -\beta (p_j - m_j)$$

$$P_{\lambda m}(n) = \lambda^n e^{-\lambda} / n!$$

$$\Delta m / \Delta t = \text{rand}(P_{\lambda m}(n))$$

$$\frac{dp_k}{dt} = -\beta (p_k - m_k)$$

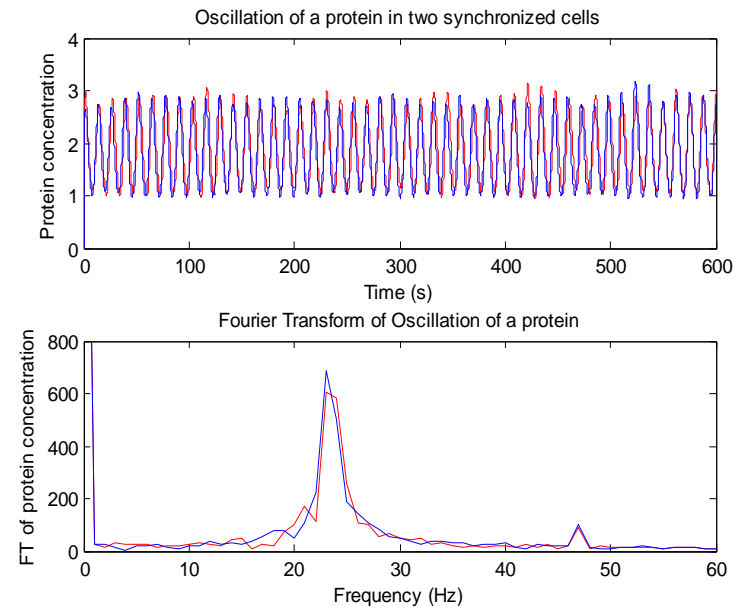
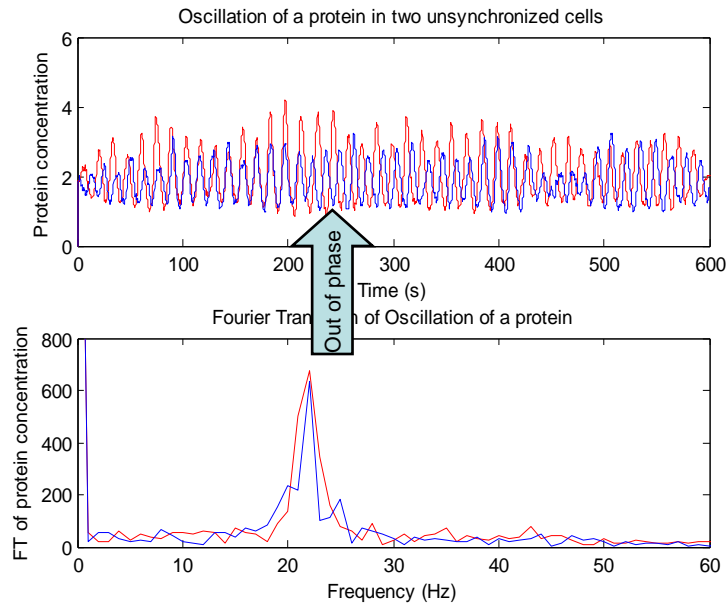
$$P_{\lambda m}(n) = \lambda^n e^{-\lambda} / n!$$

$$\Delta m / \Delta t = \text{rand}(P_{\lambda m}(n))$$

$$\frac{dS_{int}}{dt} = -k_{s0} S_{int} + k_{s1} p_k - \eta (S_{int} - S_{ext})$$

$$\frac{dS_{ext}}{dt} = -k_{se} S_{ext} + \eta \sum_{n=1}^N (S_n - S_{ext})$$

Unsynchronized vs. Synchronized



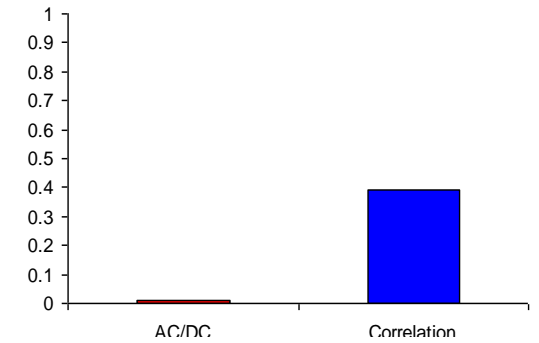
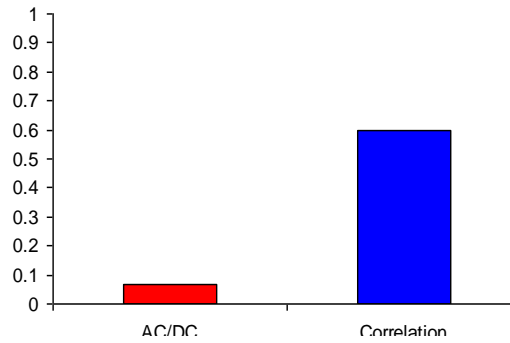
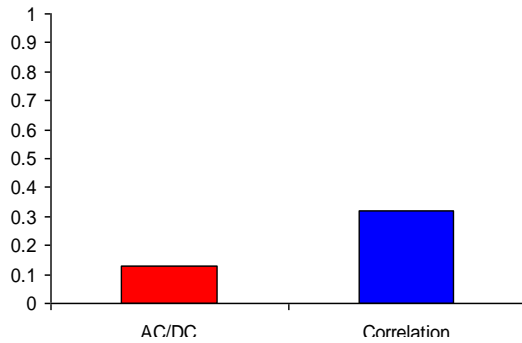
Topology Analysis

1

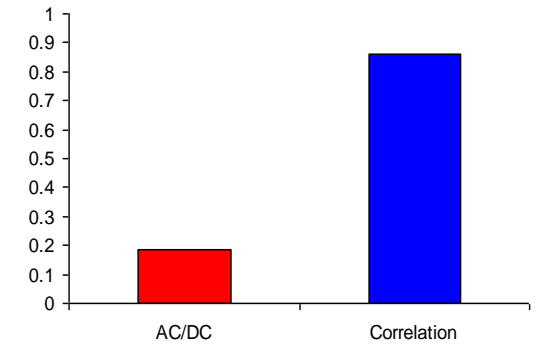
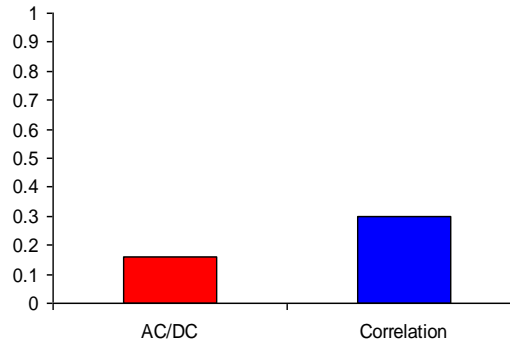
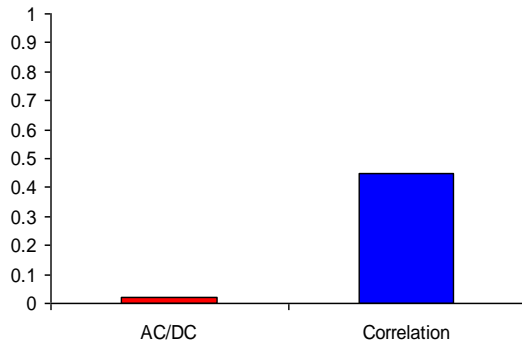
2

3

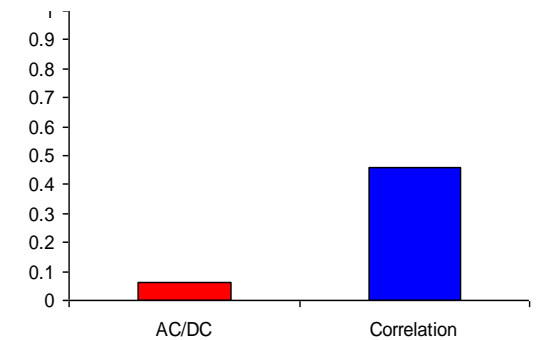
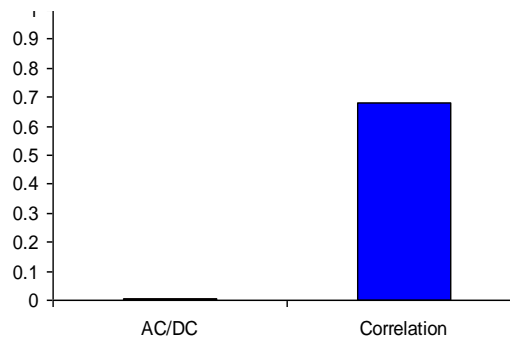
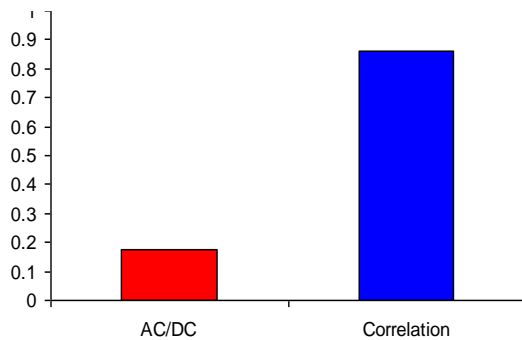
A



B



C



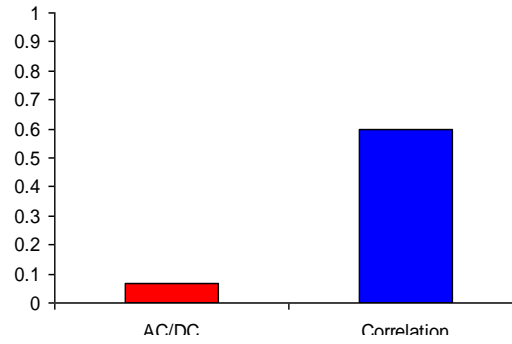
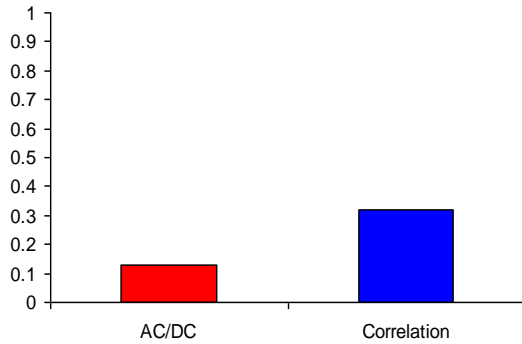
Strong Oscillation

1

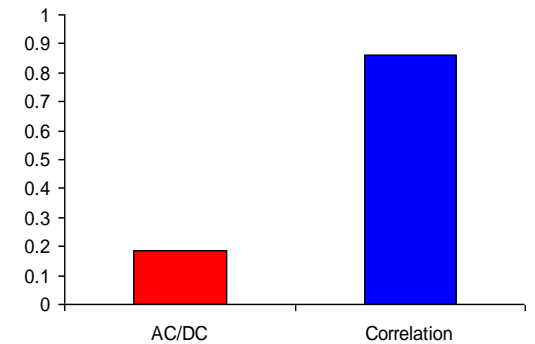
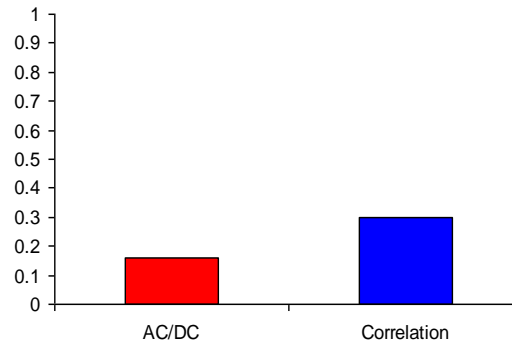
2

3

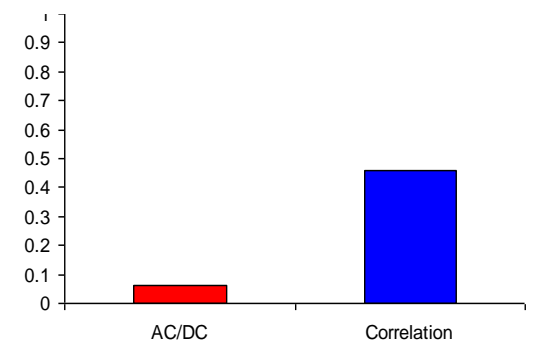
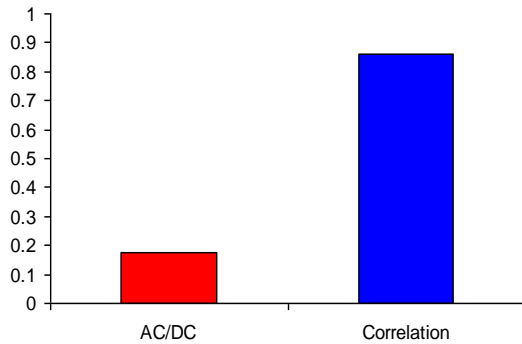
A



B



C



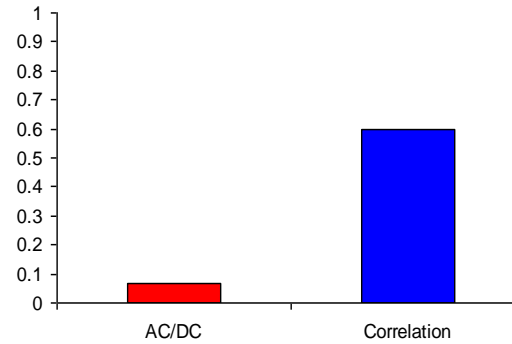
Good Synchronization

1

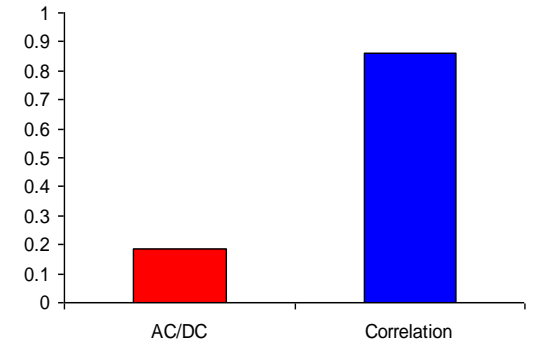
2

3

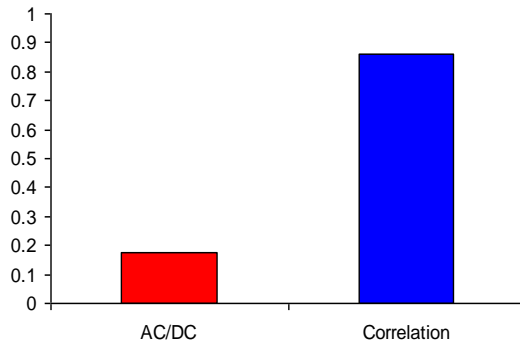
A



B

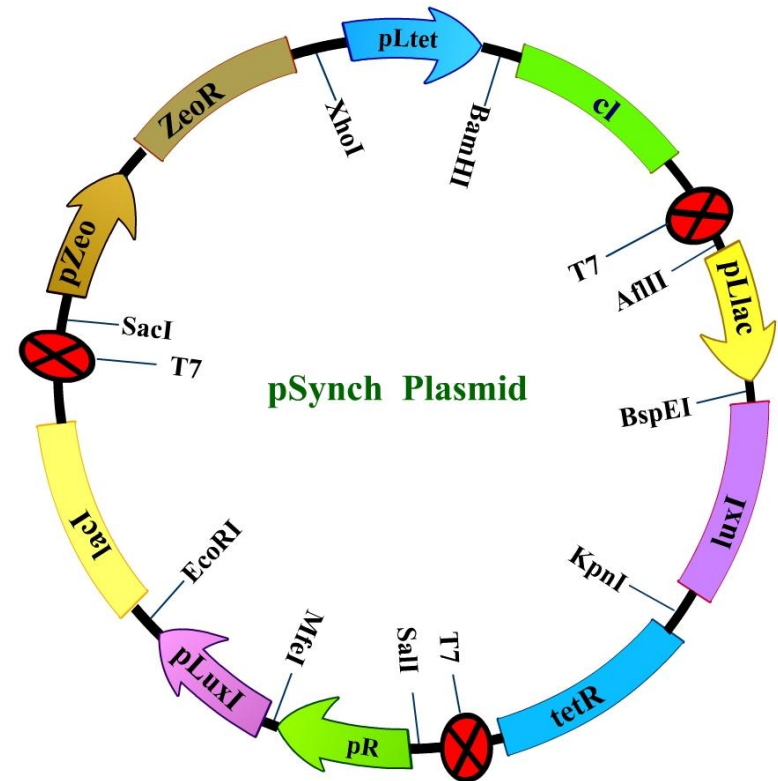


C



New target plasmid: pSynch

- We have the genes and preliminary PCR products
- Need to develop construction blueprint



Development

- Engineering multicellular organisms
 - ✓ Zebrafish
 - ☐ Humans?

–Scientific Issues

- Synchronization
- Different O/S

–Regulatory Issues

–Ethical Issues



Regulatory & Ethical Issues

- Regulations

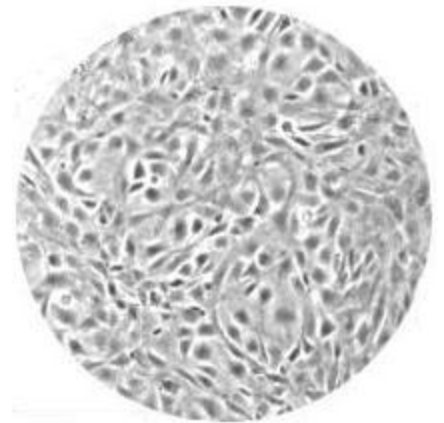
Bodies like IACUC have final say

- Justifiable use of animals
- Minimal pain
- Purpose

- Ethical Concerns

Should we be doing this? Why?

- Understanding complex biomedical systems
- What is life?



Engineered Zebra Fish
Culture Cells

Thank you!

