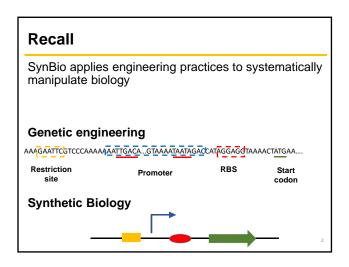


Instructor: Prof. K. Solomon Ph.D. Assistant Professor Agricultural & Biological Engineering Laboratory for Renewable Resources Engineering

Fall 2018





This lecture....

- Intro to Engineering design
- SynBio hierarchy
- · Biological parts
- Part databases

Key features of engineering design

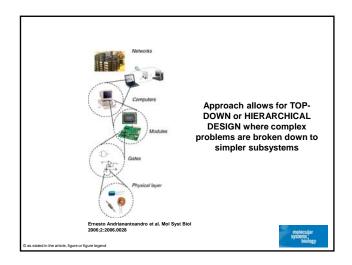
 Abstraction – "black boxing" the system to focus on core properties/functions of a system



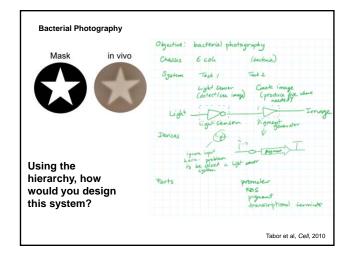
- 2. Modularization creation of essential parts with a single function that can be recombined in diverse contexts
- 3. Standardization parts have fixed interfaces for rapid assembly & prototyping

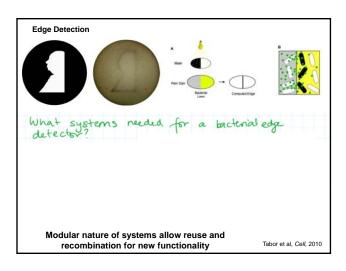
BE/ChE – Unit ops e.g. Distillation columns, CSTRs **EE** – electronic components e.g. resistors, capacitors

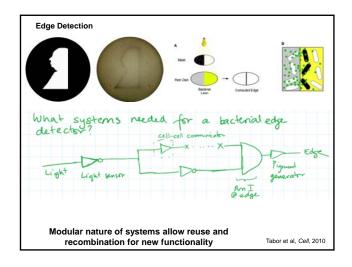
4

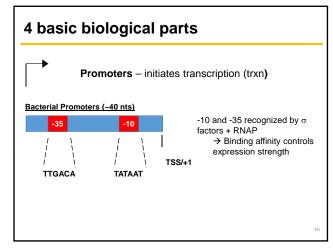


Synthetic Biology Hierarchy Problem Microbial production of insulin 4. Chassis: Host or organism 3. System: Combination of devices Insulin producing gene circuit that execute a specified task Insulin protein expression, gene regulation cassette (e.g. YES 2. Device: Combination of parts that achieve a single human specified gate) 1. Part: Minimal DNA required to A promoter, CDS, ribosome achieve a biological function binding site, regulatory protein, operator sites, 0. DNA: Physical specification









Controlling the regulatory output of a promoter

Regulator proteins (transcription factors) bind to **operator sites** within/around a promoter

- → Repressor proteins bind within or downstream of promoter and compete with RNAP for binding
 - e.g. tetR/tetO, araC/araO
 - can also form a trxn roadblock
 - e.g. lacl/lacO



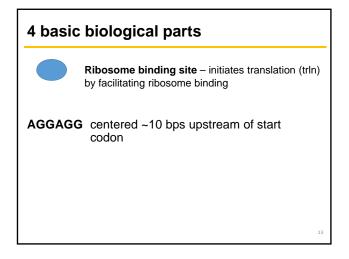
→ Activator proteins bind to upstream operators to recruit RNAP or change DNA conformation

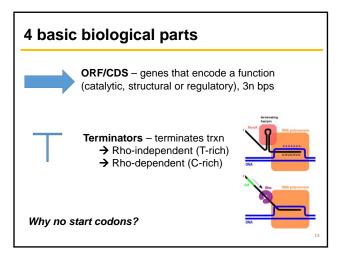
e.g. CAP, araC

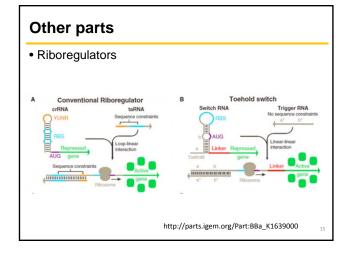
Regulator proteins

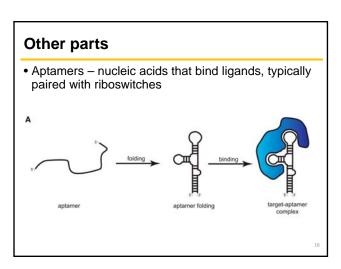
- Regulator proteins change conformation when exposed to ligands
 - lacl and lactose/IPTG
 - araC and arabinose
 - tetR and tetracycline/doxycycline
- Conformation determines whether bound to DNA or not

12







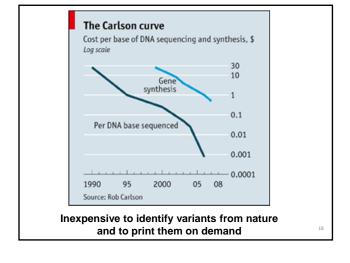


Parts have properties

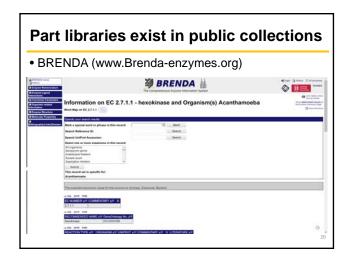
- Parts have distinct input/output
- Parts have 'strengths' that control their output
 - E.g. strong promoter == high expression
- Properties controlled by architecture and sequence

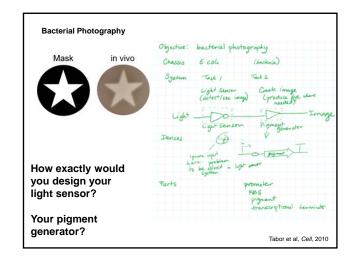
What does a promoter that responds to arabinose look like? Tetracycline?

How would you engineer a weaker tetracycline-inducible promoters?



Part libraries exist in public collections Registry of Standard Biological Parts (parts.igem.org) Registry of Standard Biological Parts (parts.igem.org) Registry of Standard Biological Parts From the Standa





Next time....

• What are the input-output relationships between parts?

22