Synthetic Biology = design and engineering of biological systems that aren't found in nature

Why would we want to do this?

- Want to understand natural systems. One of the best ways to understand a system is to change it or make new, related ones
- To fully "understand" a system, we should be able to predict the outcome when we change the system
- For molecular biology, this means:
 - designing new gene circuits and networks
 - modeling the designed systems & predicting their properties
 - making & testing the designs
 - updating our understanding from the model/test agreement

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Engineers often look at biological systems & think that the systems are equivalent to electronic circuits

e.g,

fluorescent proteins light bulbs or LEDs transcription factors transistors or logic gates

repressors NOT gates activators OR/AND gates

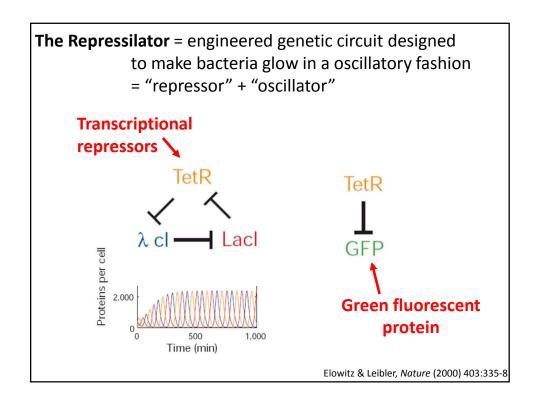
polymerases

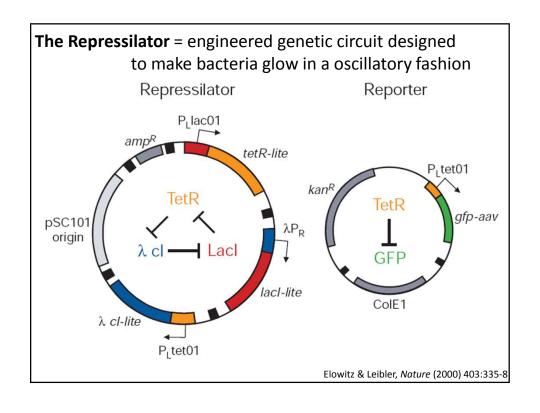
(transcriptional machinery) batteries

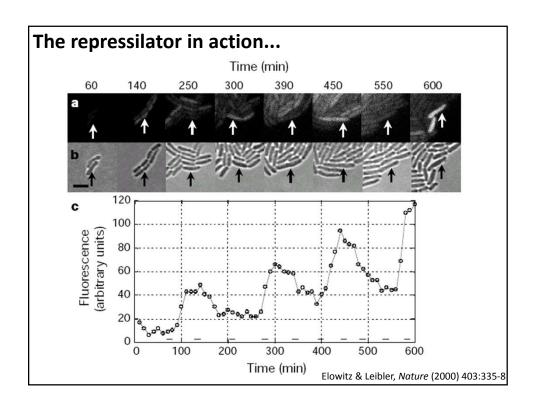
and so on...

Are they right?

→ raises the possibility that biological parts (genes, proteins, etc.) could be combined using the rules established for analog/digital circuits



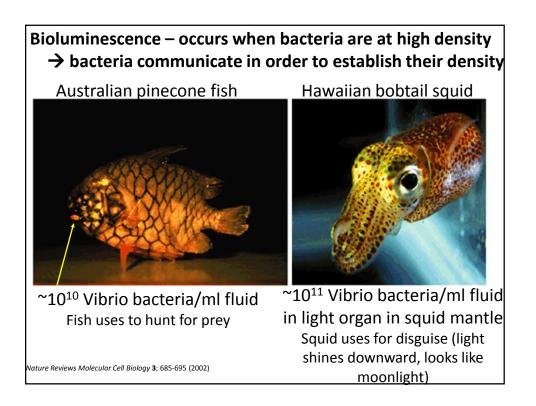


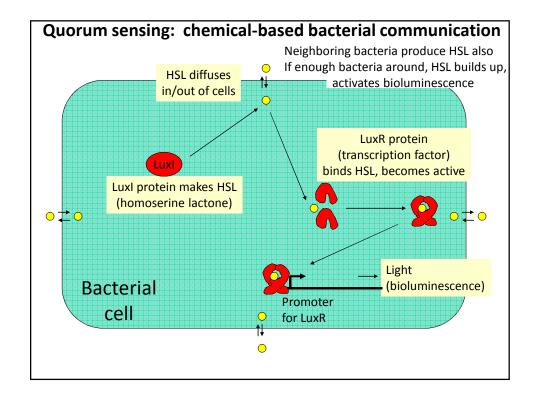


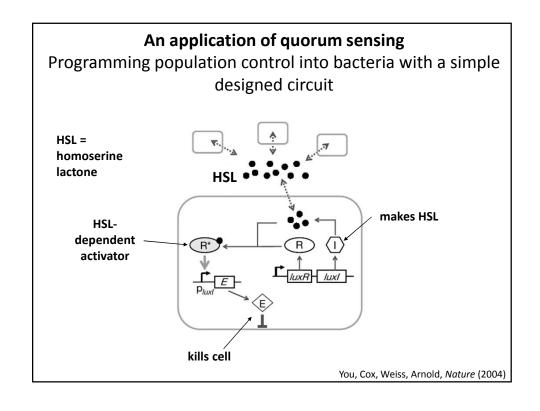
What other kinds of circuits can be built? First, we need some more parts!

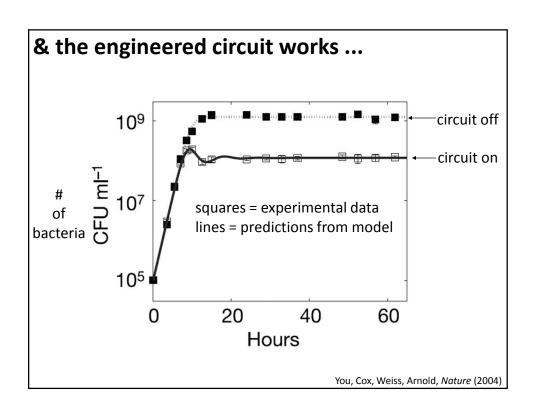
Some of the other parts available include:

- various sensors
 - light, dark, heat, cold
- more switches, logic gates
 - more repressors, activators
- parts for intracellular communication
 - helpful if cells could tell each what condition they're in
 - → quorum sensing
- parts for signaling the output of circuits
 - fluorescent & luminescent proteins







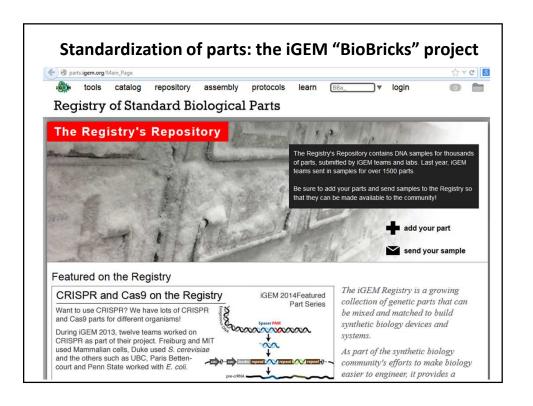


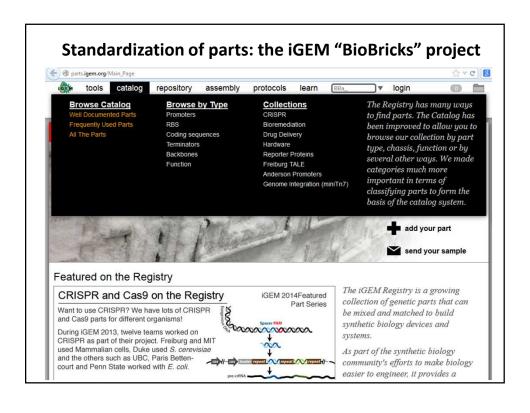
The behaviour can be predicted with a simple model
$$\frac{dN}{dt} = \frac{\text{cell growth rate}}{kN(1-N/N_m) - dEN} \text{ amount of killer protein growth}$$

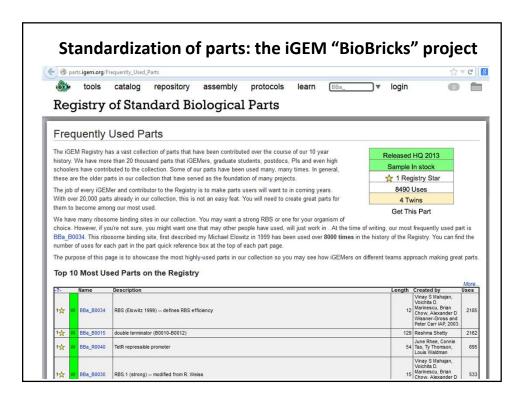
$$\frac{dE}{dt} = \frac{k_E A - d_E E}{killer protein} \text{ synthesis rate} \text{ degradation rate}$$

$$\frac{dA}{dt} = v_A N - d_A A$$

$$\frac{dA}{dt} = v_A N - d_$$







iGEM: A synthetic biology contest

from iGEM's web site

Can simple biological systems be built from standard, interchangeable parts and operated in living cells? Or is biology simply too complicated to be engineered in this way?

iGEM's broader goals include:

- To enable systematic engineering of biology
- To promote open & transparent development of tools for engineering biology
- To help construct a society that can productively apply biological technology

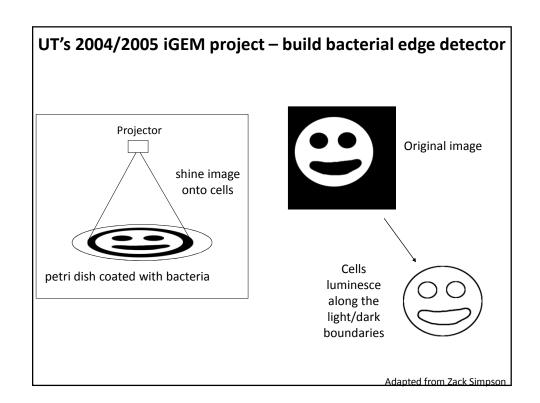
2004: MIT, UT, Princeton, Boston University, Cornell

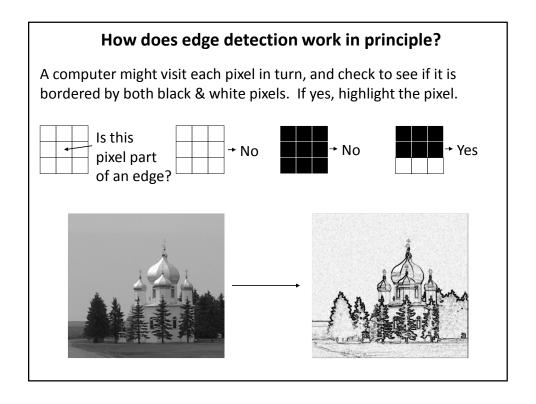
2005: 13 teams (the above + UK, Germany, more...)

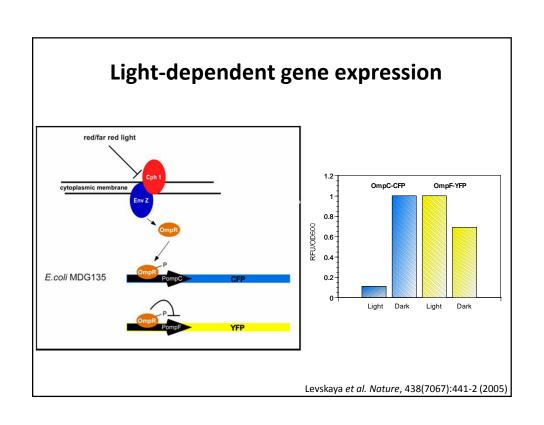
2006: 32 teams, incl. Japan/Latin America/Korea/India/more Europe

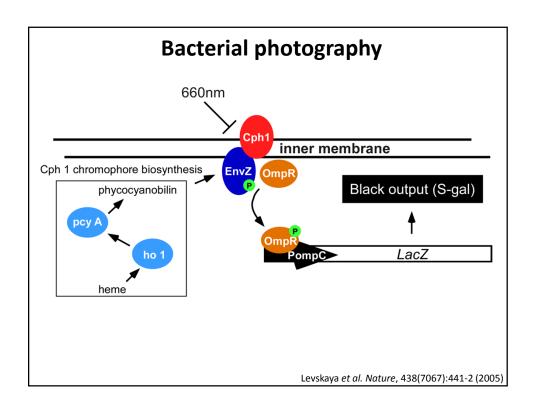
54 teams in 2007, 84 teams in 2008, 112 teams in 2009, 130 teams in

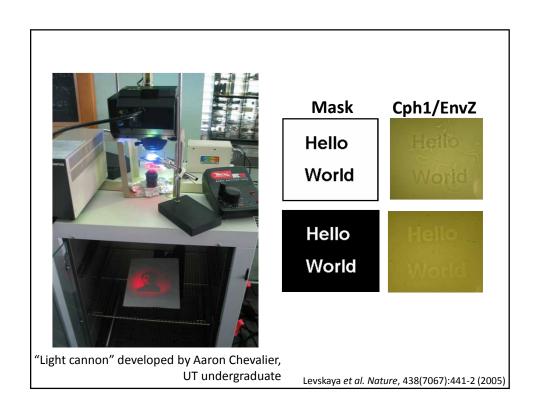
2010, 165 teams in 2011, and 245 teams in 2012 and 2013...

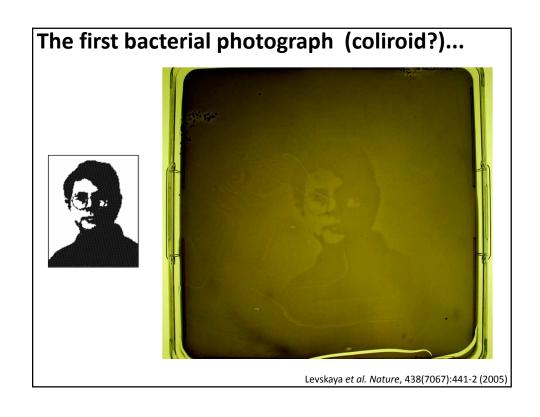


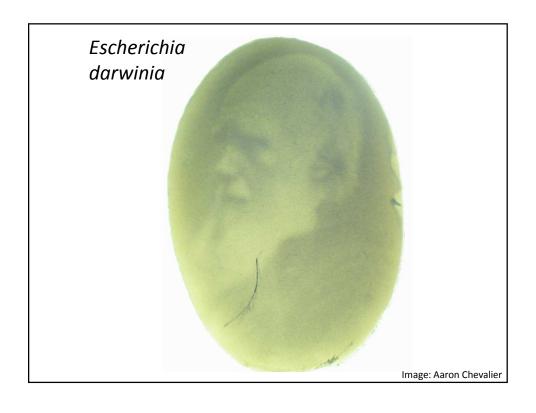


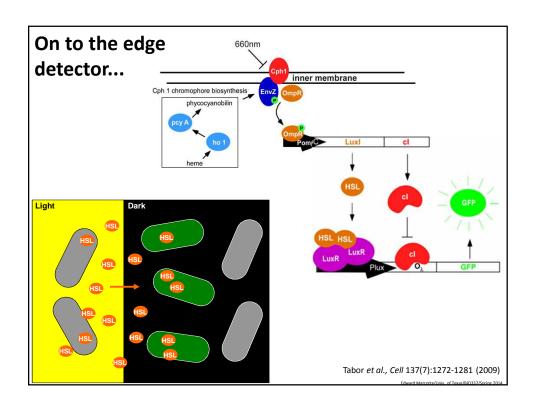


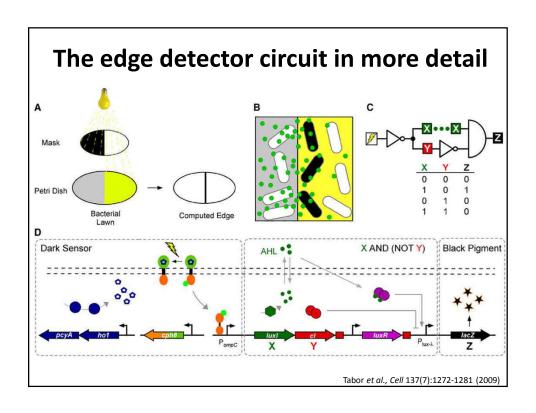


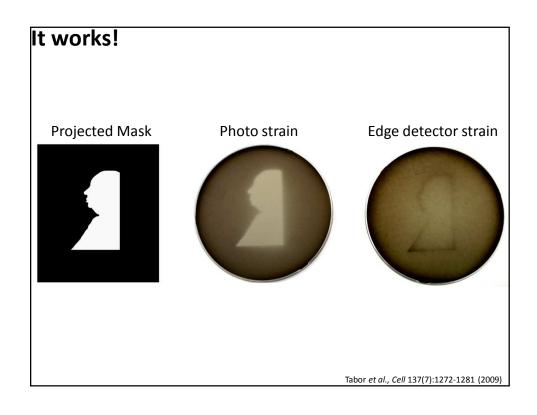


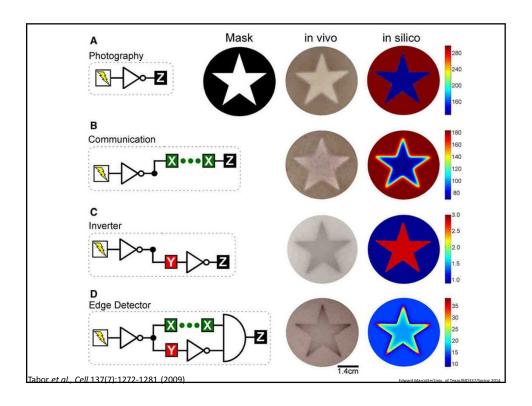












UT's 2012 iGEM project – build caffeine biosensor Synthetic Biology-Basic idea Decaffeination and Measurement of Caffeine Content by Addicted Escherichia coli with a Refactored N-Demethylation Operon from Block de novo guanine synthesis Convert caffeine to xanthine Pseudomonas putida CBB5 Erik M. Quandt, Michael J. Hammerling, Ryan M. Summers, Peter B. Otoupal, Ben Slater, Razan N. Altahhas, Aurko Dasgupta, James L. Bachman, Mani V. Subramanian, and Jeffrey E. Barrick, and Jeffrey E Addict E. coli bacteria to caffeine Xanthine methylxanthine (e.g. caffeine) Decaffeination pathway Guanine xanthine Xanthine salvage gpt Theophylline De novo guanine biosynthesis 11 steps PRPP --- IMP -----> XMP **-**➤ GTP and dGTP ——➤ RNA and DNA

