

Software for plant Synthetic Biology

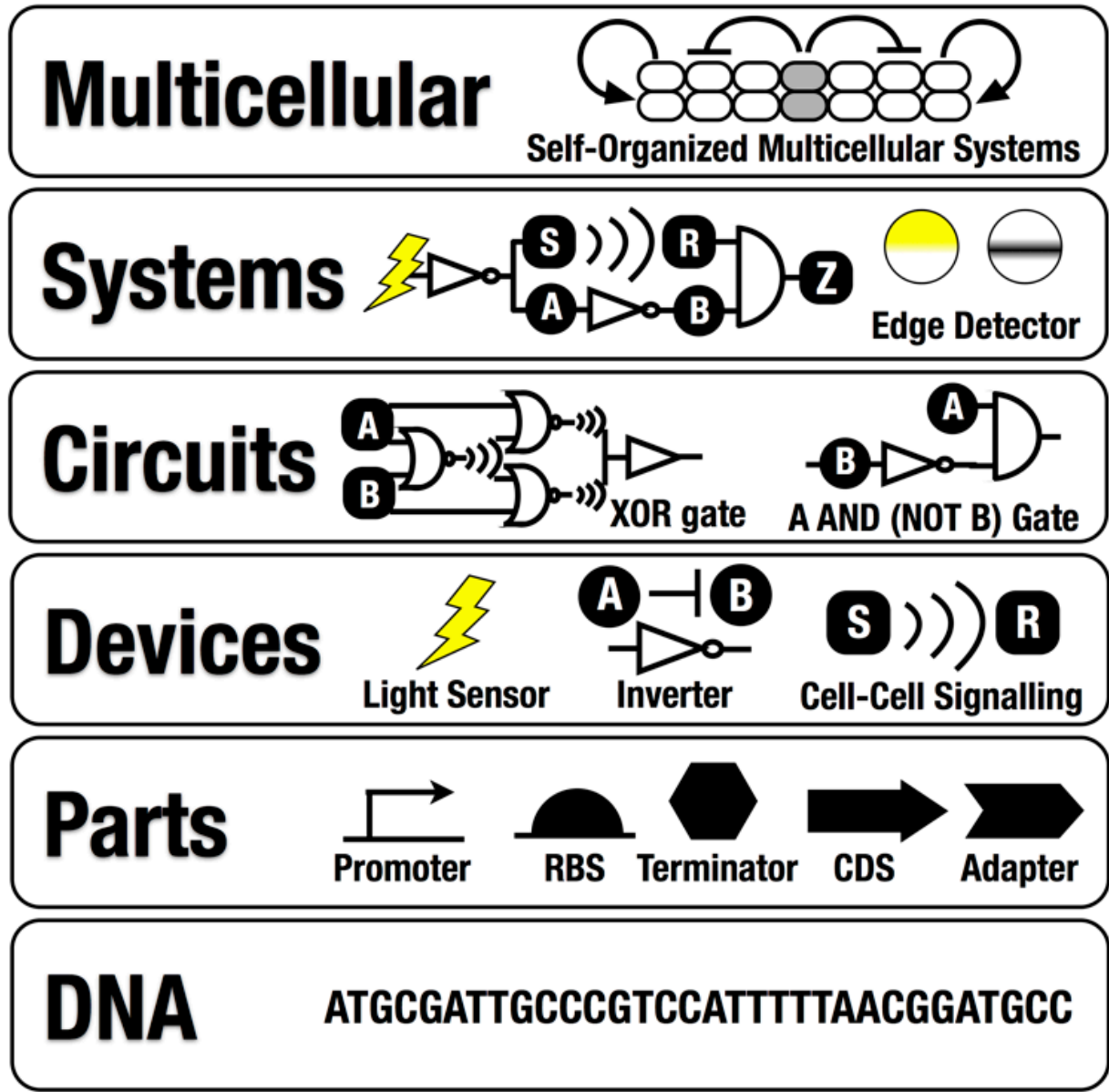
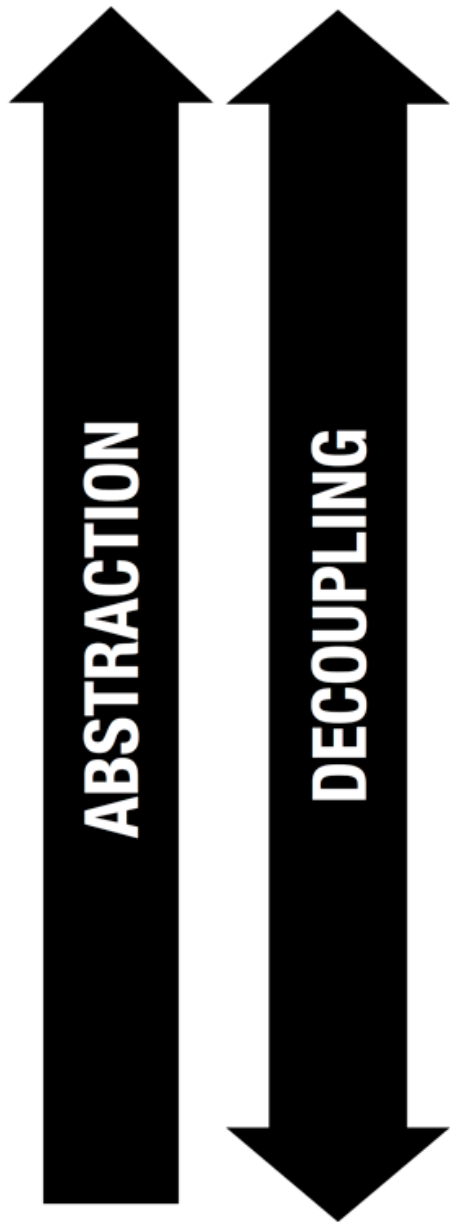
Jim Haseloff
University of Cambridge

www.haseloff-lab.org
www.synbio.org.uk

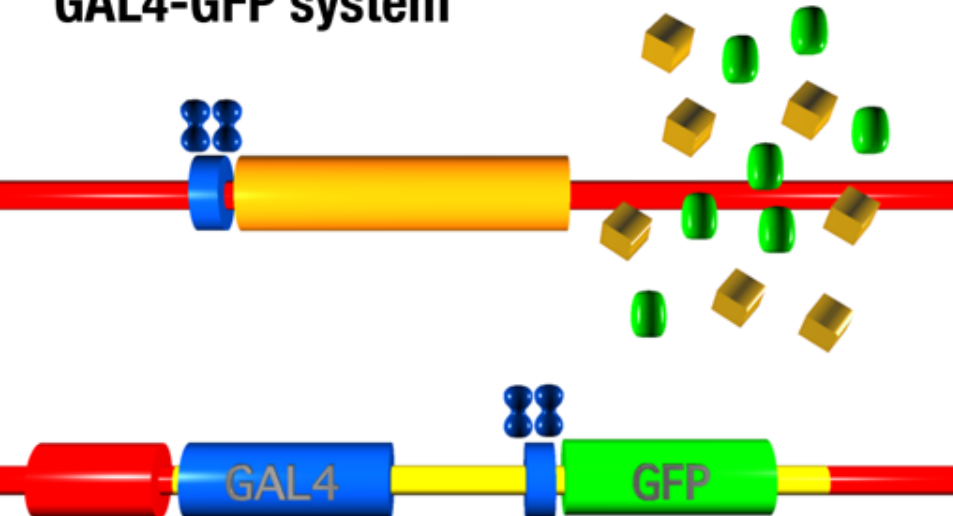
What is *Synthetic* Biology?

syn·thet·ic [sin-thet-ik] –adjective

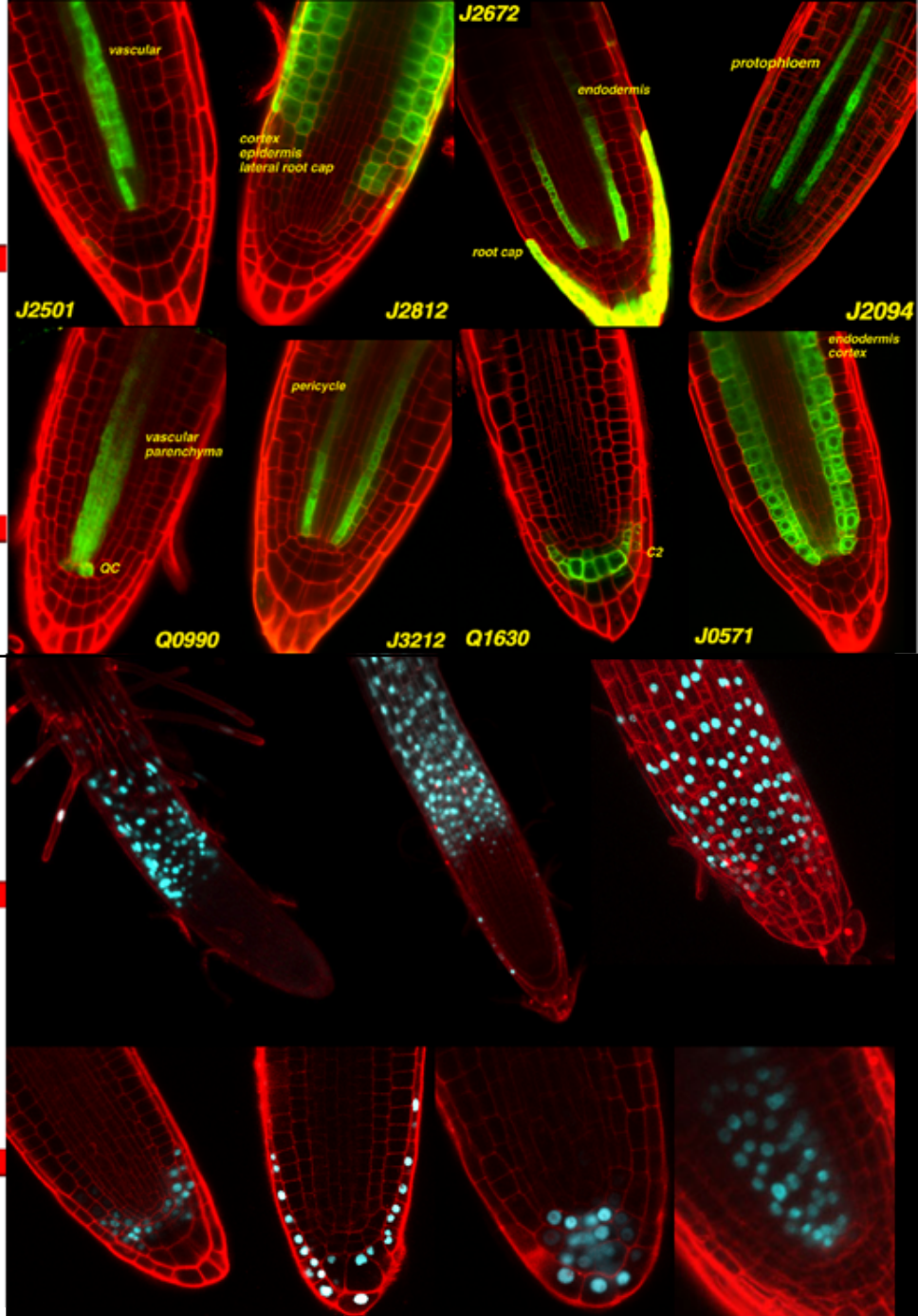
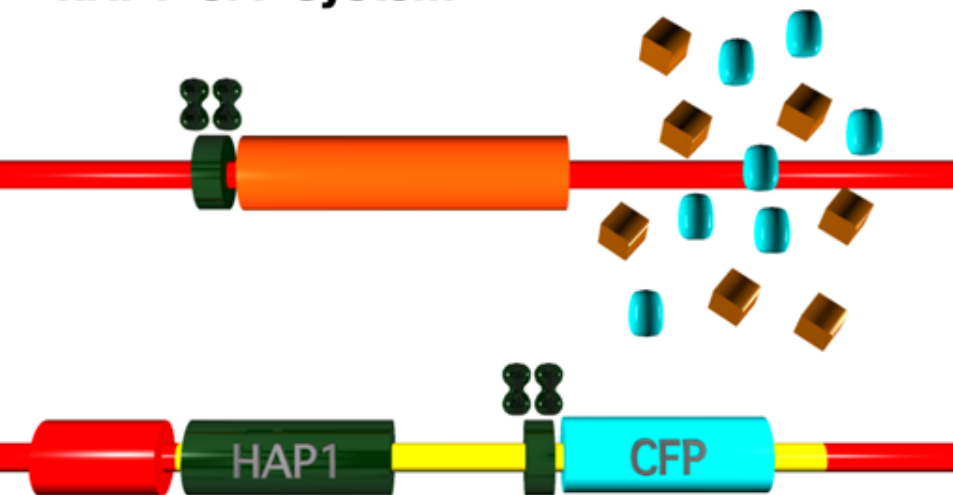
1. Prepared or made artificially, not of natural origin.
2. Relating to, or involving synthesis
(construction of a coherent whole from
separate elements)



GAL4-GFP system



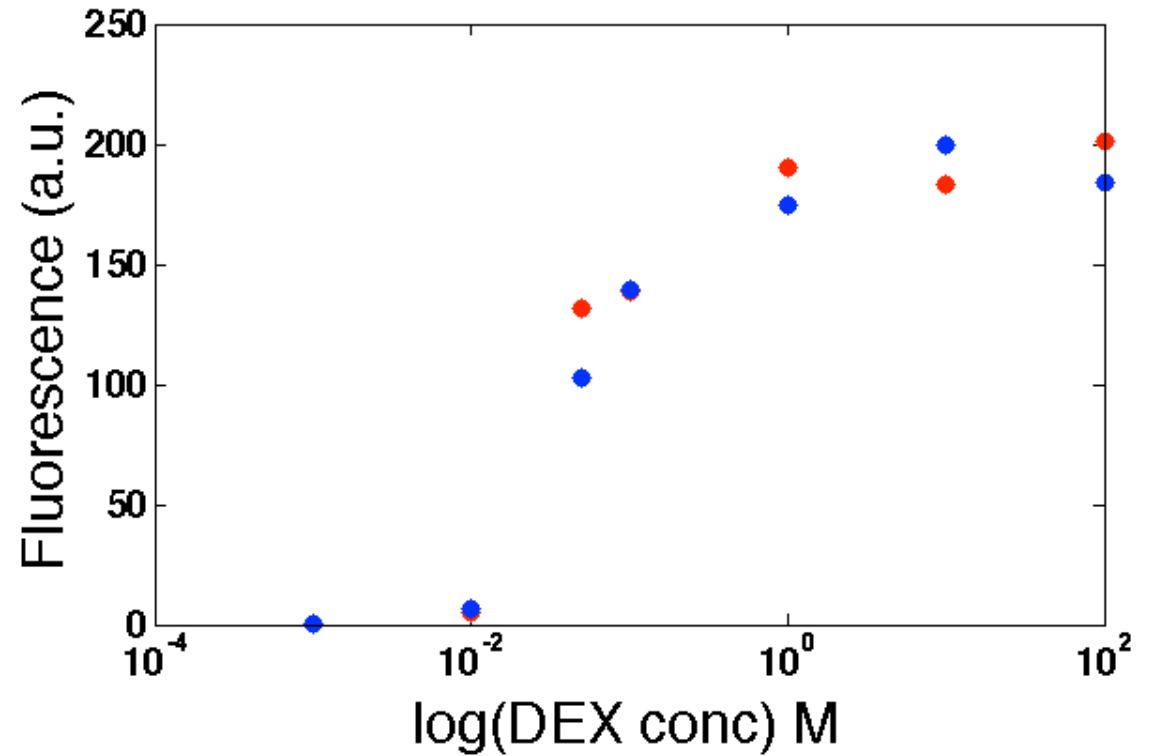
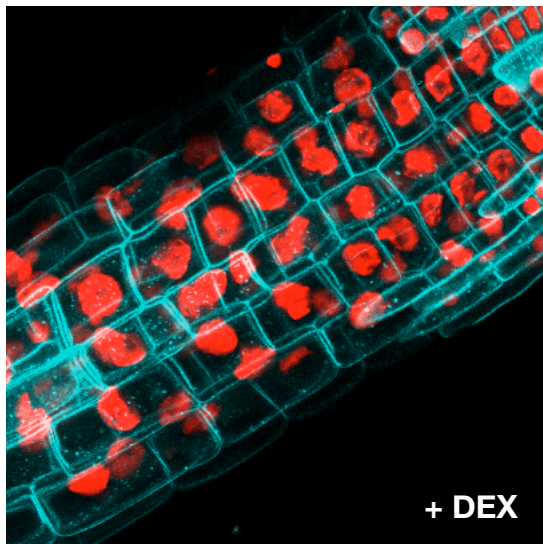
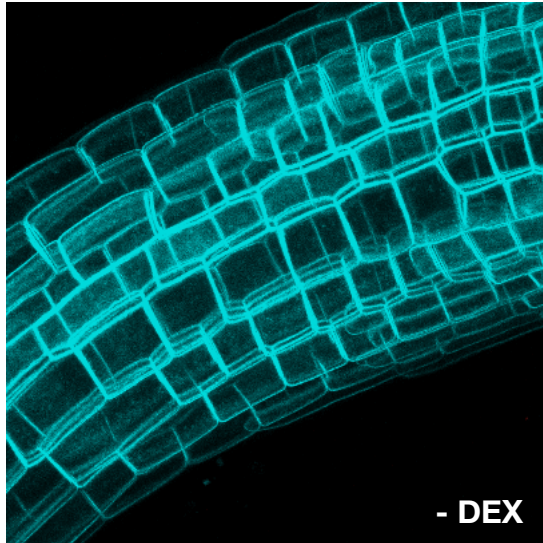
HAP1-CFP system



Triggers for gene expression

Temporal switching

Inducible control of HAPI-mediated gene expression

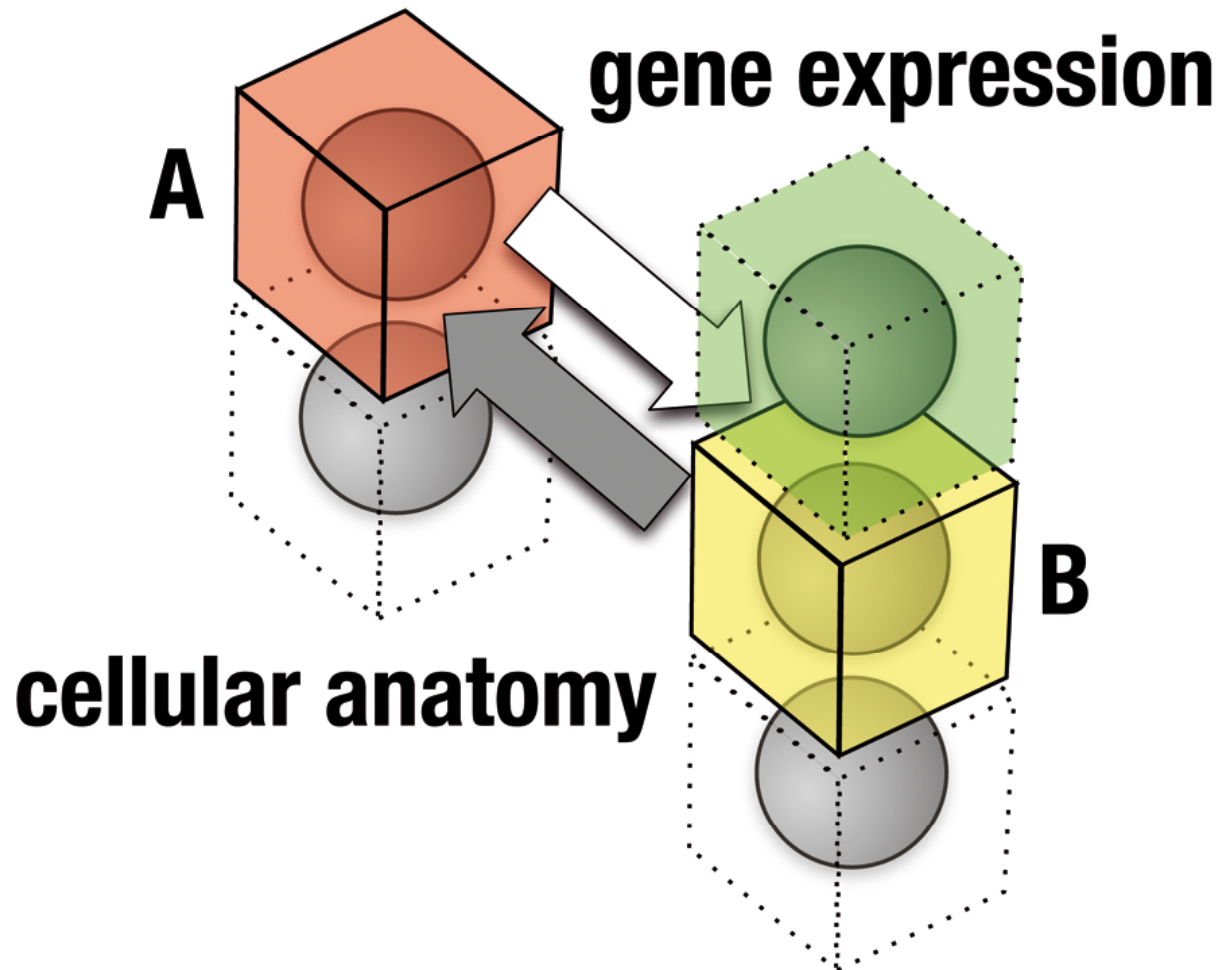


Testbed for synthetic biology in plants



Marchantia polymorpha

The properties of living systems are governed by parallelism and feedback



Design cycle for Synthetic Biology systems

1.

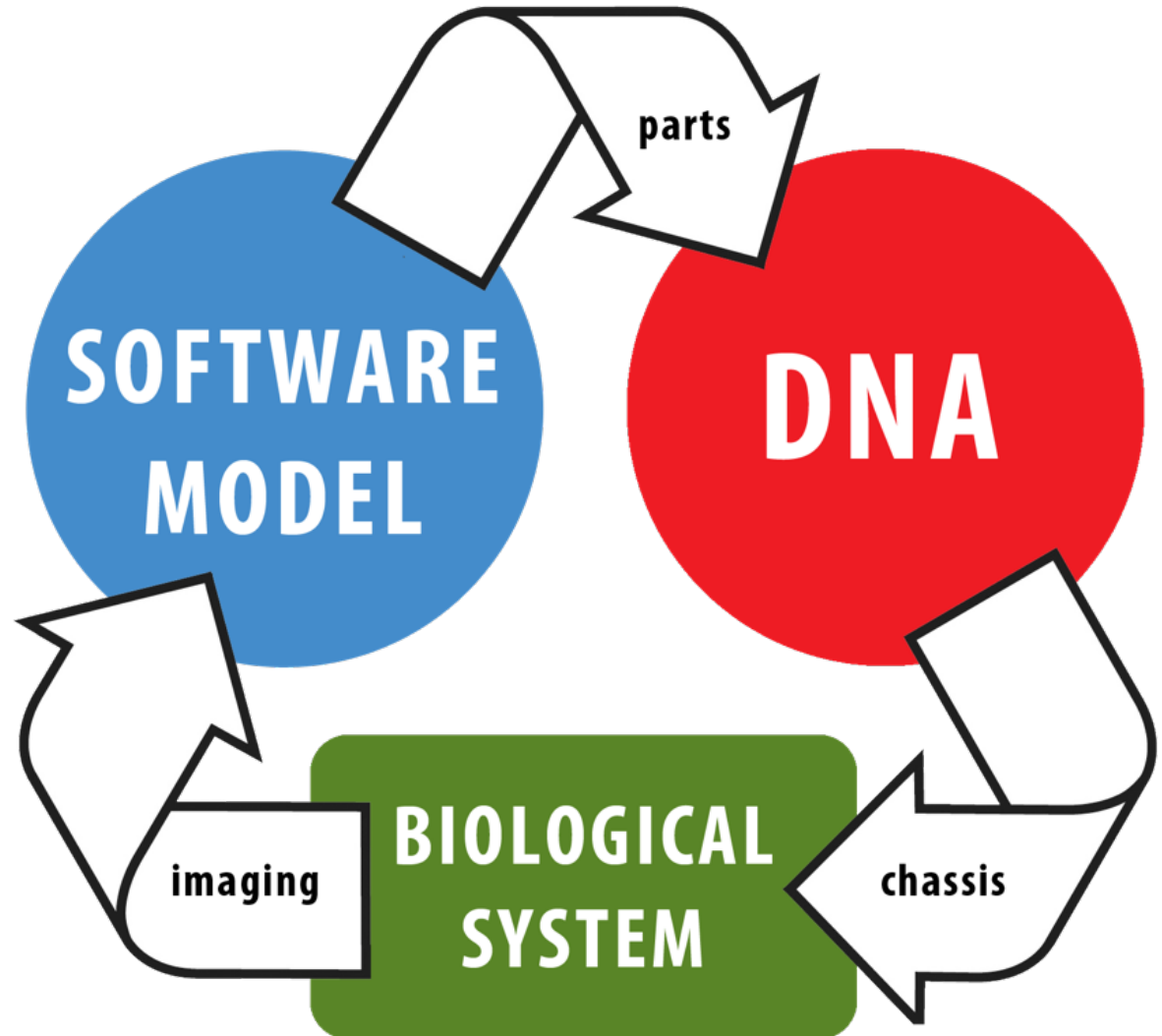
Specification and design of the system using computer models of the biological system

2.

Construction of genetic circuits using standard DNA parts and high throughput assembly techniques

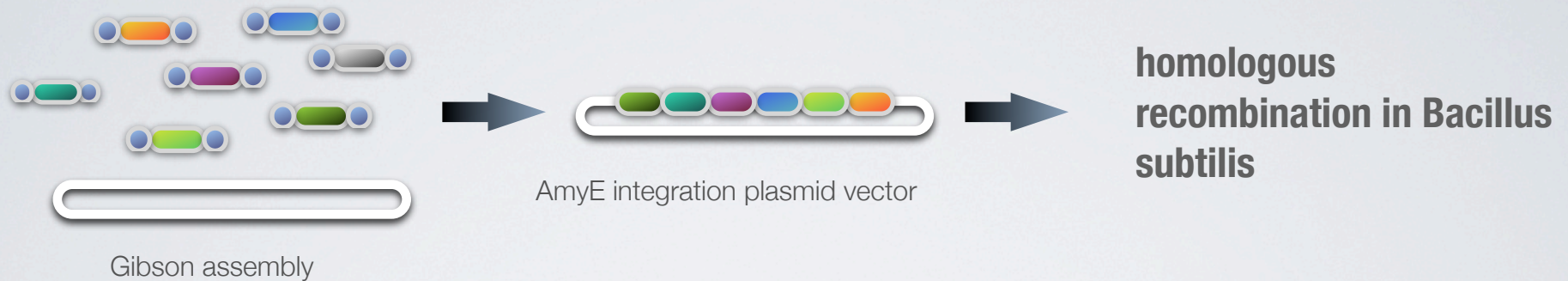
3.

Transformation of chassis and visualisation of gene expression, cell states and phenotype



High throughput chromosome engineering

Bacterial chromosome integration

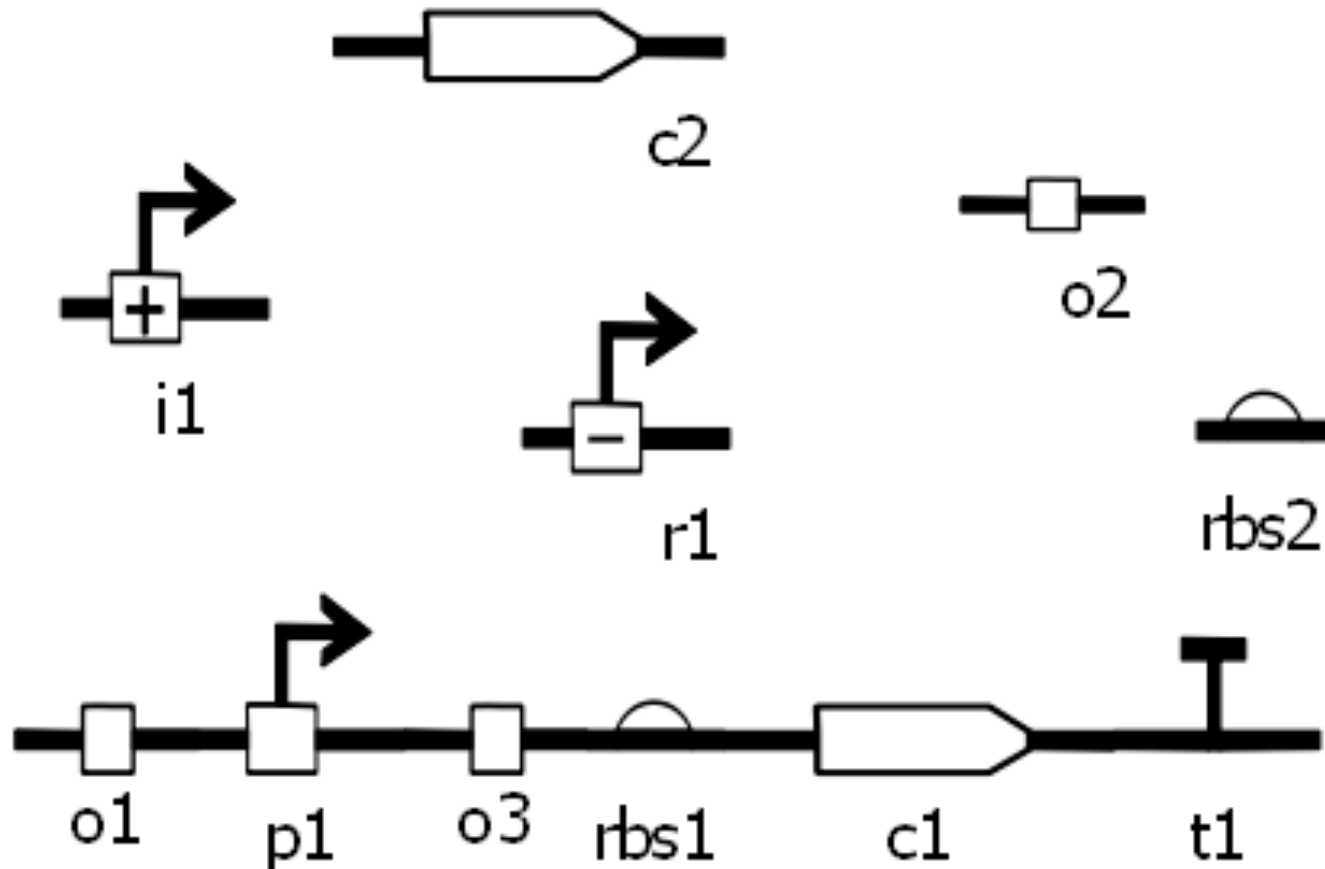


Plant chromosome integration



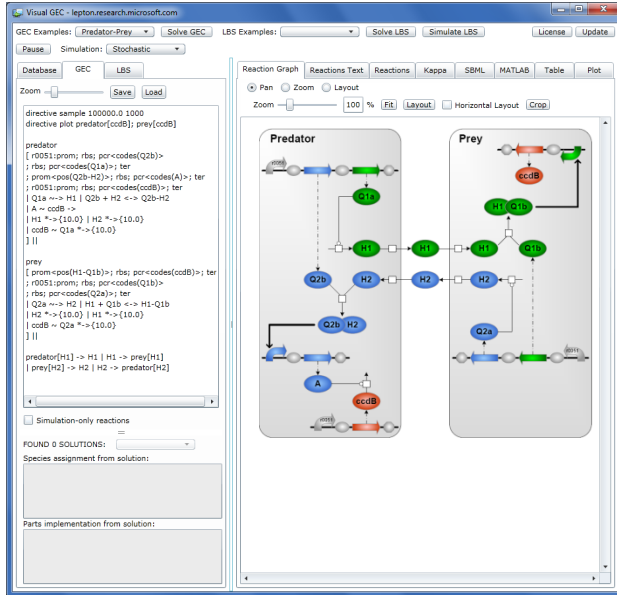
SBOL: symbolic description of DNA parts

Synthetic Biology Open Language: is a software standard for the electronic exchange of specifications and descriptions of genetic parts, genetic devices, genetic modules, genetic systems, and engineered genomes.

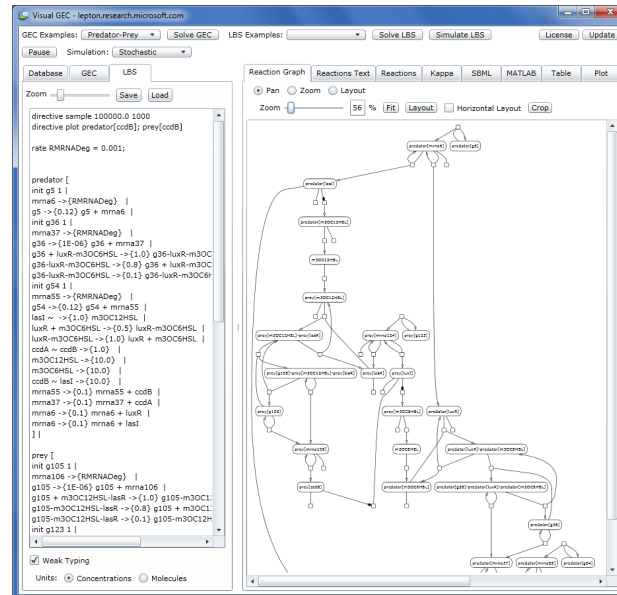


Software for compilation of DNA circuits

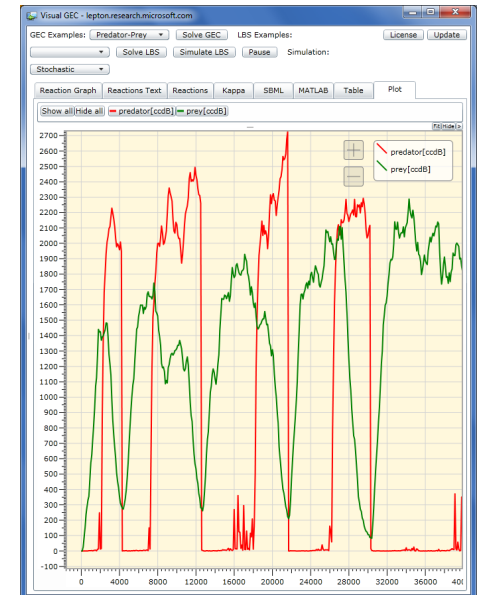
Step 1: Program device design



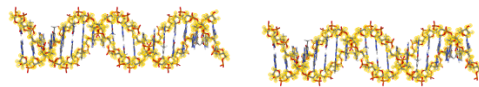
Step 2: Compile device behaviour



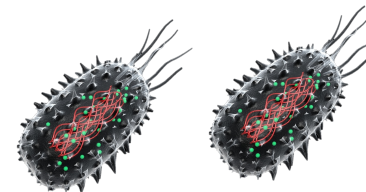
Step 3: Simulate device



Step 4: Compile device to DNA



Step 5: Insert DNA into cells



Genetic Engineering of Cells (GEC): a tool for programming cells

Microsoft Research Cambridge

CELLMODELLER

Software framework for modelling cell interactions

CellModeller version 4

CellModeller is a framework for modelling multicellular systems, including biophysics, regulatory dynamics, intercellular signalling, and rule-based behaviours. It is implemented using OpenCL, a parallel computation standard, allowing very large numbers of cells to be simulated

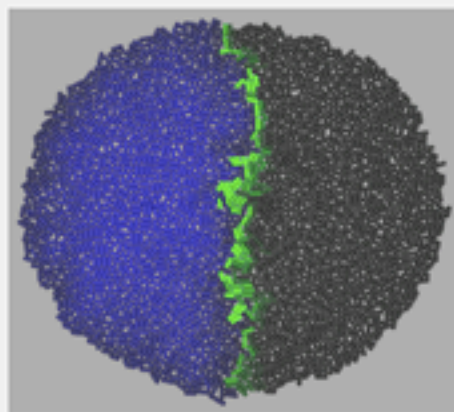


Morphogenesis is the process of creating and maintaining form through coordinated cell-division and growth. We have developed a software environment for modelling plant morphogenesis – an interactive virtual laboratory with a front-end called CellModeller4.

We have implemented cellular models of rod-shaped bacteria (3D) and plant tissues (2D). The bacterial model is a novel growing rigid-body method, described here. The plant model is a finite element method based on the work of Lionel Dupuy in the lab, and implemented using a matrix-free approach. The plant model is in the final testing stage, and will be released soon. The bacterial model is a constraint solver, and so can incorporate growth conditions like a flat substrate or microfluidic chamber by including additional constraints.

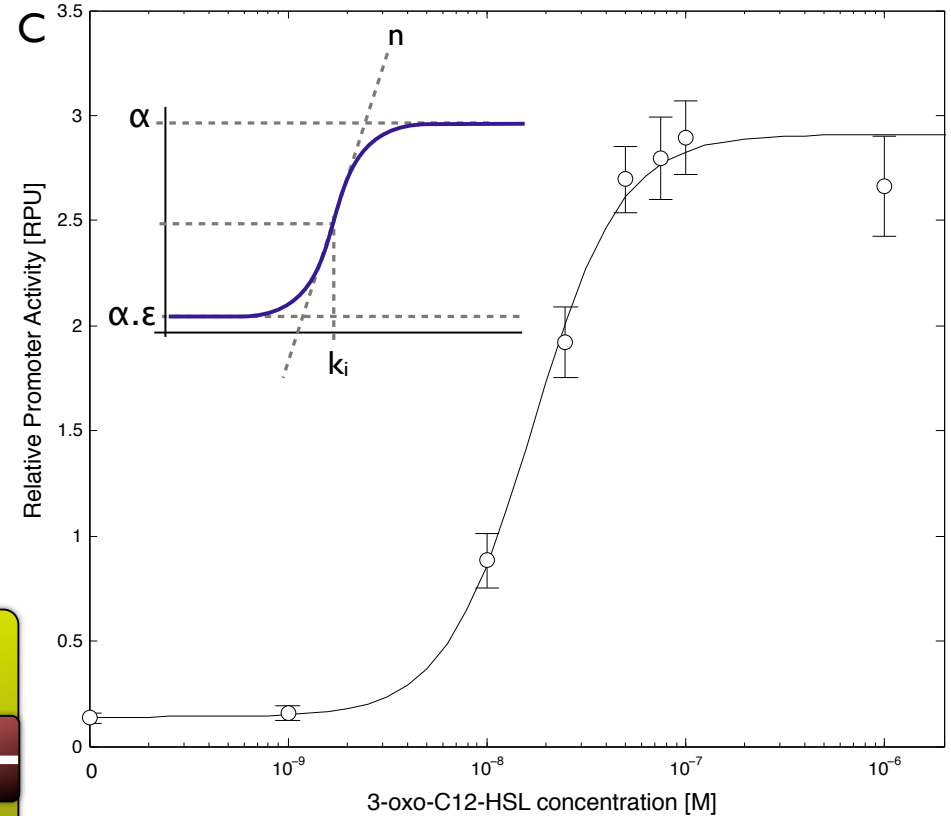
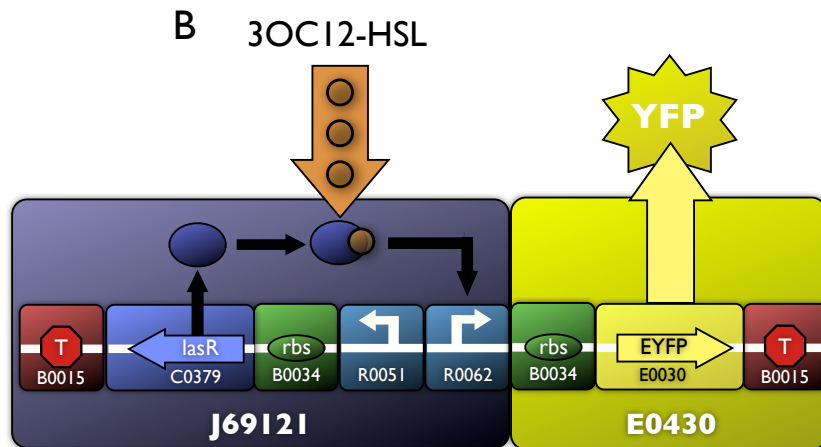
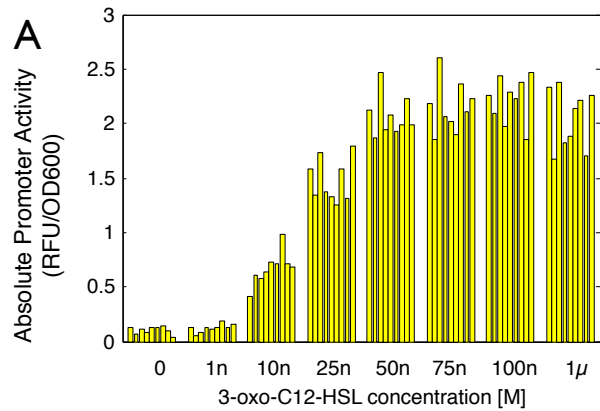
This is based on a spatial/mechanical model of cells coupled to an arbitrary user-defined set of reaction-diffusion systems and a genetic script. Essentially the system allows us to grow plant-like synthetic tissue which generates patterns of morphogens. The morphogens influence the behaviour, growth and division of the cells via the genetic script, forming a feedback loop between patterning and growth.

The 2D model is a matrix of linear walls controlled by the adjacent cells. Using the model we can design patterns, shapes and structures by constructing genetic scripts, effectively a kind of genetic sculpture. This is of great value in testing hypotheses about how genes and signalling molecules direct the growth and development of

[CellModeller](#)[Design](#)[Plant biophysics](#)[Cellular automata](#)[Genetic system](#)[Download software](#)[Cambridge-Microsoft](#)[Archiroot site](#)

Measurement of circuit behaviour *in vivo*

I22:pSB3K3CI Las Receiver



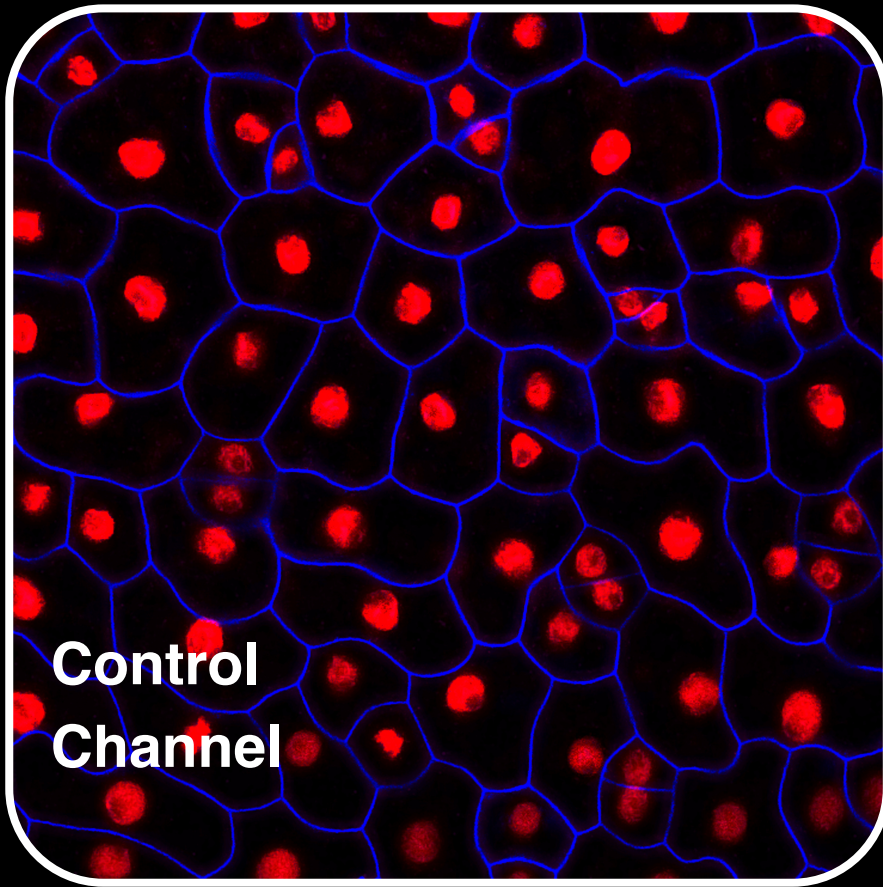
Characterisation of DNA parts for plants



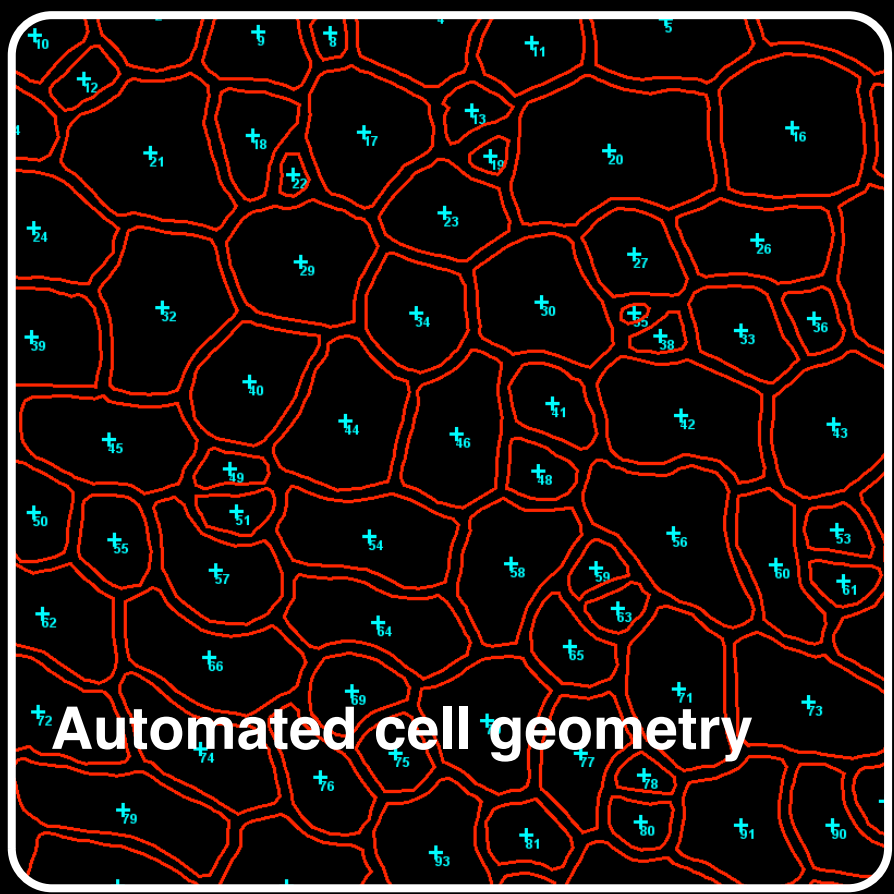
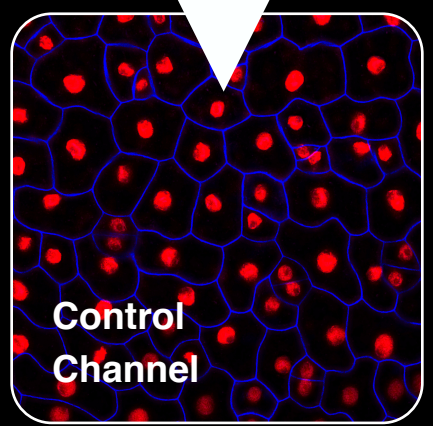
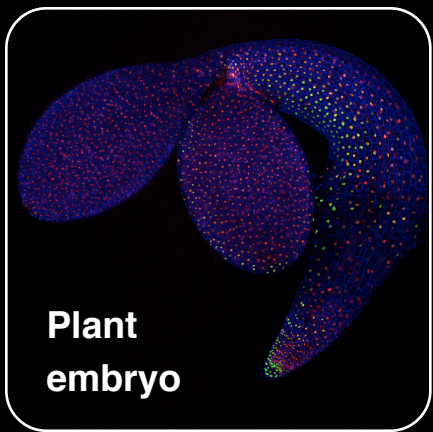
In planta cytometry

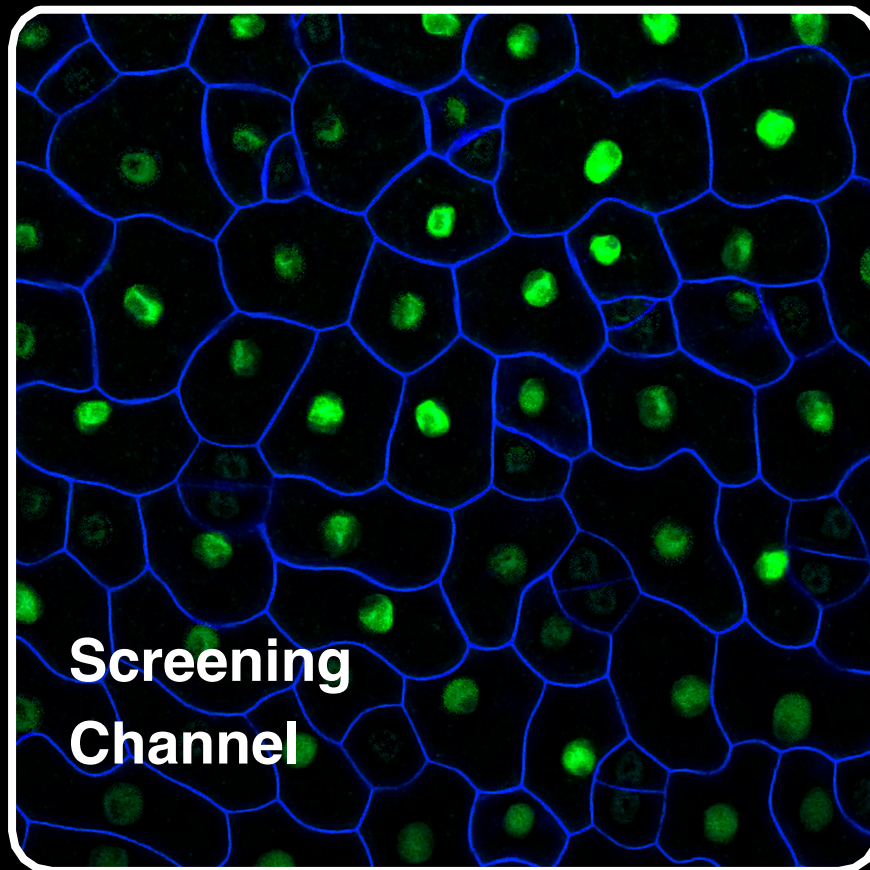
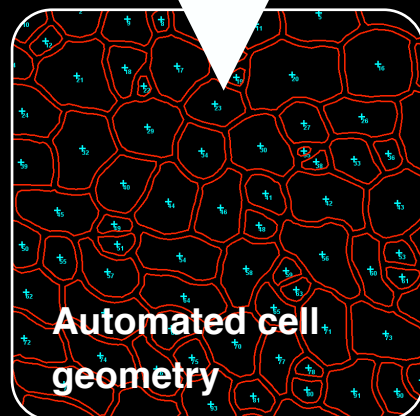
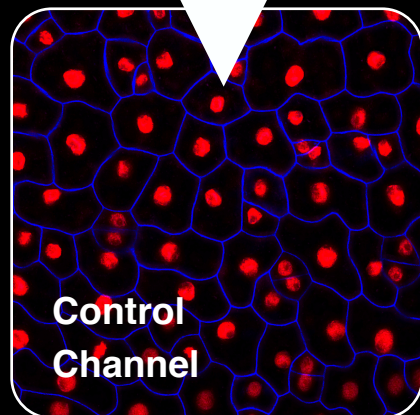
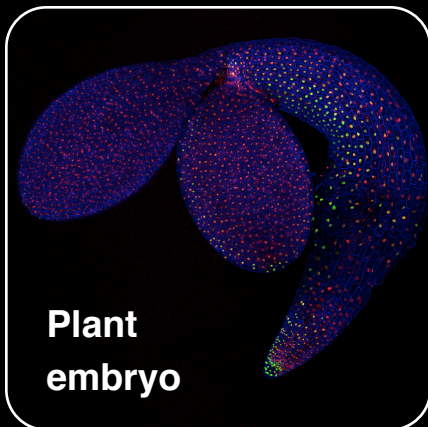


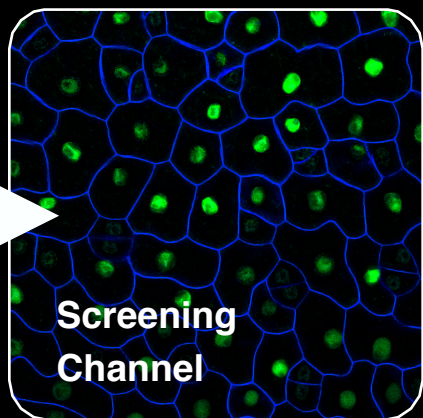
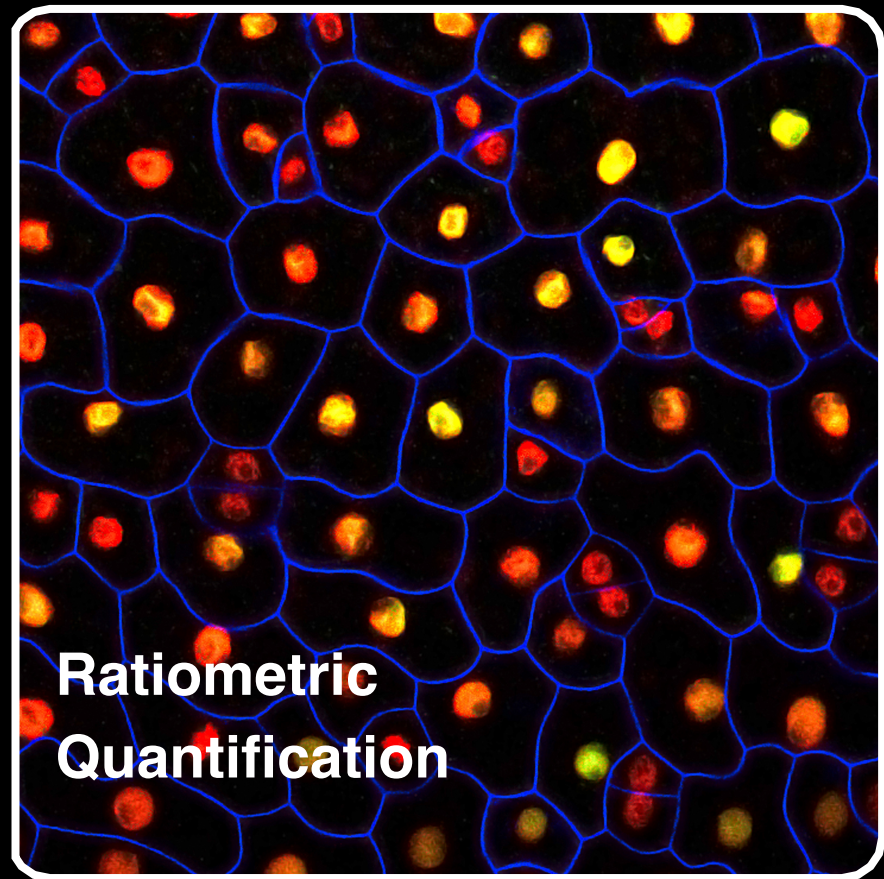
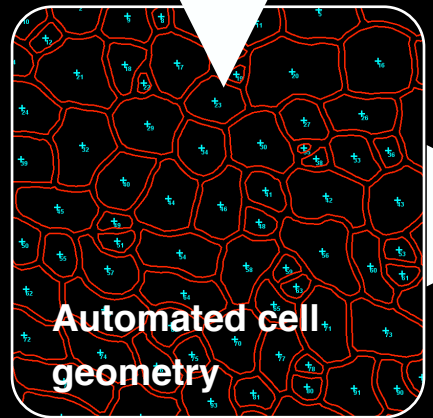
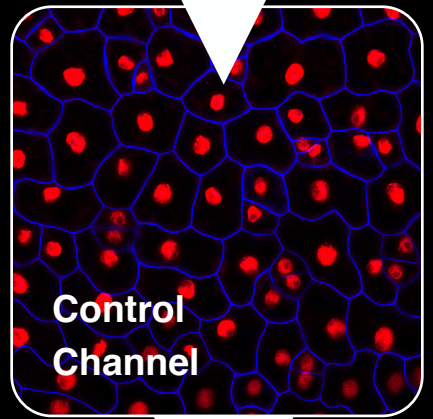
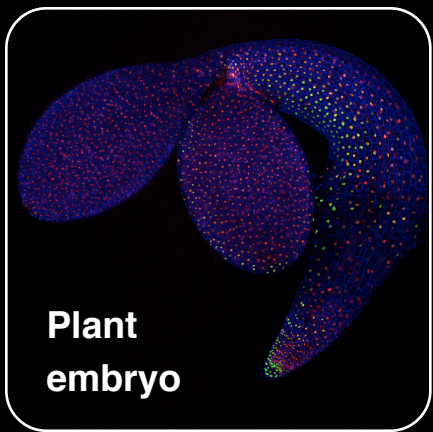
Plant embryo

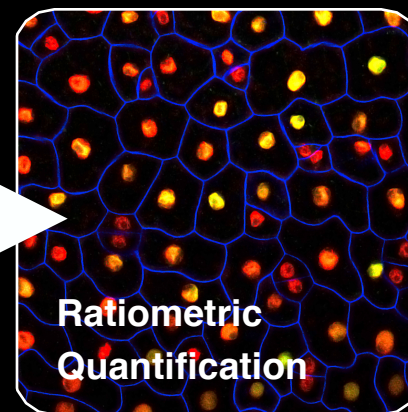
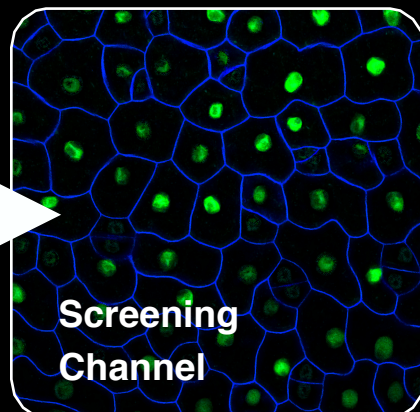
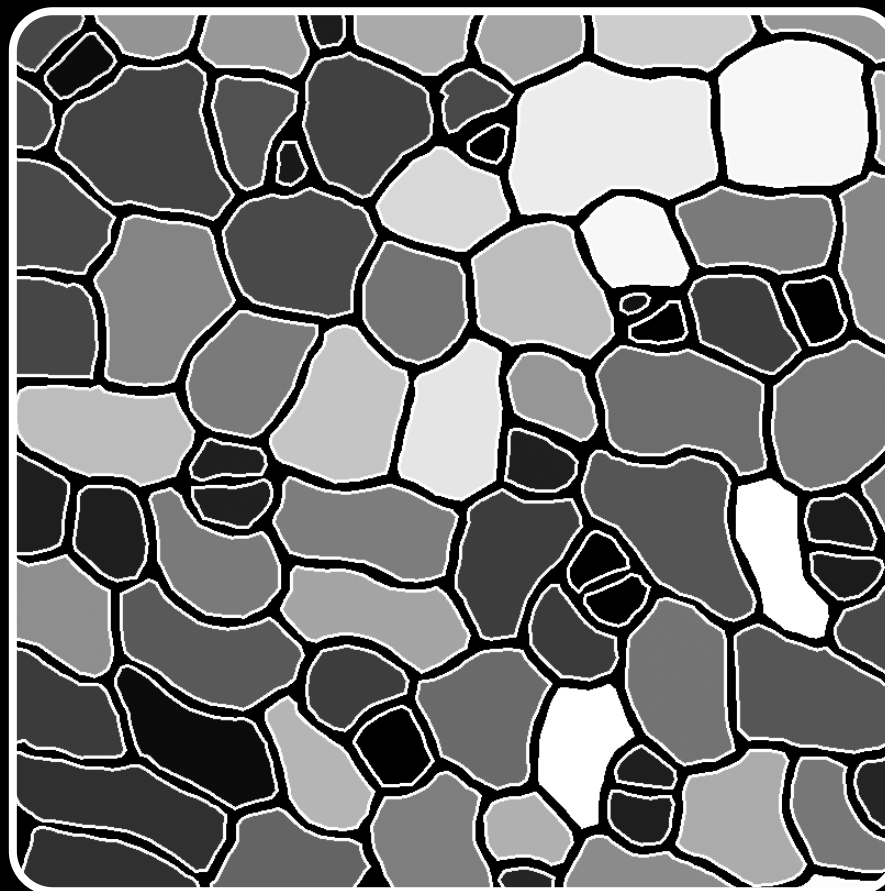
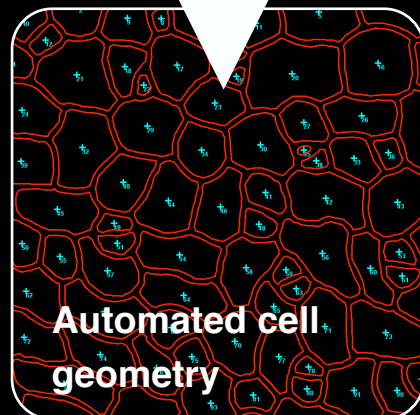
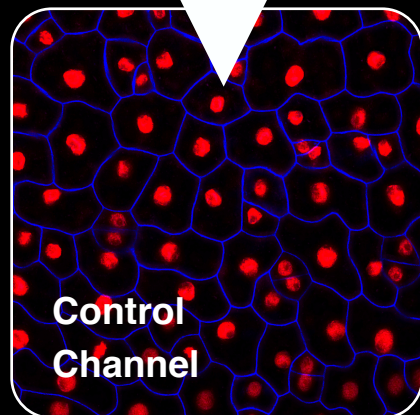
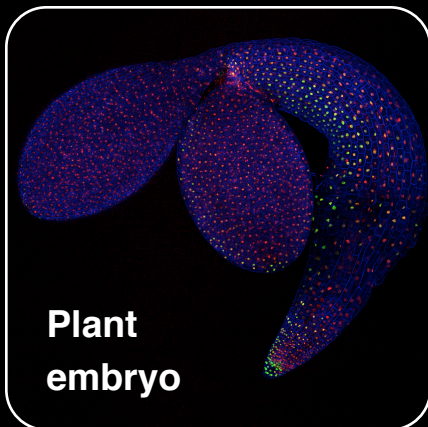


Control Channel



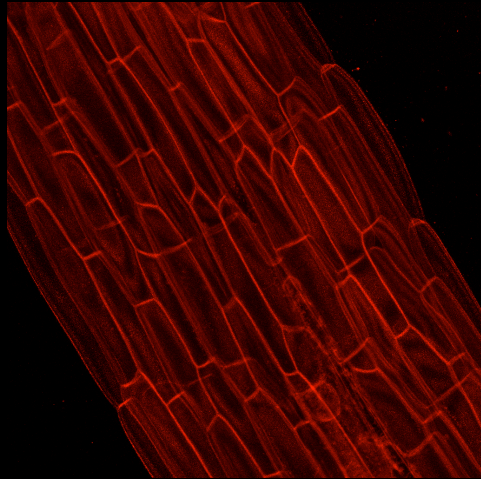






In planta cytometry II

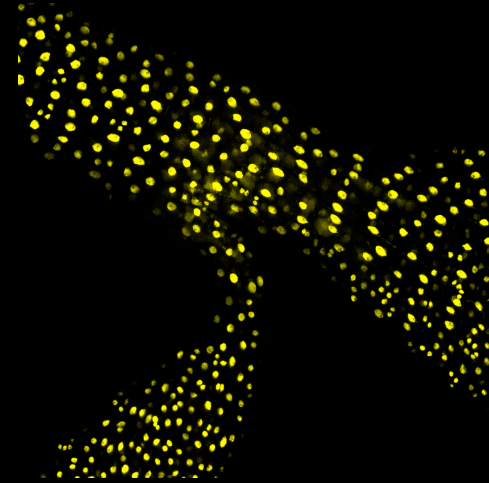
Sharing numbers



mKate2-LTI6b

TagRFP-T-LTI6b

sfGFP-LTI6b



H2B-mKate2

H2B-mTurquoise2

H2B-sfGFP

H2B-Venus

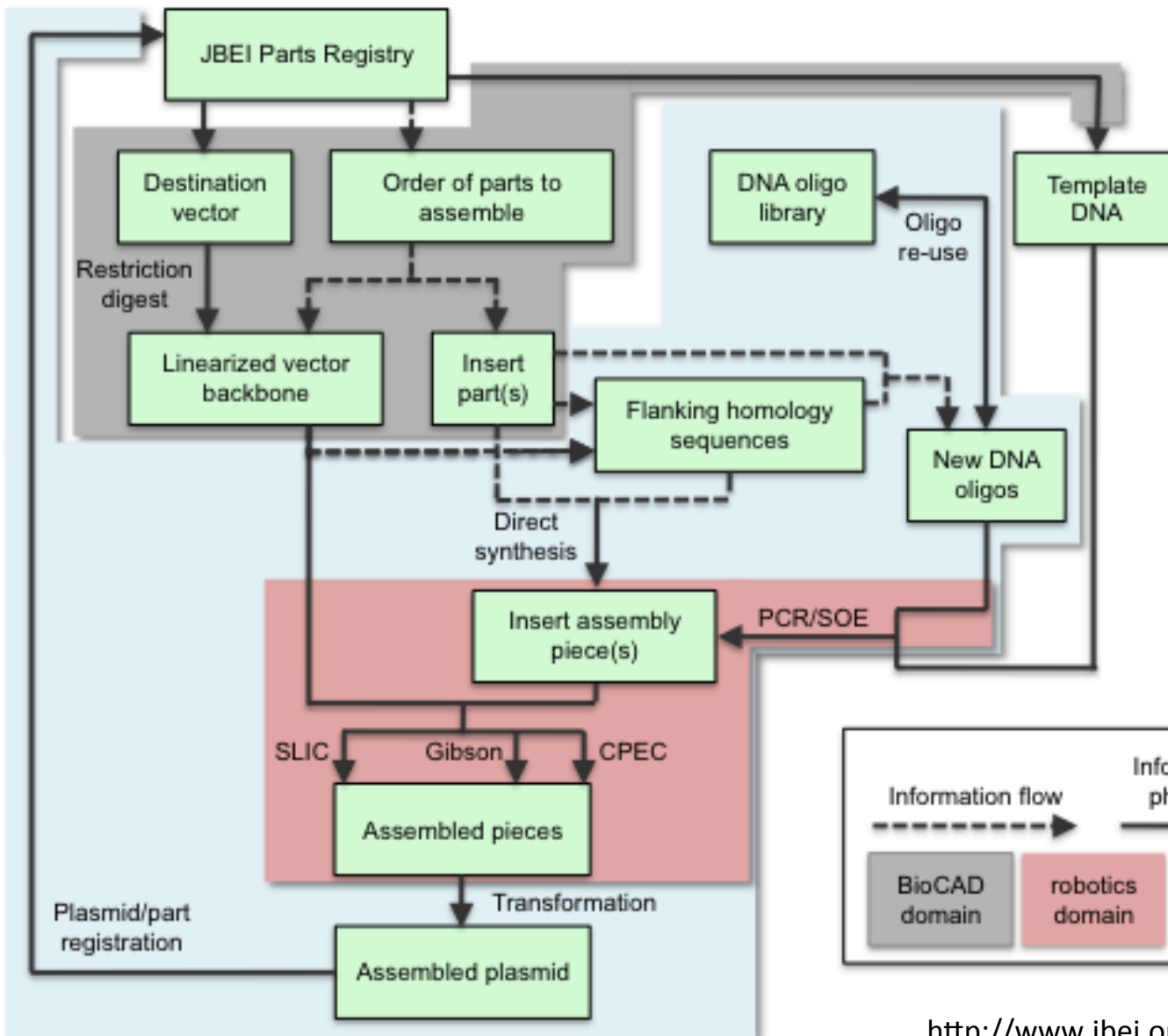
Reference Standard Promoters for Plants

UBQ10

Calculation of transcription rate

$$p^{ss} = \frac{\gamma_M (a + \gamma_I + \mu)(\gamma_F + \mu)[F]_{cell}^{ss}}{\rho_{an}}$$

Smart Registry for DNA parts



JBEI Inventory for Composable Elements (ICE)

Open source registry software for DNA parts

Nathan Hillson
Tim Hamm

Plasmid: pJ241:33268 violacein biobrick - Plantfab Registry

https://registry.plantfab.org/entry/view/8

PlantFab Cambridge

As of Sunday, December 25, 2011 there are 8 Entries available

Welcome, Jim Haseloff | Log Out

Home Collections Add new entry Bulk Import

Search Advanced Search

Plasmid: pJ241:33268 violacein biobrick

General Seq. Analysis (36) Report a Problem

Attachments (2) | Edit
QC_33268_64663.pdf
PM_33268_SEQ_64662.p...

Samples | Edit
No Samples

Permissions | Edit
Read Allowed: Everyone
Read and Write Allowed: Only you

Part ID: PLANTFAB_00008
Name: pJ241:33268 violacein biobrick
Markers: Backbone: pUC ori
Origin of Replication: pTR rrmB1 B2 T1 bxn terminator
Creator: Jim Haseloff
Status: Complete
Owner: Jim Haseloff
Created: Dec 14, 2011
Modified: Dec 14, 2011

Keywords:
Summary: The complete violacein biosynthesis operon synthesized by DNA2.0 for the Cambridge iGEM2009 team.

References:
Bio Safety: 1

IP Information:
Principal Investigator: Haseloff, Ajoka and Moklem
Funding Source:

Parameters:
Samples:

Download: (Original) | GenBank | FASTA Open in VectorEditor Delete

Done Read only 10135

Notes

registry.plantfab.org
DNA parts for Synthetic Biology in Plants and Microbes
Haseloff Laboratory - University of Cambridge

© JBEI Registry
All rights reserved.
Feature Request | Report a Bug | Feedback | API

Display page for a plasmid entry in the PlantFab Registry.

Import/export compatibility with Genbank sequence annotations.

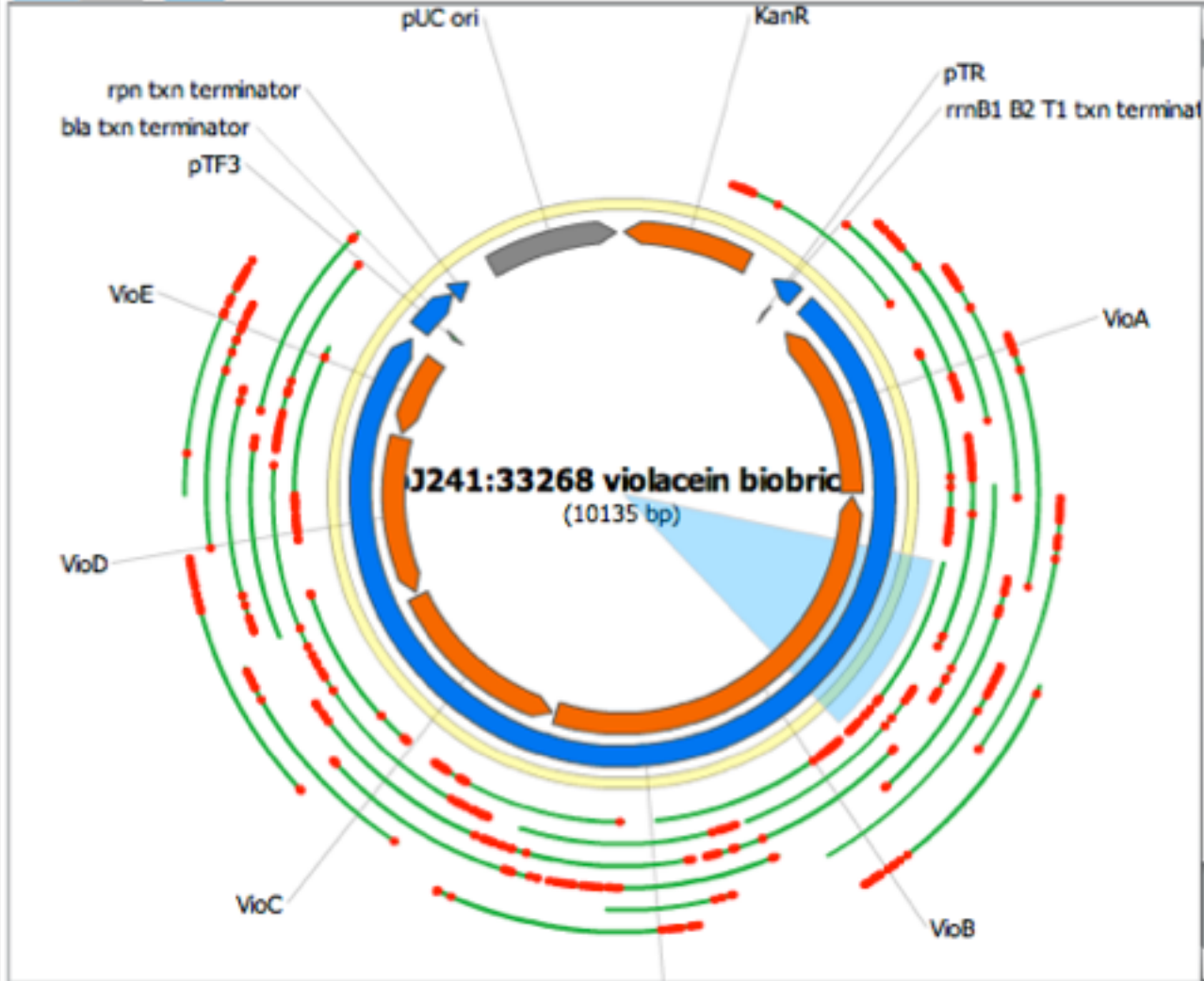
The screenshot displays the PlantFab Vector Editor web interface. The main window shows a circular plasmid map for '41:33268 violacein biotik (10135 bp)'. The map is annotated with various features, including promoters (pUC ori, pTF3, pTR), terminators (rpn txn terminator, bla txn terminator, rrnB1 B2 T1 txn terminator), and coding sequences (KanR, VioB, VioA, VioC, VioD, VioE). Restriction enzyme sites are also indicated around the map.

A 'Properties' window is open, showing the 'GenBank' tab. The window contains a table of features with the following data:

Name	Type	Position	Strand
Synthetic vio operon	misc_feature	1238 - 8623	+
pUC ori	rep_origin	9292 - 10095	+
rpn txn terminator	misc_feature	9007 - 9120	+
bla txn terminator	misc_feature	8700 - 9000	+
rrnB1 B2 T1 txn terminator	misc_feature	988 - 1162	-
pTF3	promoter	8791 - 8816	-
pTR	promoter	1063 - 1079	-
KanR	CDS	9 - 803	-
VioB	CDS	2549 - 5545	-
VioA	CDS	1275 - 2531	-
VioC	CDS	5569 - 6858	-
VioD	CDS	6882 - 8003	-
VioE	CDS	8027 - 8602	-

The 'Properties' window also includes a search field 'Enter Name or Type' and buttons for 'New', 'Edit', 'Remove', and 'OK'. The background shows the DNA sequence with annotations for features like pUC ori, pTR, KanR, and VioA-VioE.

Storage and automatic display of sequence trace files.



Name	Score	
1657501.ab1	1866	<input checked="" type="checkbox"/>
1669855.ab1	1938	<input checked="" type="checkbox"/>
1669858.ab1	1922	<input checked="" type="checkbox"/>
1674875.ab1	1760	<input checked="" type="checkbox"/>
1657494.ab1	1866	<input checked="" type="checkbox"/>
1657504.ab1	1802	<input checked="" type="checkbox"/>
1669837.ab1	1880	<input checked="" type="checkbox"/>
1669848.ab1	1886	<input checked="" type="checkbox"/>
1669849.ab1	1990	<input checked="" type="checkbox"/>
1657482.ab1	1892	<input checked="" type="checkbox"/>
1657490.ab1	1734	<input checked="" type="checkbox"/>
1669846.ab1	1984	<input checked="" type="checkbox"/>
1669850.ab1	1956	<input checked="" type="checkbox"/>
1669856.ab1	1944	<input checked="" type="checkbox"/>
1674872.ab1	1774	<input checked="" type="checkbox"/>
1657500.ab1	1810	<input checked="" type="checkbox"/>
1669838.ab1	1972	<input checked="" type="checkbox"/>
1669847.ab1	966	<input checked="" type="checkbox"/>
1669852.ab1	1910	<input checked="" type="checkbox"/>

Expected Trace 2876: GGTGACGGTCTGGTCGGTGTCTCGTTTGGCACTGTGGGGTCACTACAATGATTATCTGCGT
 Trace 28: GGTGNNGGTCTGGTCGGTGTCTCGTTTGGCACTGTGGGGTCACTACAATGATTATCTGCGT

ACCACCTTCAATCGTGCTCGTTGGGTCGACAGCGACCCGACGCGCCGTGACGCTGCACAA
 ACCACCTTCAATCGTGCTCGTTGGGTCGACAGCGACCCGACGCGCCGTGACGCTGCACAA

Built-in vector editing functions.

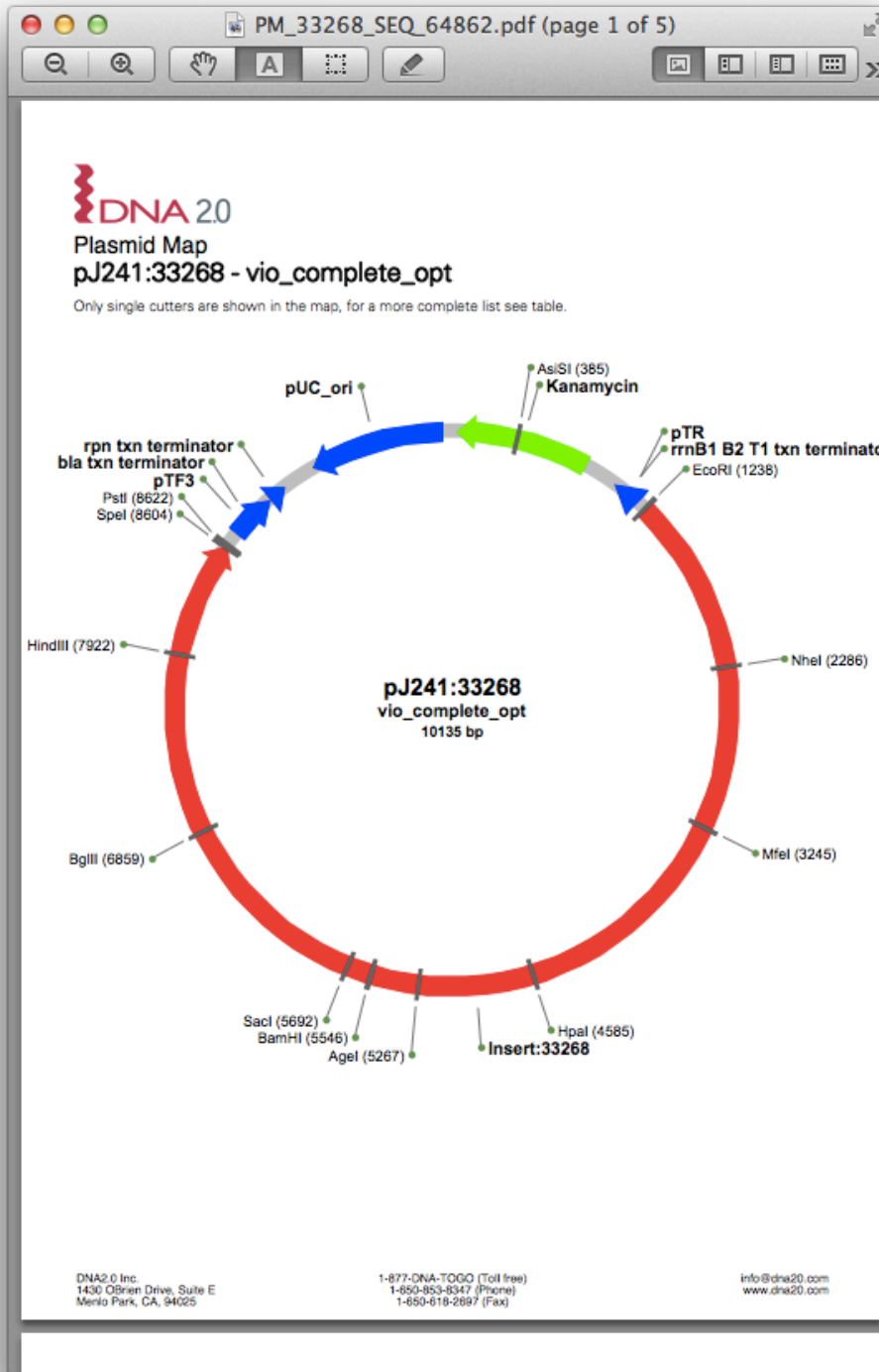
Browser address bar: <https://registry.plantfab.org/7wicket:bookmarkablePage=:org.jbel.ice.web.pages.VectorEditorPage&entryId=da34b8c5-3fe6-46f1-ae1>

File Edit View Tools Help

pJ241:33268 violacein biobrick
(10135 bp)

1	TAGAAAACT	CATCGAGCAT	CAAATGAAAC	TGCAATTTAT	TCATATCAGG	ATTATCAATA
61	CCATATTTT	GAAAAAGCCG	TTTCTGTAAT	GAAGGAGAAA	ACTCACCGAG	GCAGTTCCAT
121	AGGATGGCAA	GATCCTGGTA	TCGGTCTGCG	ATTCCGACTC	GTCCAACATC	AATACAACCT
181	ATTAATTTCC	CCTCGTCAAA	AATAAGGTTA	TCAAGTGAGA	AATCACCATG	AGTGACGACT
241	GAATCCGGTG	AGAATGGCAA	AAGTTTATGC	ATTTCTTTCC	AGACTTGTTT	AACAGGCCAG
301	CCATTACGCT	CGTCATCAAA	ATCACTCGCA	TCAACCAAAC	CGTTATTCAT	TCGTGATTGC
361	GCCTGAGCGA	GGCGAAATAC	GCGATCGCTG	TAAAAAGGAC	AATTACAAC	AGGAATCGAG
421	TGCAACCGGC	GCAGGAACAC	TGCCAGCGCA	TCAACAATAT	TTTCACCTGA	ATCAGGATAT
481	TCTTCTAATA	CCTGGAACGC	TGTTTTTCCG	GGGATCGCAG	TGGTGAGTAA	CCATGCATCA
541	TCAGGAGTAC	GGATAAAATG	CTTGATGGTC	GGAAAGTGGA	TAAATCCGT	CAGCCAGTTT
601	AGTCTGACCA	TCTCATCTGT	AACATCATTG	GCAACGCTAC	CTTTGCCATG	TTTCAGAAAC
661	AACTCTGGCG	CATCGGGCTT	CCCATAACAAG	CGATAGATTG	TCGCACCTGA	TTGCCCGACA
721	TTATCGCGAG	CCCATTTATA	CCCATATAAA	TCAGCATCCA	TGTTGGAATT	TAATCGCGGC
781	CTCGACGTTT	CCCGTTGAAT	ATGGCTCATA	TCTTCTCTTT	TTCAATATTA	TTGAAGCATT
841	TATCAGGGTT	ATTGTCTCAT	GAGCGGATAC	ATATTTGAAT	GATTTTAGAA	AAATAAACAA
901	ATAGGGGTCA	GTGTTACAAC	CAATTAACCA	ATTCTGAACA	TTATCGCGAG	CCCATTATA

Writable | - | 10135



Documents can be attached to sequence entries.

www.plantfab.org

Registry of DNA parts for Plants



DNA parts for the fabrication of new genetic systems in plants and related microbes.

(construction in progress, Dec 2011)



Registry of DNA parts

A new resource for sharing details of DNA parts for use in Synthetic Biology experiments in plants.

The recent development of **genome-scale DNA assembly techniques** allows fast, efficient multiplex fabrication of large-scale DNA assemblies - with no need for restriction enzymes or low-efficiency ligation reactions. This places new demands on the availability of reliable and well characterised DNA parts for assembly.

This site contains a registry of DNA parts dedicated to use in plant and related microbial systems. The formation of the registry was driven by the lab's need to keep our own DNA parts in an orderly manner - many of the parts will be "work-in-progress", and not accessible without password. Details of published DNA constructs will be available.

The PlantFab Registry is based on the **Inventory for Composable Elements (ICE)**, developed as an open source registry for biological parts by the Joint Bio Energy Institute (JBEI), UC Berkeley (with many thanks to Tim Ham



biofab

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Search

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[Home](#)

about the biofab

The BIOFAB: International Open Facility Advancing Biotechnology (BIOFAB) was founded in December 2009 as the world's first biological design-build facility. This professionally staffed public-benefit facility was initiated by a grant from the National Science Foundation (NSF) and is led by bioengineers from UC Berkeley and Stanford University. The BIOFAB is operated in partnership with [Lawrence Berkeley National Laboratory \(LBNL\)](#), the [BioBricks Foundation \(BBF\)](#), and the [Synthetic Biology Engineering Research Center \(SynBERC\)](#).

BioFab projects will be designed to produce broadly useful collections of standard biological parts that can be made freely available to both academic and commercial users, while also enabling the rapid design and prototyping of genetic constructs needed to support specific needs of partner efforts such as SynBERC Testbeds. The BioFab will thus also represent the first significant focused investment in the development of open technology platforms underlying and supporting the next generation of biotechnology. Once fully operational the BioFab facility will be capable of producing tens of thousands of professionally engineered, high quality standard biological parts each year.

Read the full announcement here:

http://www.berkeley.edu/news/media/releases/2010/01/20_biofab_synthetic_biology.shtml

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<http://www.biofab.org>



The BioBricks Foundation

- Develop and implement legal strategies to ensure that BioBrick™ standard biological parts remain freely available to the public.
- Support the development of open technical standards that define BioBrick™ standard biological parts.
- Develop and provide educational and scientific materials to allow the public to use and improve existing BioBrick™ standard biological parts, and contribute new BioBrick™ standard biological parts.



BioBricks
FOUNDATION

Biotechnology in the public interest

biobricks.org


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 BioBricks Foundation
@biobricks

The BioBricks Foundation SB6 Conference registration is now open! Space is limited. <http://t.co/LLLOCaVL>
25 Jan 2013 · reply · retweet · favorite

Thanks, @GerdMoeBehrens for a really important paper:<http://t.co/FFdN5oTH>
25 Jan 2013 · reply · retweet · favorite

Tom Knight, BBF Founding Director, Emeritus, profiled in "New Scientist" on @biobricks, @igem & craziest synbio: <http://t.co/FB8yWXpIF>
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BIOFAB success to continue in partnership with the BBF. [Learn more »](#)



Synthetic Biology 6.0 Conference



BioBrick™ Public Agreement



Stanford BIOFAB



The BBF's mission is to ensure that the engineering of biology is conducted in an open and ethical manner to benefit all people and the planet.

We believe fundamental scientific knowledge belongs to all of us and must be freely available for ethical, open innovation. This is a new paradigm.

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LATEST FROM THE BBF BLOG

March 20, 2013
Free! Reliable! Parts! Use 'Em! Courtesy Drew Endy
»

January 18, 2013
Genspace Gives Great TED Talk »

January 17, 2013
SB6.0 Conference Registration is Now Open! »

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 The Synthetic Biology Network 

Applications

- **open source and free-to-operate tools**
- **open standards to promote innovation**
- **international scientific exchange**



Cambridge website for Synthetic Biology resources

Compiled by Jim Haseloff at the University of Cambridge.

This site contains details of recent papers and activity in Synthetic Biology, with particular emphasis on: (i) development of standards in biology and DNA parts, (ii) microbial and (iii) plant systems, (iv) research and teaching in the field at the University of Cambridge, (v) hardware for scientific computing and instrumentation, (vi) tools for scientific productivity and collected miscellany.

Similar to the Cambridge-based Raspberry Pi and OpenLabTools initiatives, we promote the use of low cost and open source tools - in our case for use in biological engineering.



SYNBIO CALENDAR

11
MAY

First International Mammalian Synthetic Biology Workshop

12:00 AM to 12:00 AM

Recognizing the fast emergence and potential significance of this field, the aim of this workshop is to bring together practitioners of mammalian synthetic biology together with experts from other relevant

20
MAY

Introduction to opportunities in plant synthetic biology

07:00 PM to 07:00 PM

This meeting will introduce and showcase the many and varied applications of synthetic biology at the molecular, cell, and whole plant level to introduce synthetic biology to a broad UK plant science base.

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RESEARCH STUDIES

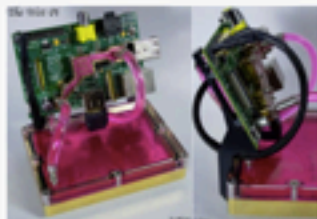
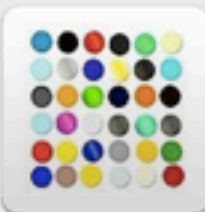
PhD Studentships in Cambridge

The Board of Graduate Studies manages admission of the University's graduate students. Prospective students should start [here](#) - for an introduction to the University of Cambridge, the courses we offer, how to apply for postgraduate study, how your



Scientific instrumentation and computing hardware

Latest news in scientific computing and development of low-cost hardware for laboratory instrumentation. The pages in this section contain links to web sites with DIY instructions for building hardware, specialised components and open-source or low-cost software, with a focus on Arduino and Raspberry Pi.



Water Cooled Raspberry Pi Case

Hacker Phame has taken a low powered Raspberry Pi mini PC, which is available to purchase for around \$35, and added a pimped up cooling system creating a water cooled Raspberry Pi mini computer. To re-cap the Raspberry Pi mini computer which is manufactured in Wales, UK, is equipped with a a 700 MHz ARM11 processor, which can safely be overclock it to 1 GHz if desired. (...) Original Story...

| 14 hits | [Read more](#)



Pick the Right Electronics Board for Your DIY Projects

The Arduino, Raspberry Pi, and BeagleBone are all low-cost controllers that are great for your DIY projects, but it's a little confusing when you're trying to figure out which one is best suited for you. Make has a breakdown of all three and what types of projects they're best for. Read more... Pick the Right Electronics Board for Your DIY Projects (Via Lifehacker.)

| 20 hits | [Read more](#)

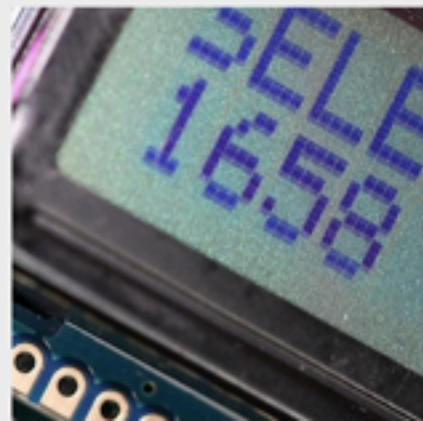


Eyes-on with Ninja Blocks 'home automation for hackers' (video)

Yet another Kickstarter success story is here on the Hardware Alley Floor

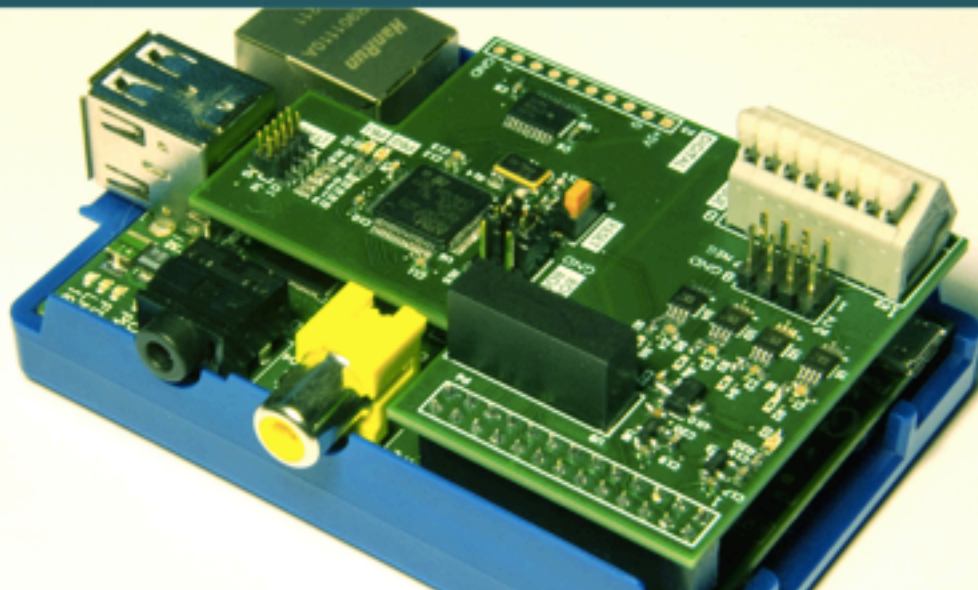
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OPENLABTOOLS IN CAMBRIDGE



OpenLabTools

Home Projects Resources Contacts



Data acquisition system for Raspberry Pi computers (prototype)

1 of 2

The OpenLabTools project

The OpenLabTools initiative aims to provide a forum and knowledge centre for the development of low cost and open access scientific tools, with an emphasis on undergraduate and graduate teaching and research. The programme will officially start in October 2013. In order to bootstrap this initiative, a number of MEng projects (4-5 per year) will be offered to establish the core components required for such tools; these include data acquisition, sensing, actuating, processing and 3D manufacturing. Protocols, designs and tutorials will be published on this website. These components will be subsequently combined to establish a documented collection of instruments, to be developed and maintained by a community of undergraduate and graduate students of the University.

We anticipate that the core tools will be rolled out in undergraduate laboratories from 2014 onwards. We would like to invite anybody interested in this programme to get in touch with us and explore ways to contribute, either by using the tools (once ready) or

Academic team

Anurag Agarwal

(Fluids - Acoustics)

Oliver Haderl

(CamBridgeSens)

Jim Haseloff

(Synthetic Biology)

Simone Hochgreb

(Fluids - Reacting flows)

Alexandre Kabla

Related initiatives

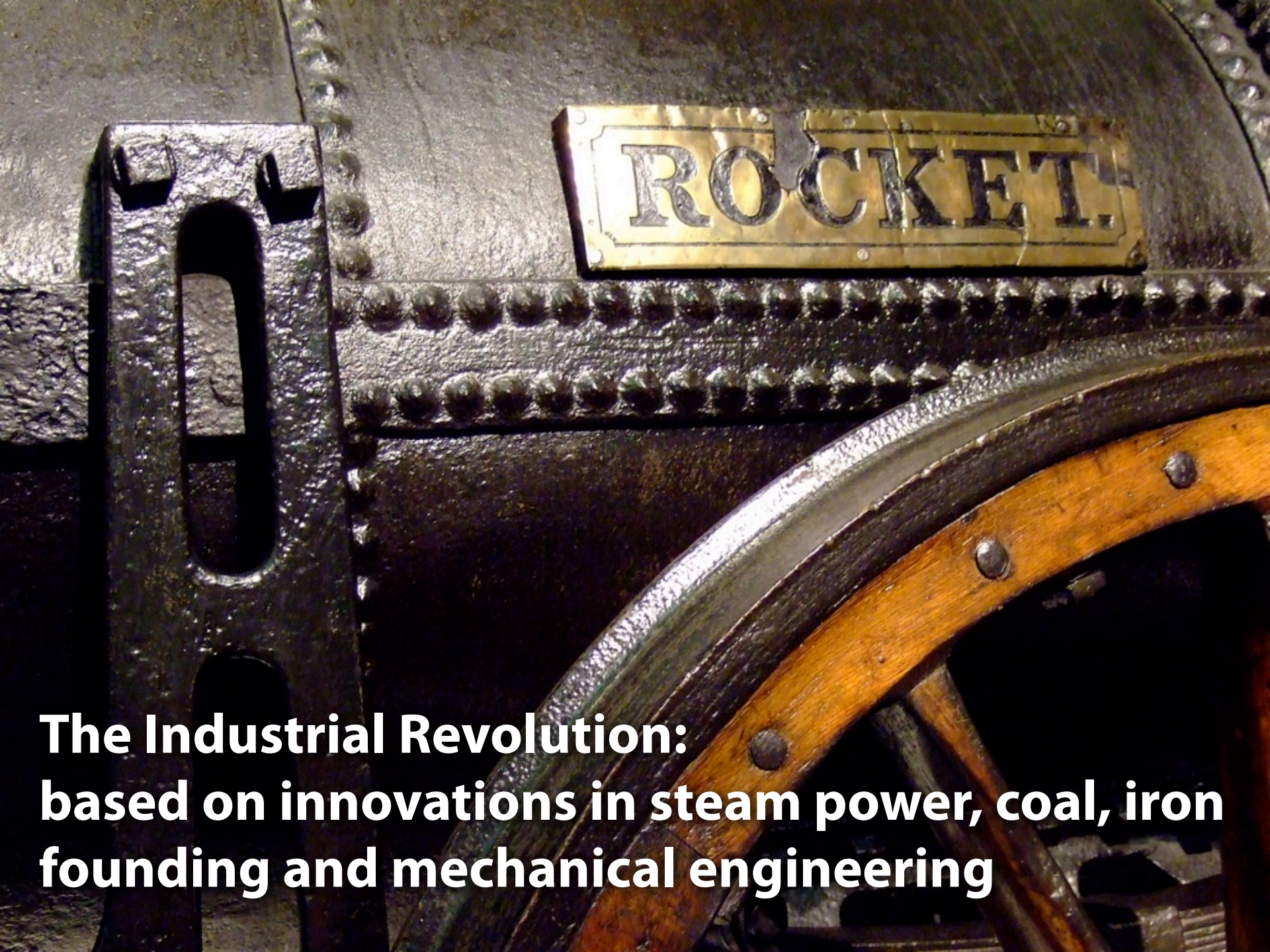
[The Raspberry Pi foundation](#)

[Arduino](#)

[Synbio.ork.uk](#)

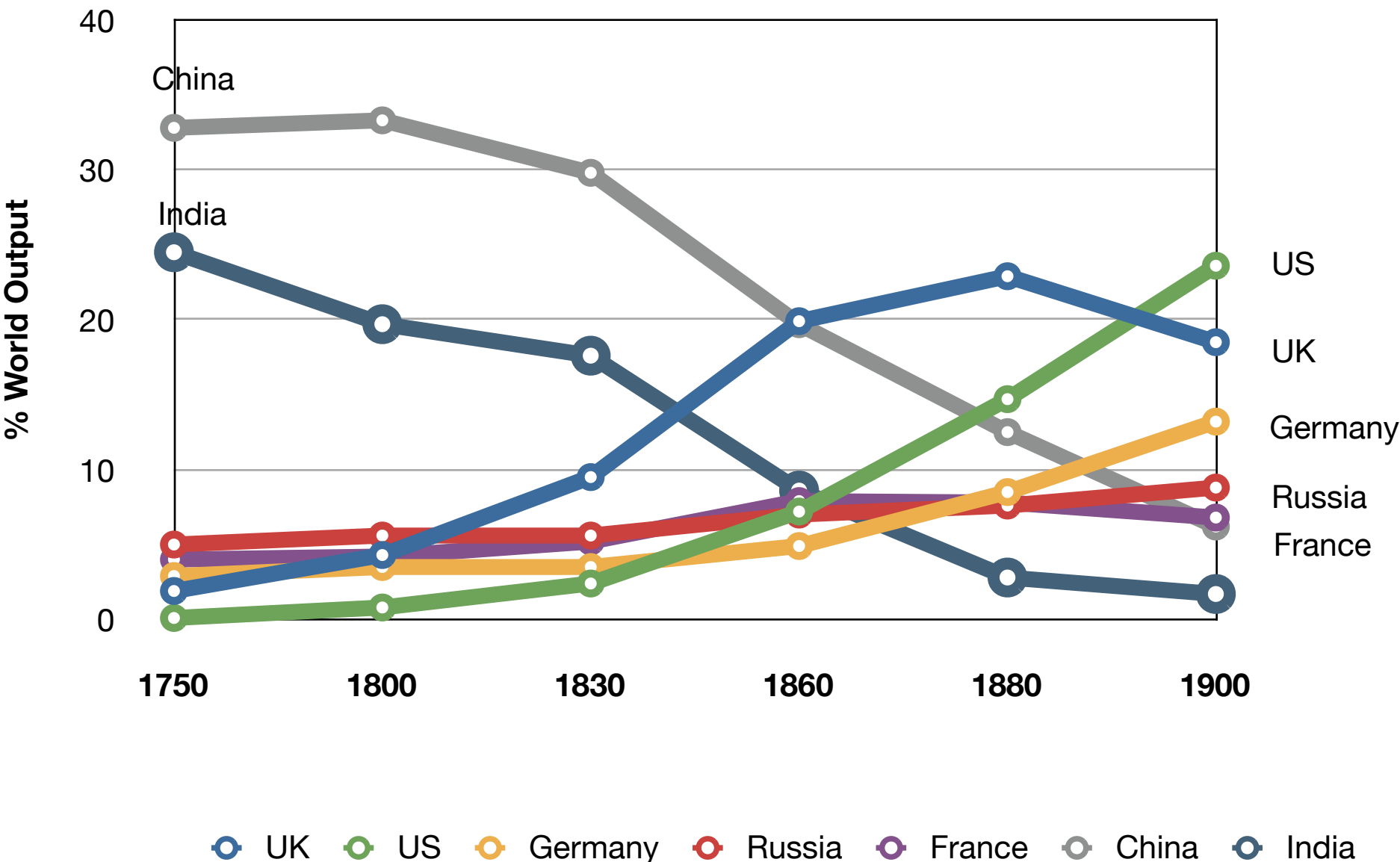
[Makespace \(Cambridge, UK\)](#)

[CamBridgeSens](#)

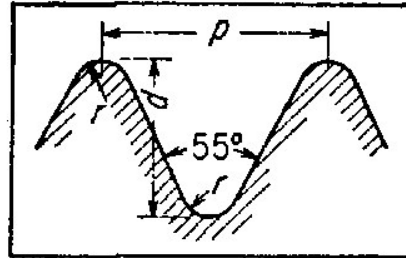
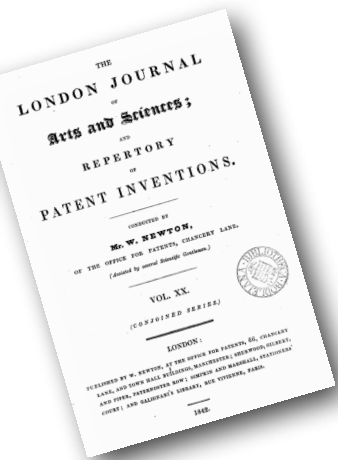


**The Industrial Revolution:
based on innovations in steam power, coal, iron
founding and mechanical engineering**

Relative shares of world manufacturing output 1750-1900



Standardisation of parts for construction

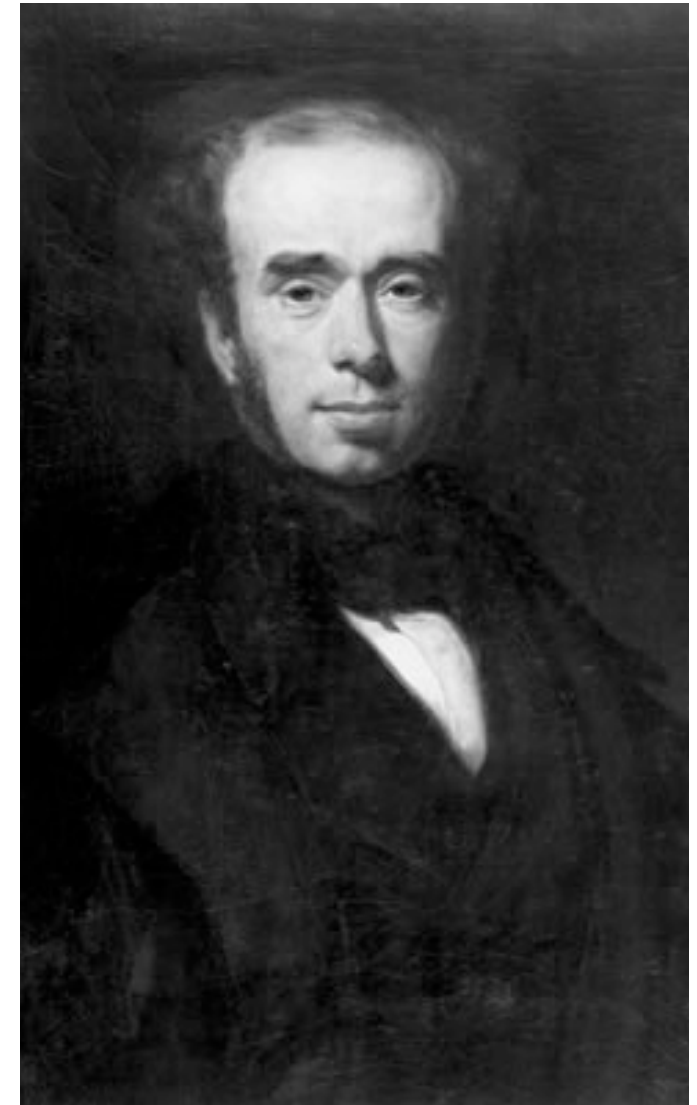


“On an uniform system of Screw Threads.”

By Joseph Whitworth, Assoc. Inst. C. E.

The subject considered in this paper, is the importance of having a constant thread for a given diameter in all screws used in fitting up steam engines and other machinery. It is argued, that uniformity of thread would be productive of economy, both in the use of screwing apparatus, and in the consumption of bolts and nuts. The refitting shop of a railway or steam packet company, affords a striking instance of the advantage to be derived from the application of this principle. If the same system of screw threads were common to the different engines, a single set of screwing tackle would suffice for any repairs.

No attempt appears to have been hitherto made to attain this important object. Engineers have adopted their threads without reference to a common standard. Any such standard must be in a great measure arbitrary, and hence its absence may be accounted for.



Joseph Whitworth 1842

Interchangeable parts led to continuous production methods

