

Synthetic Botany

Engineering plant form

Jim Haseloff

University of Cambridge

But...bespoke DNA assembly techniques are still common practice in the field after 40 years



Smithsonian museum exhibit: Stanley Cohen's laboratory bench

Construction of Biologically Functional Bacterial Plasmids *In Vitro*

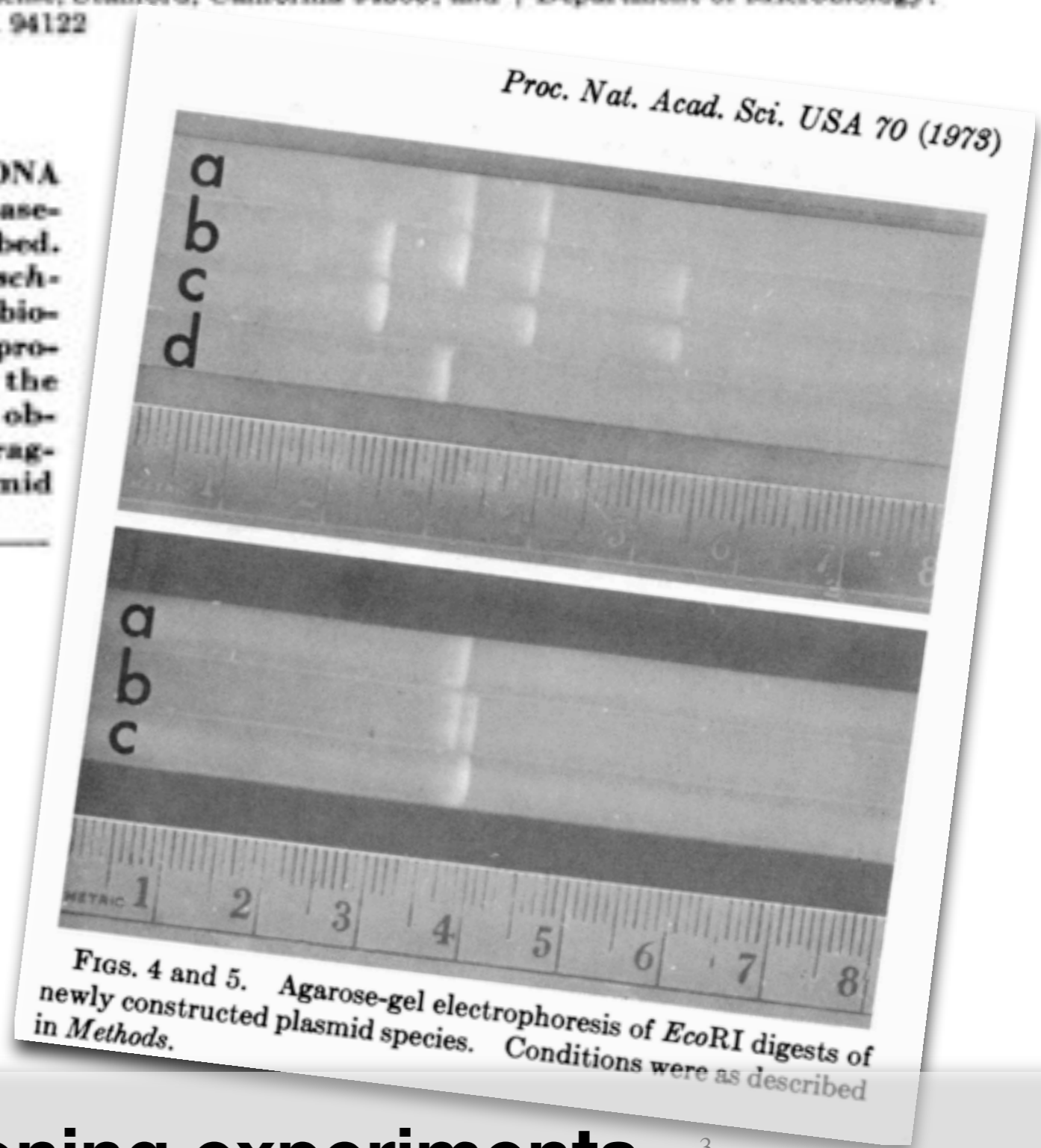
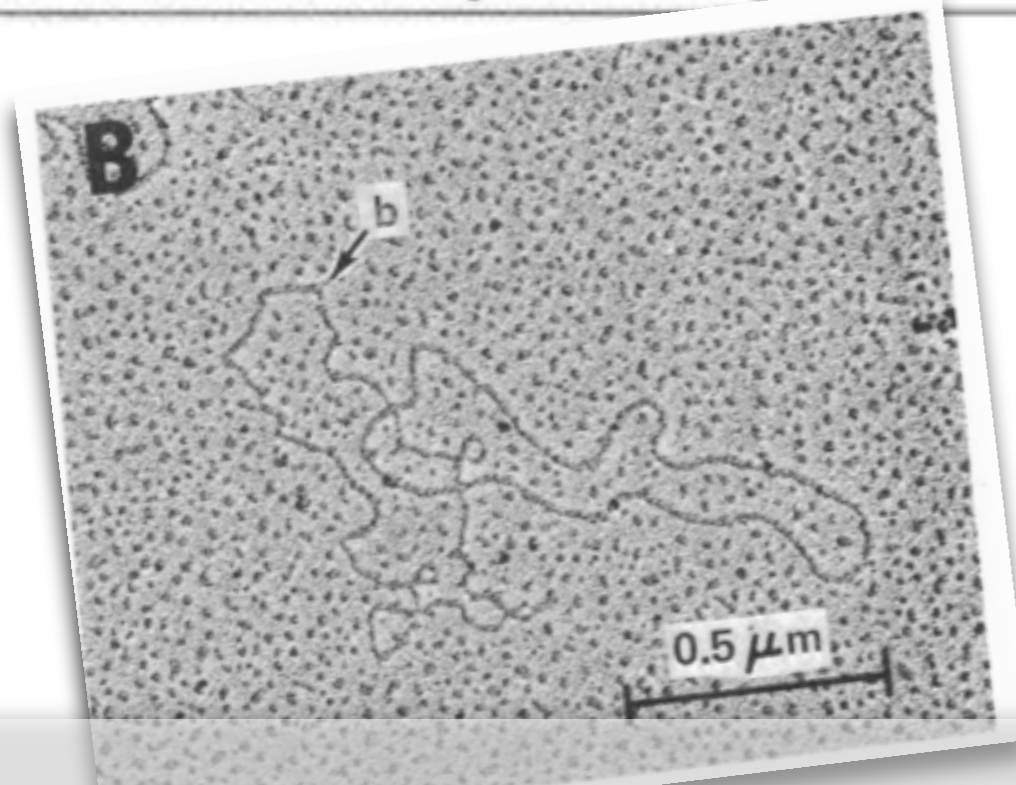
(R factor/restriction enzyme/transformation/endonuclease/antibiotic resistance)

STANLEY N. COHEN*, ANNIE C. Y. CHANG*, HERBERT W. BOYER†, AND ROBERT B. HELLING†

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Communicated by Norman Davidson, July 18, 1973

ABSTRACT The construction of new plasmid DNA species by *in vitro* joining of restriction endonuclease-generated fragments of separate plasmids is described. Newly constructed plasmids that are inserted into *Escherichia coli* by transformation are shown to be biologically functional replicons that possess genetic properties and nucleotide base sequences from both of the parent DNA molecules. Functional plasmids can be obtained by reassociation of endonuclease-generated fragments of larger replicons, as well as by joining of plasmid DNA molecules of entirely different origins.



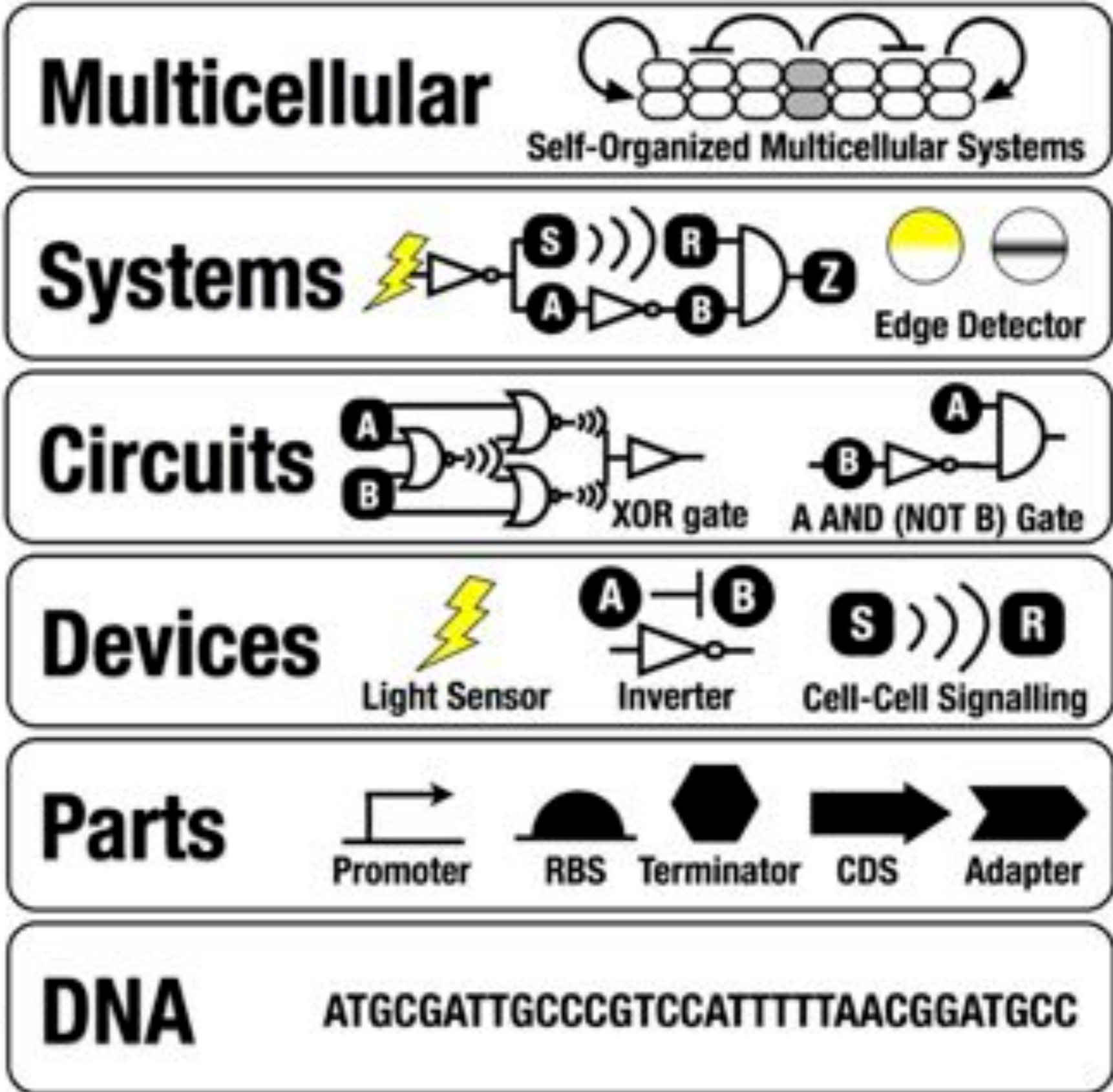
Figs. 4 and 5. Agarose-gel electrophoresis of *Eco*RI digests of newly constructed plasmid species. Conditions were as described in *Methods*.

1973 - first molecular cloning experiments...³



Tom Knight, MIT

2003 Invention of standardised DNA parts...



1950s

Silicon
Transistor



1
Transistor

1960s

TTL
Quad Gate



16
Transistors

1970s

8-bit
Microprocessor



4500
Transistors

1980s

32-bit
Microprocessor



275,000
Transistors

1990s

32-bit
Microprocessor



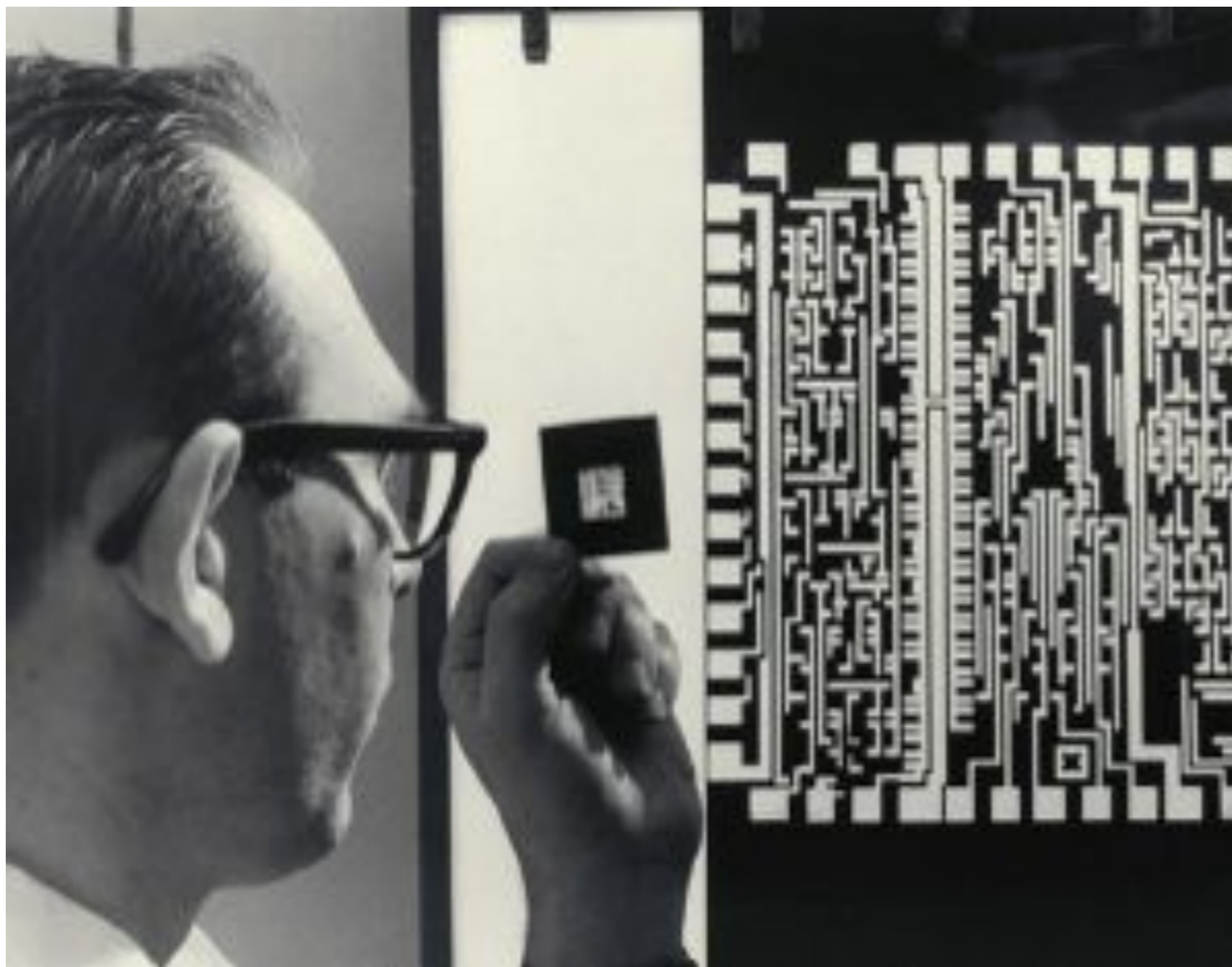
3,100,000
Transistors

2000s

64-bit
Microprocessor



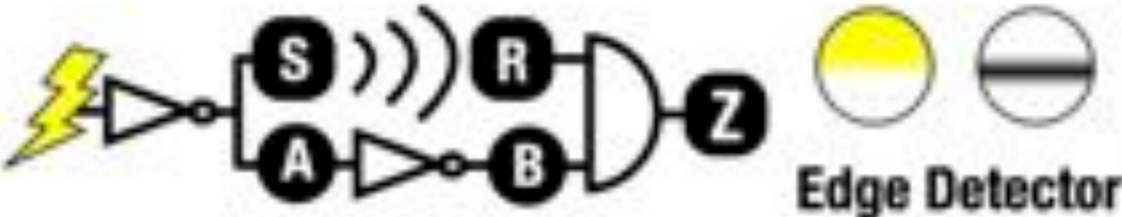
592,000,000
Transistors




2. Development of automated design tools and modular circuits to deal with increased complexity




Multicellular  Self-Organized Multicellular Systems

Systems  Edge Detector

Circuits  XOR gate A AND (NOT B) Gate

Devices  Light Sensor Inverter Cell-Cell Signalling

Parts  Promoter RBS Terminator CDS Adapter

DNA ATGCGATTGCCCGTCCATTTTAAACGGATGCC

Applications for Synthetic Biology

Cell autonomous genetic circuits with self-regulating properties

e.g. microbial engineering,
environmental and biomedical sensors
engineering novel metabolic pathways

Morphogenetic circuits with self organising properties

e.g. microbial biofilms or self-organising communities for
bioremediation and bio catalysis
novel plant and algal feedstocks for bioproduction and bioenergy
tissue engineering

Chloroplast

0.5 μm

ENERGY

METABOLISM

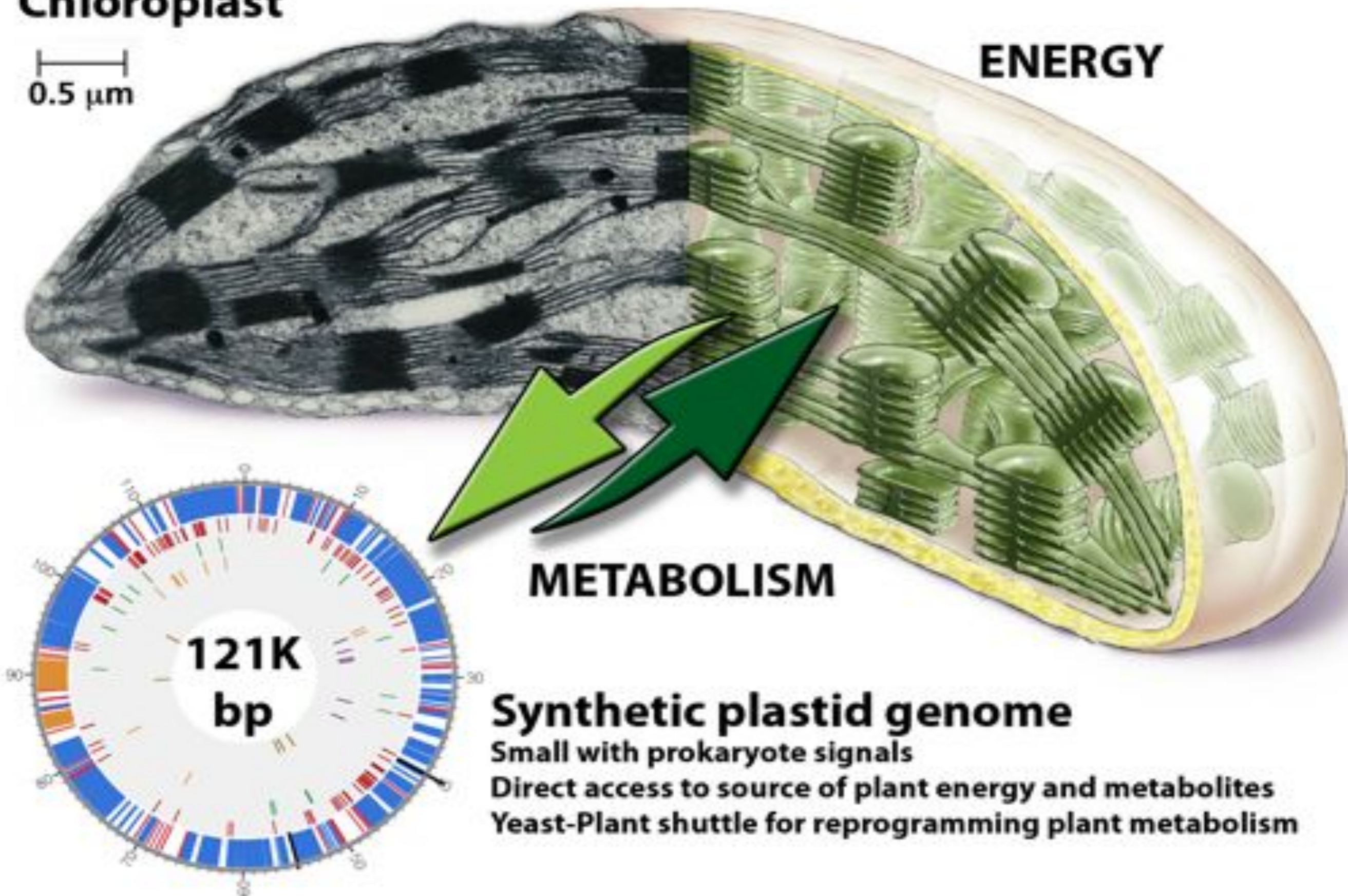
**121K
bp**

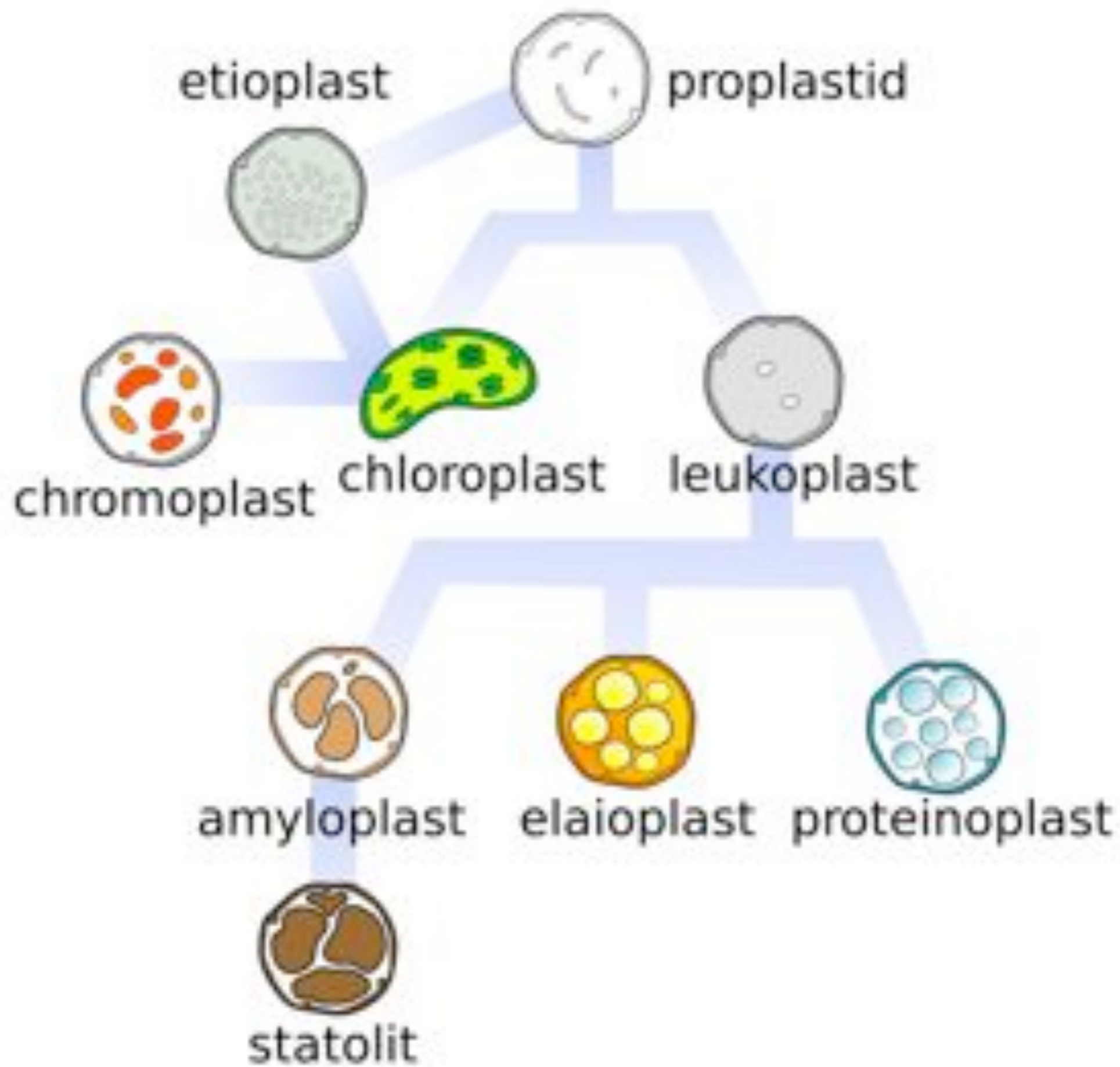
Synthetic plastid genome

Small with prokaryote signals

Direct access to source of plant energy and metabolites

Yeast-Plant shuttle for reprogramming plant metabolism





Applications for Synthetic Biology

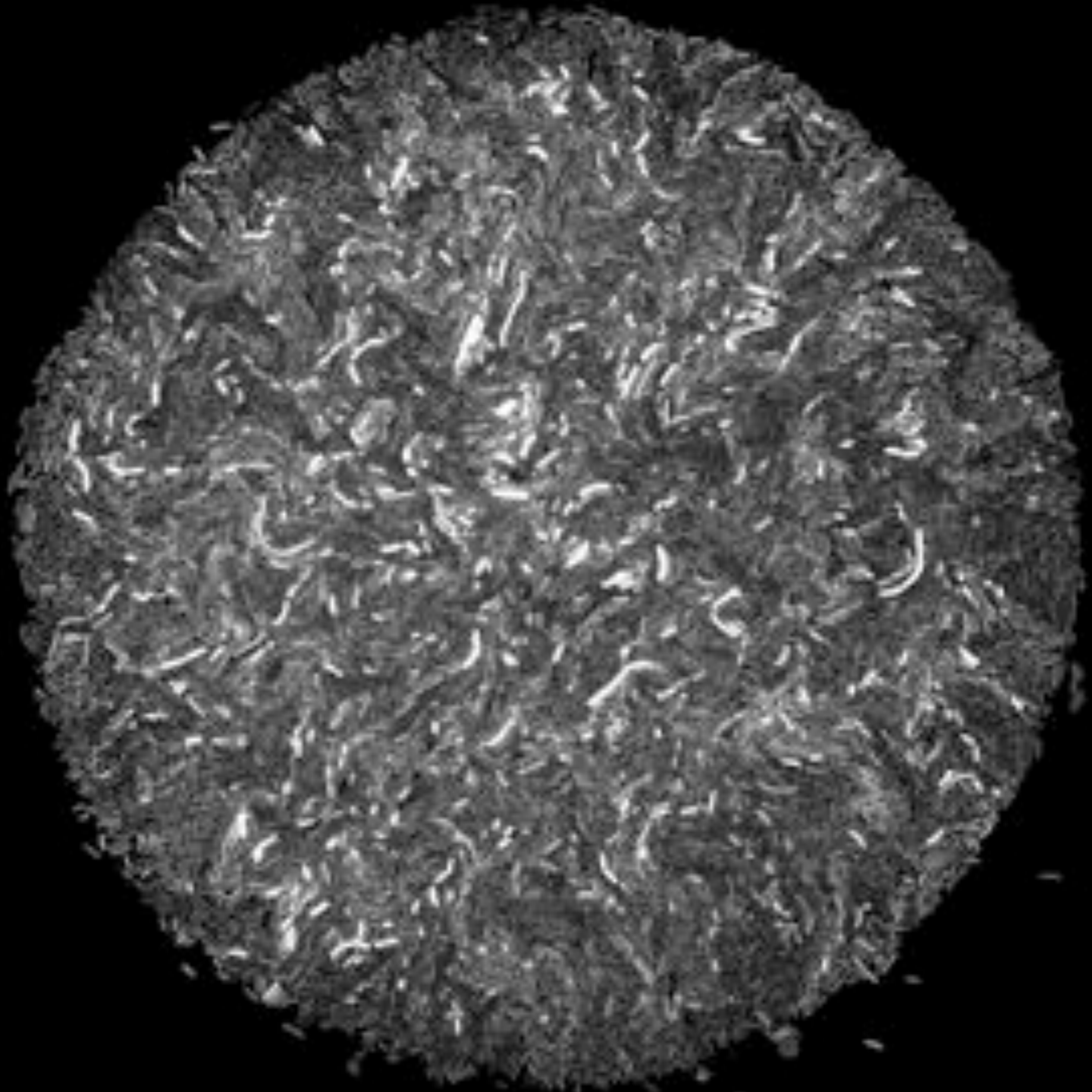
Cell autonomous genetic circuits with self-regulating properties

e.g. microbial engineering,
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engineering novel metabolic pathways

Morphogenetic circuits with self organising properties

e.g. microbial biofilms or self-organising communities for
bioremediation and bio catalysis
novel plant and algal feedstocks for bioproduction and bioenergy
tissue engineering

E. coli



Tim Rudge

Genetic program



Cellular automata model



**Finite element model
for cell wall
mechanics**

**Algorithm for
Cell Division**

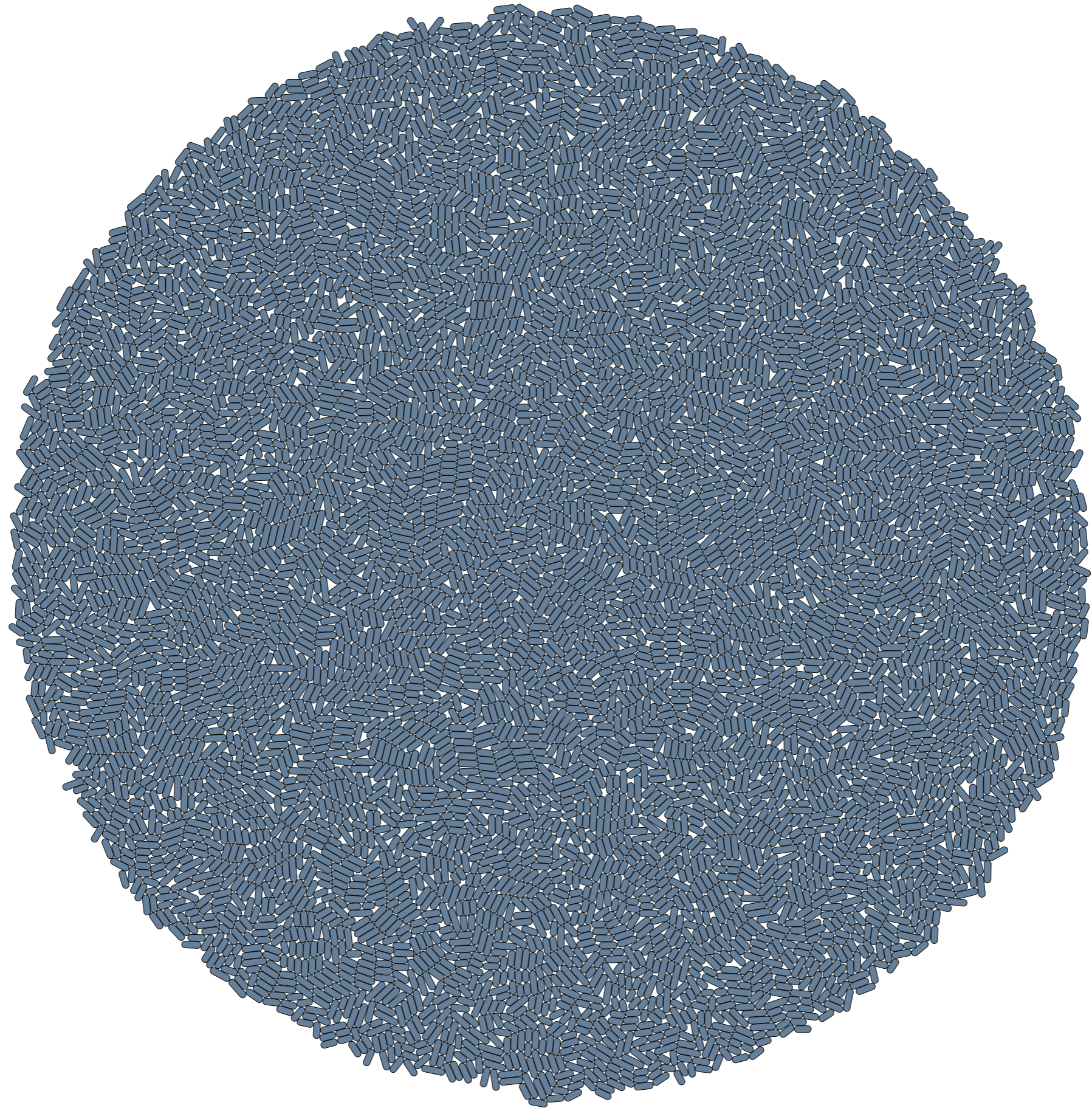


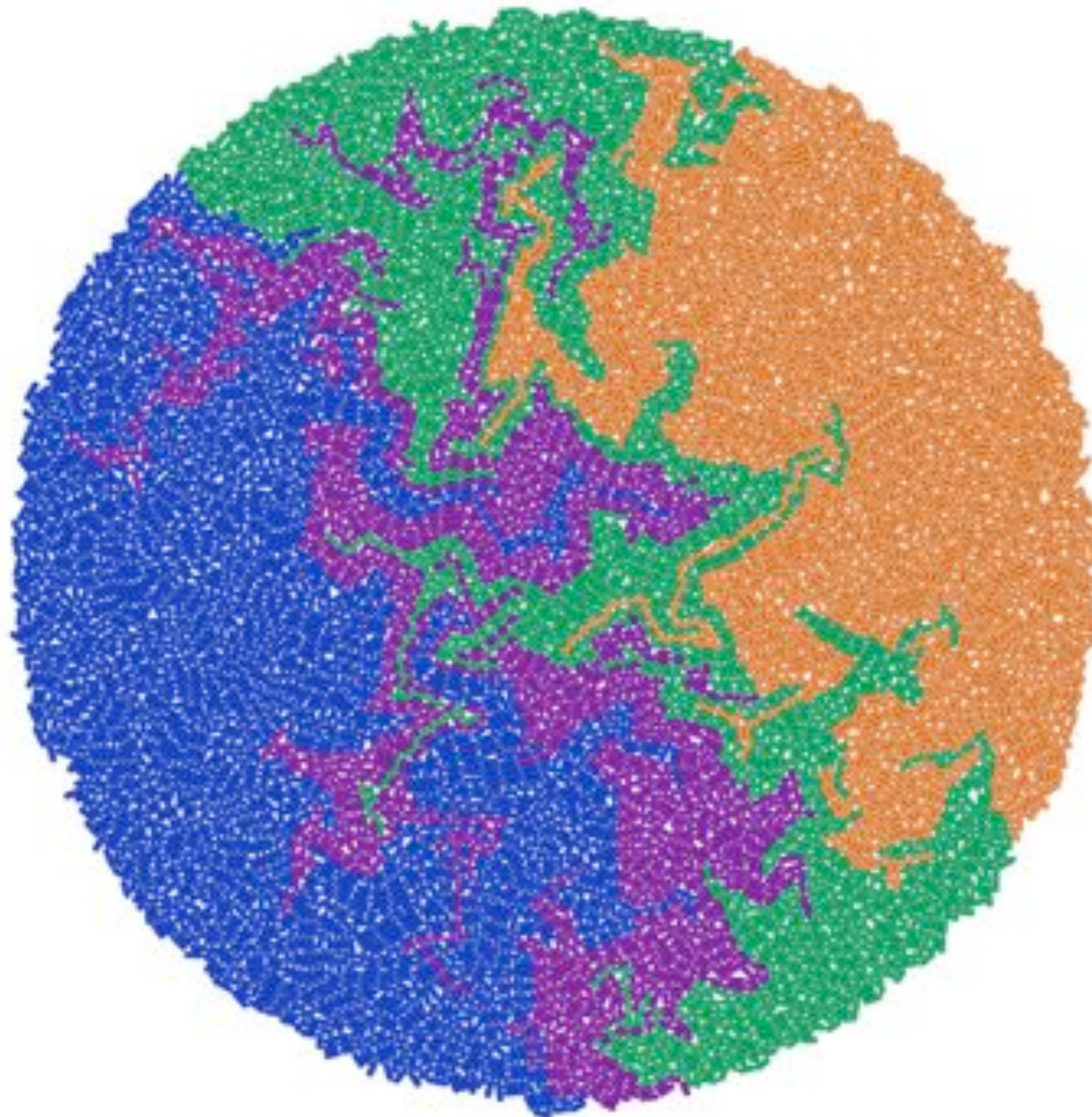
CELLMODELLER

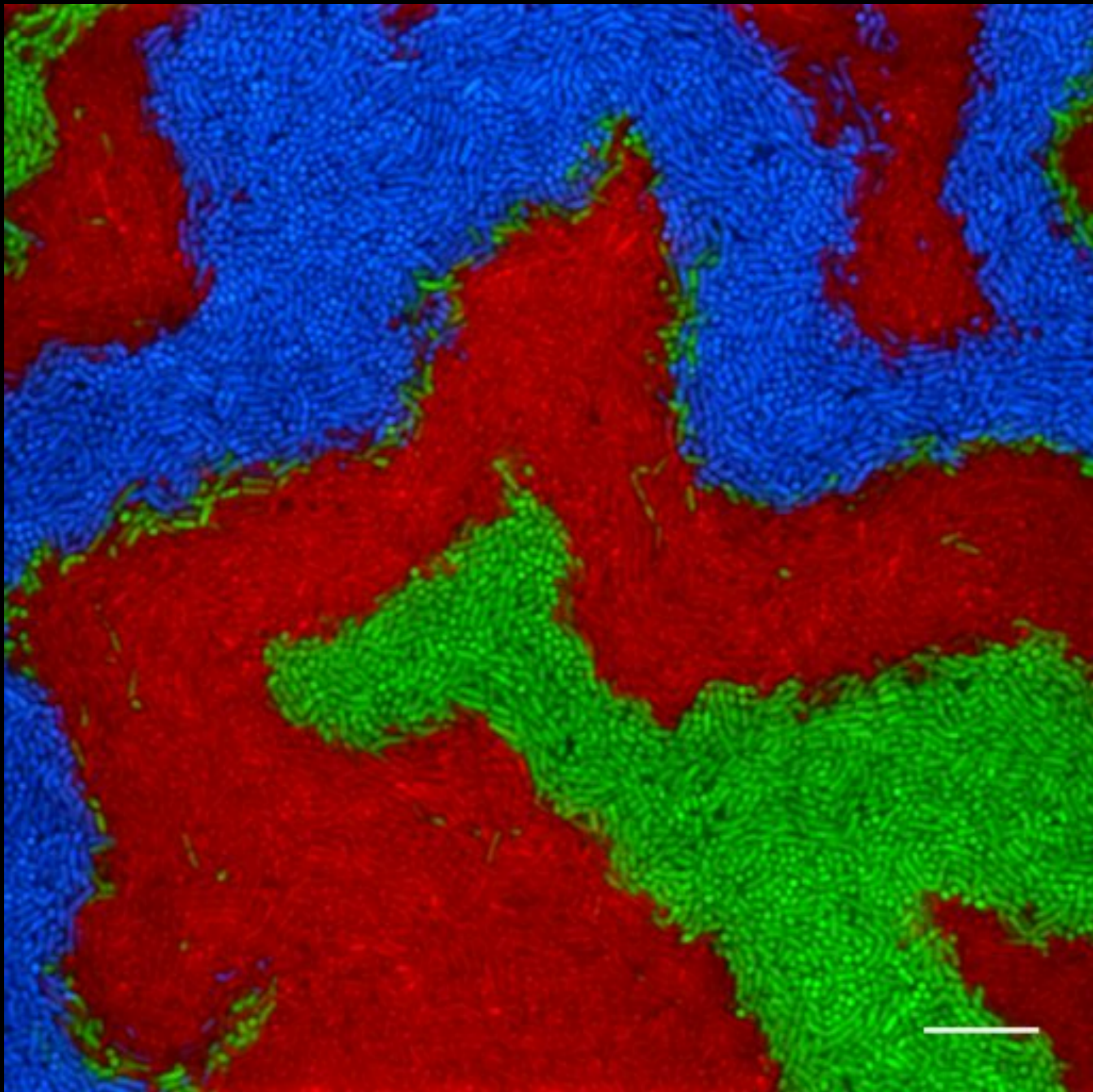
**A software engine for
programmed growth of
multicellular tissues.**

**A tool for testing prototypes
of DNA based morphogenetic
programs.**

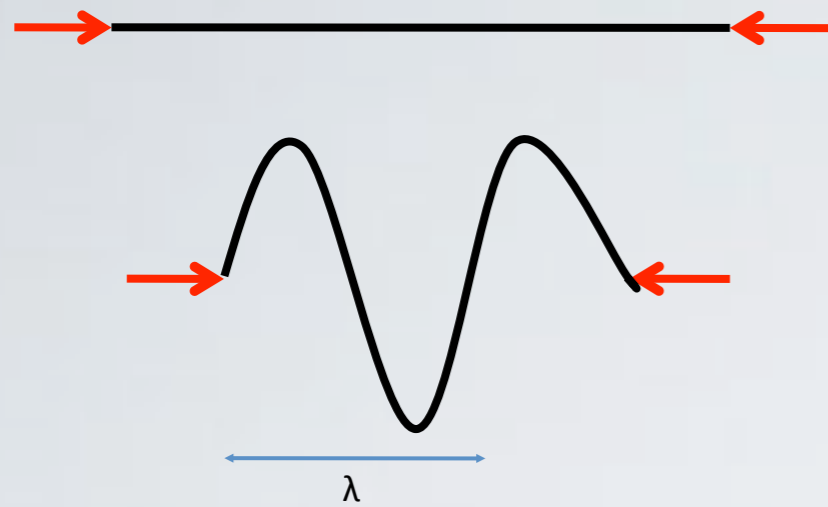
www.cellmodeller.org



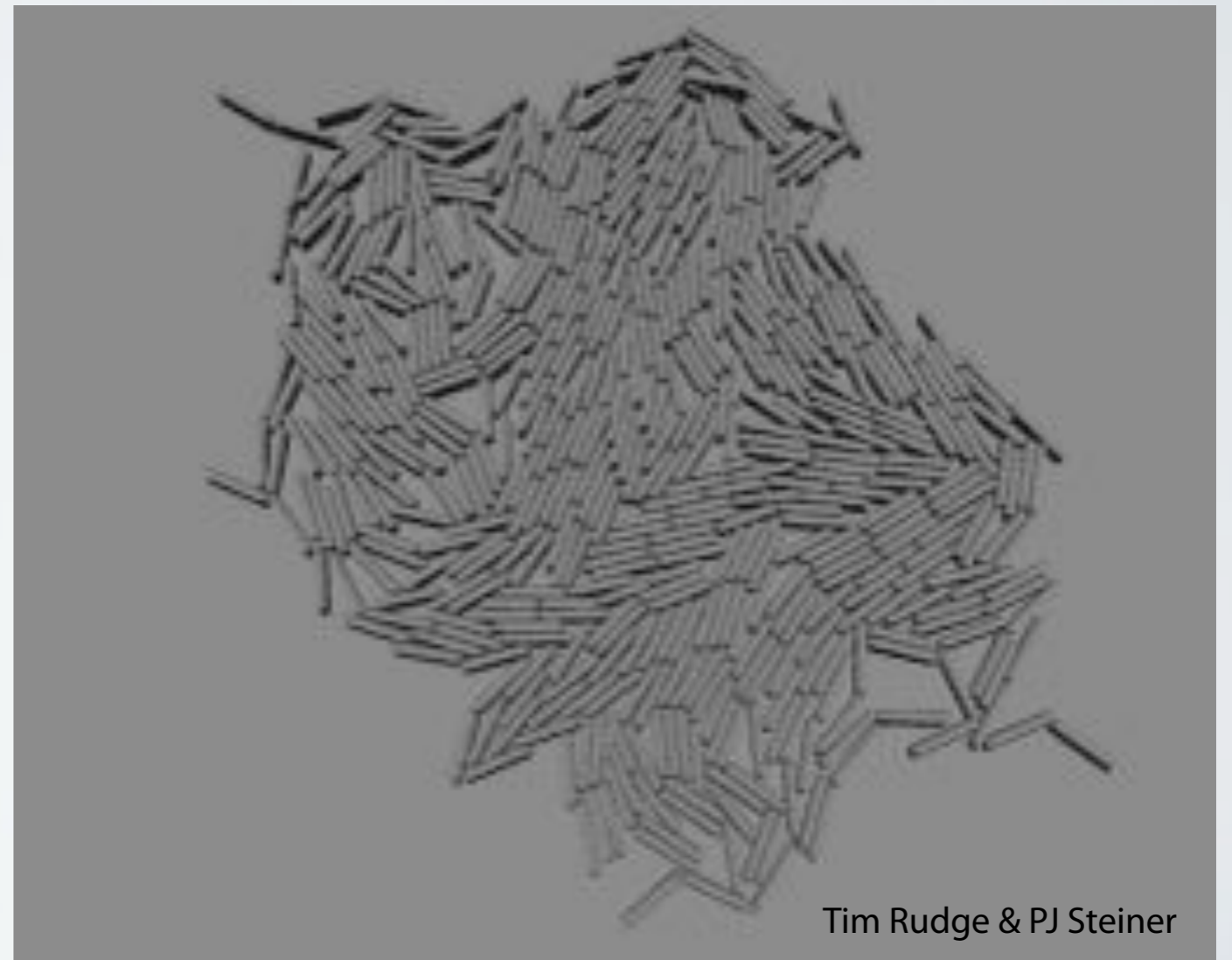
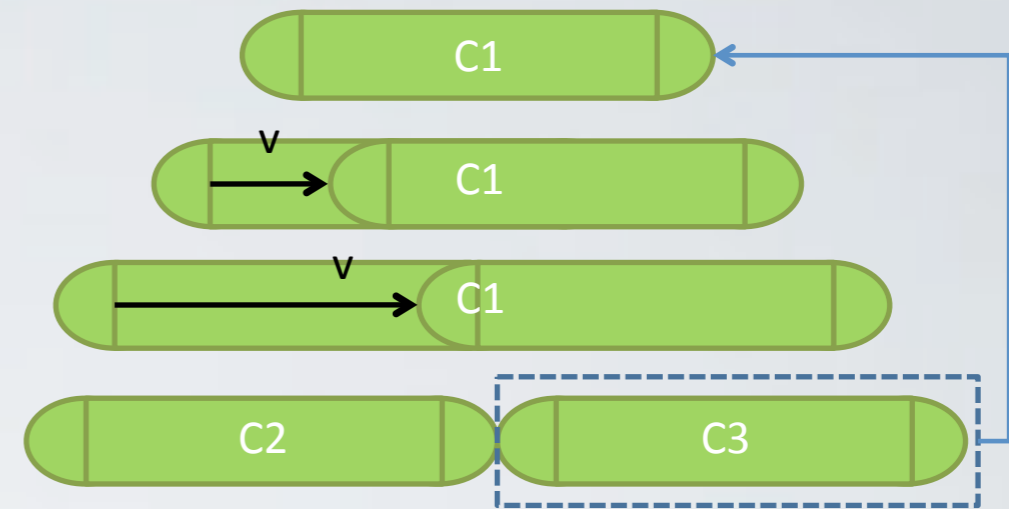




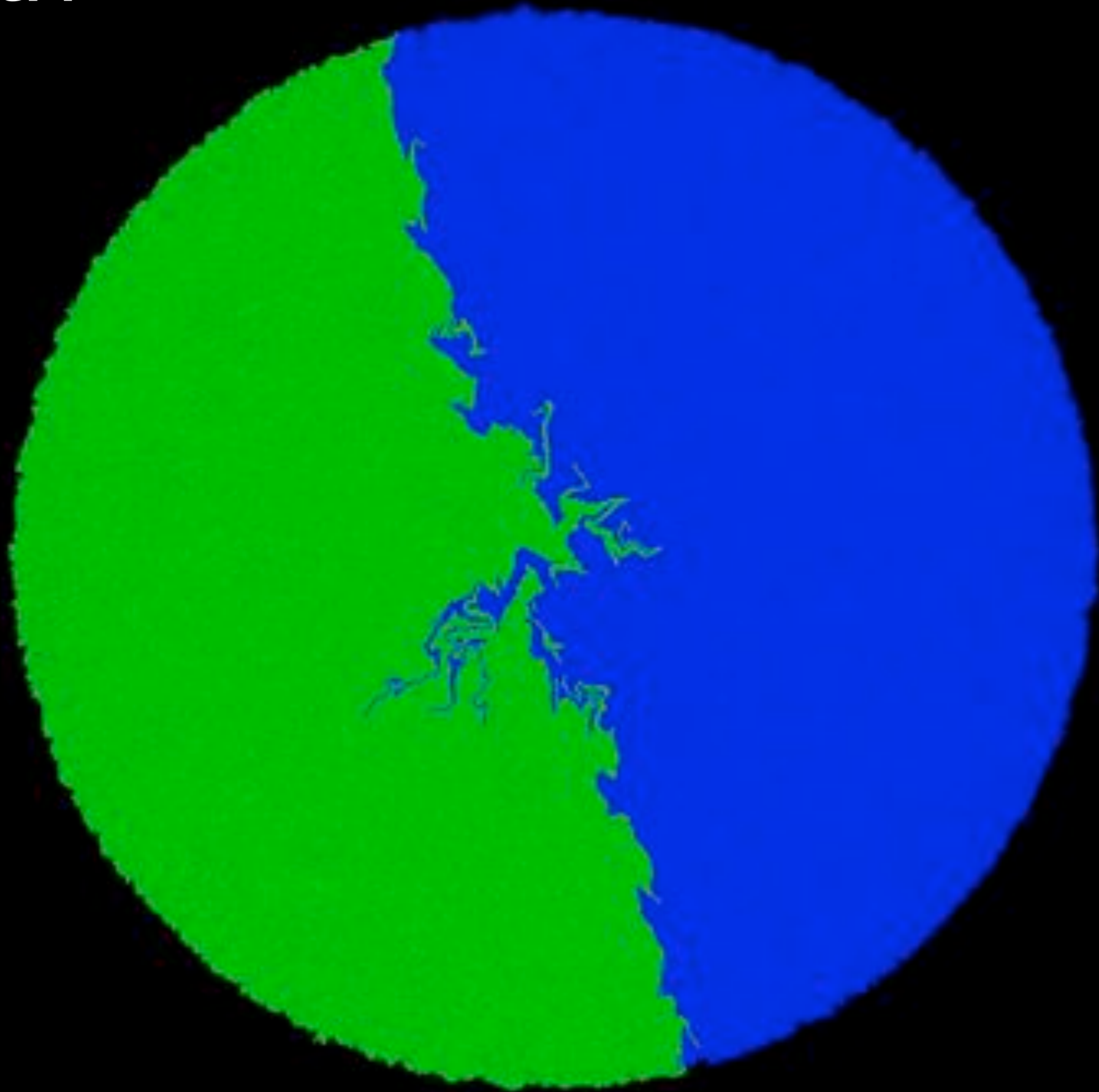
Folding growth pattern due to buckling

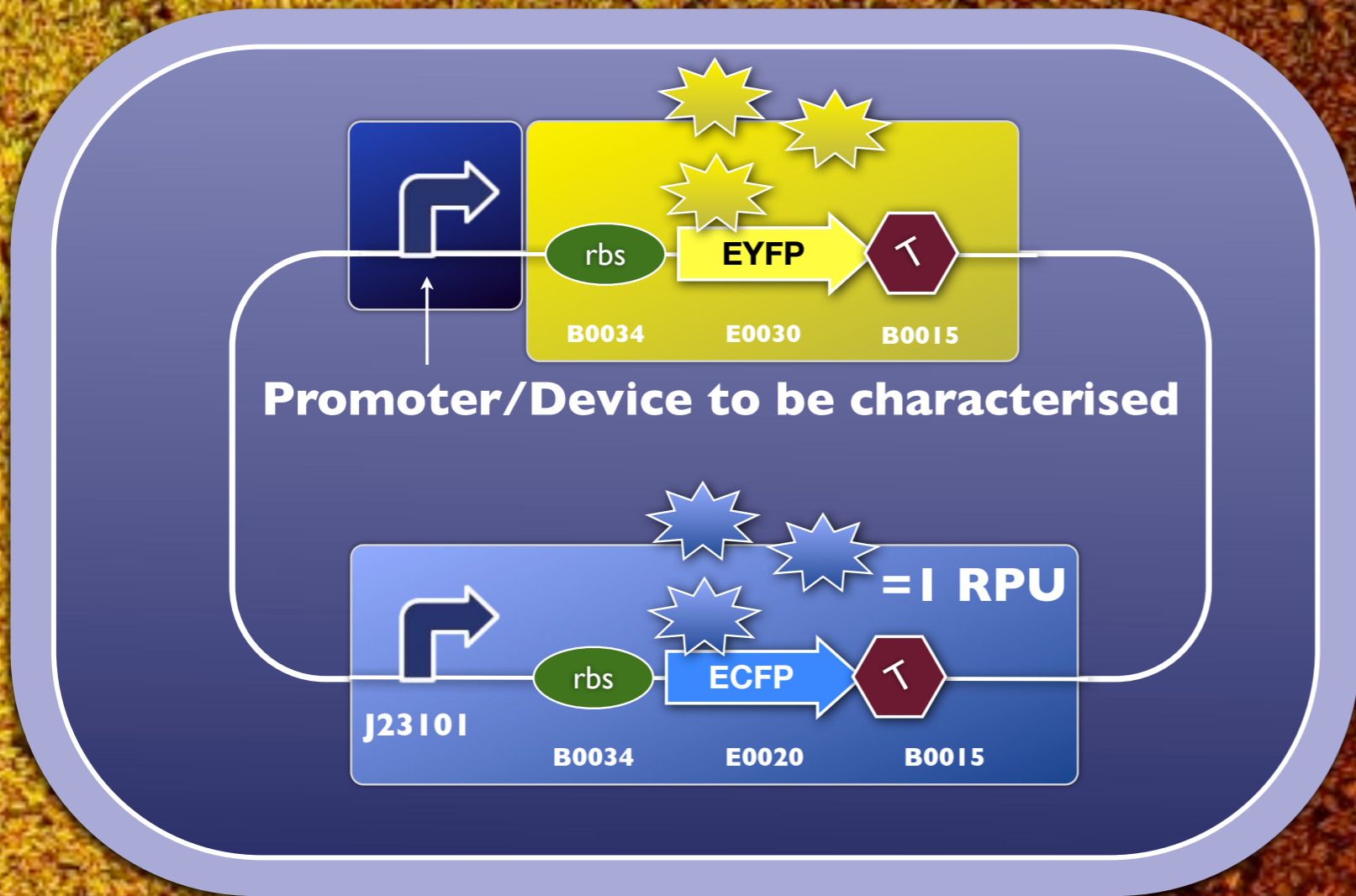


3D Rigid body dynamics



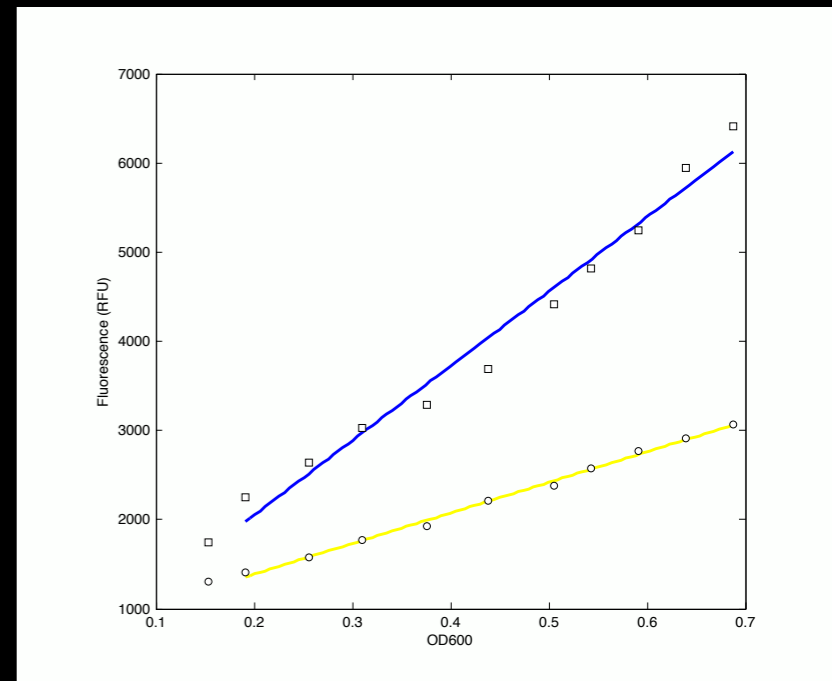
Software model for 3D growth of *Bacillus subtilis*



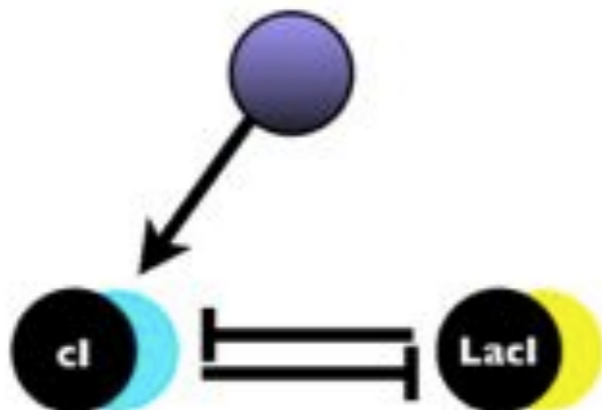
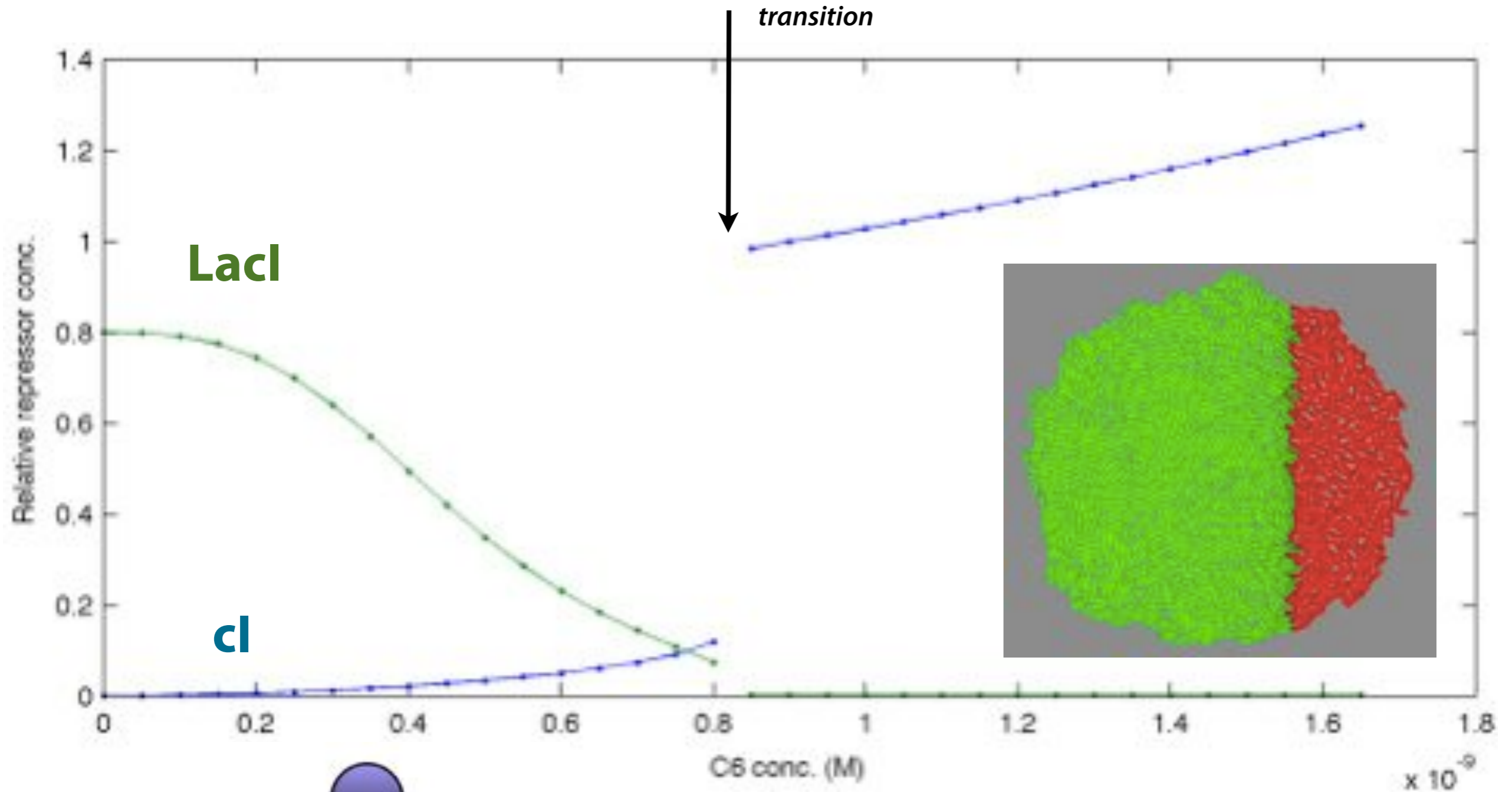


Measuring gene expression in living bacterial cells: ratiometric fluorescence measurement of promoter activity

James Brown & Fernan Federici

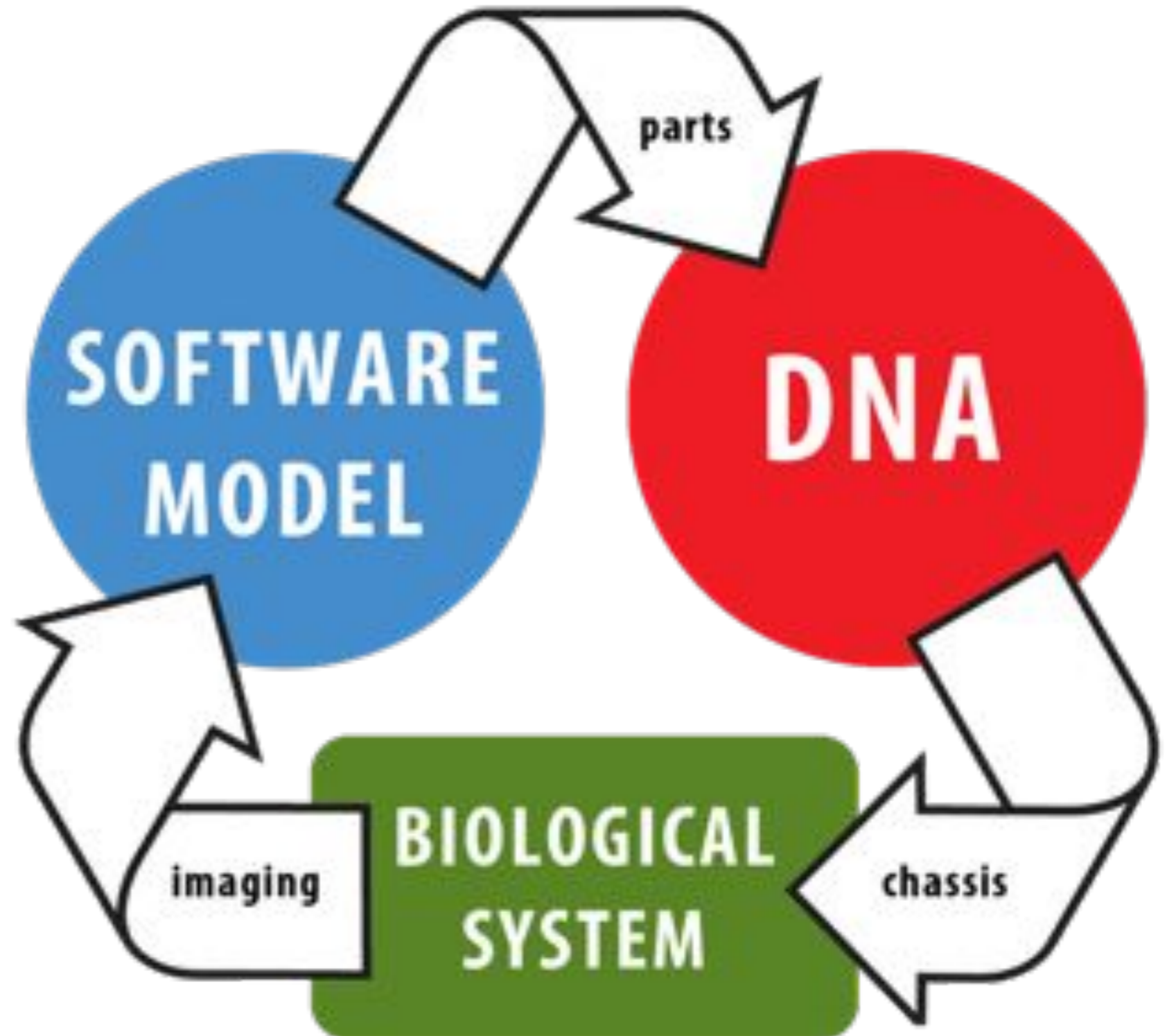


Modelling of metastable circuits

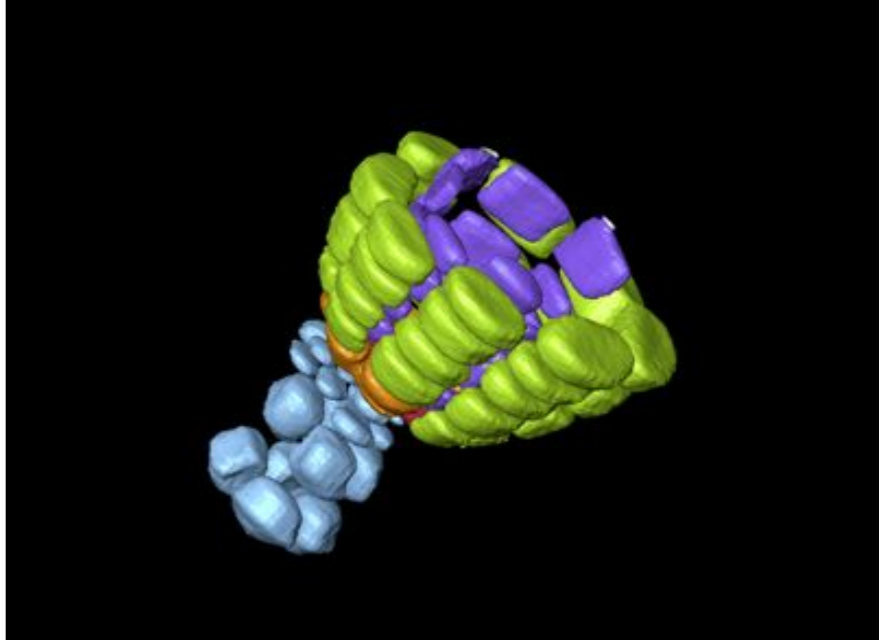
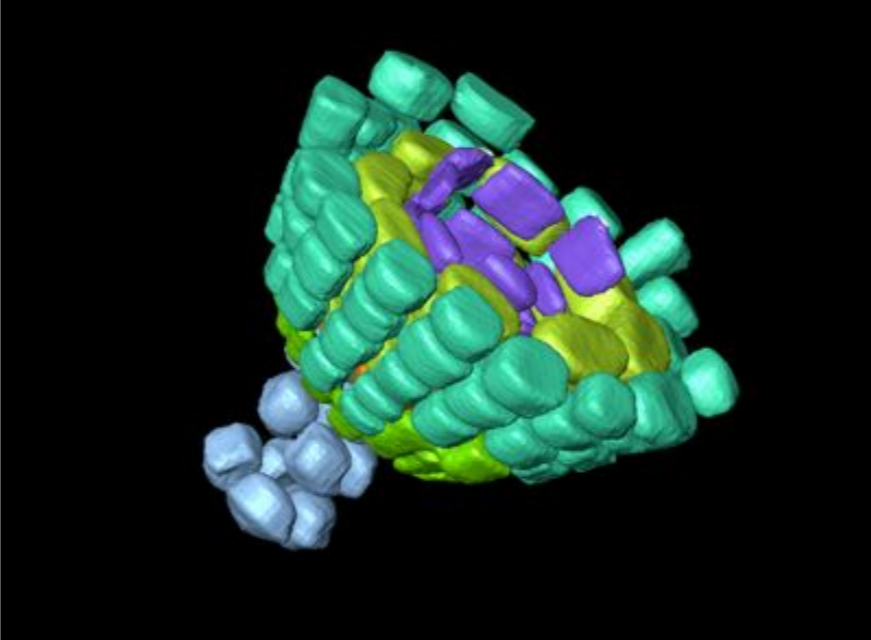
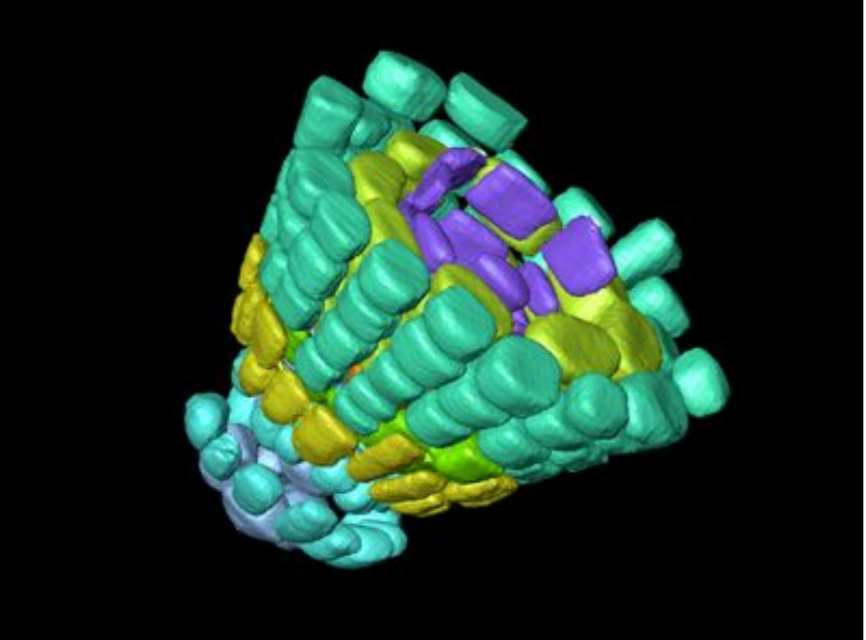
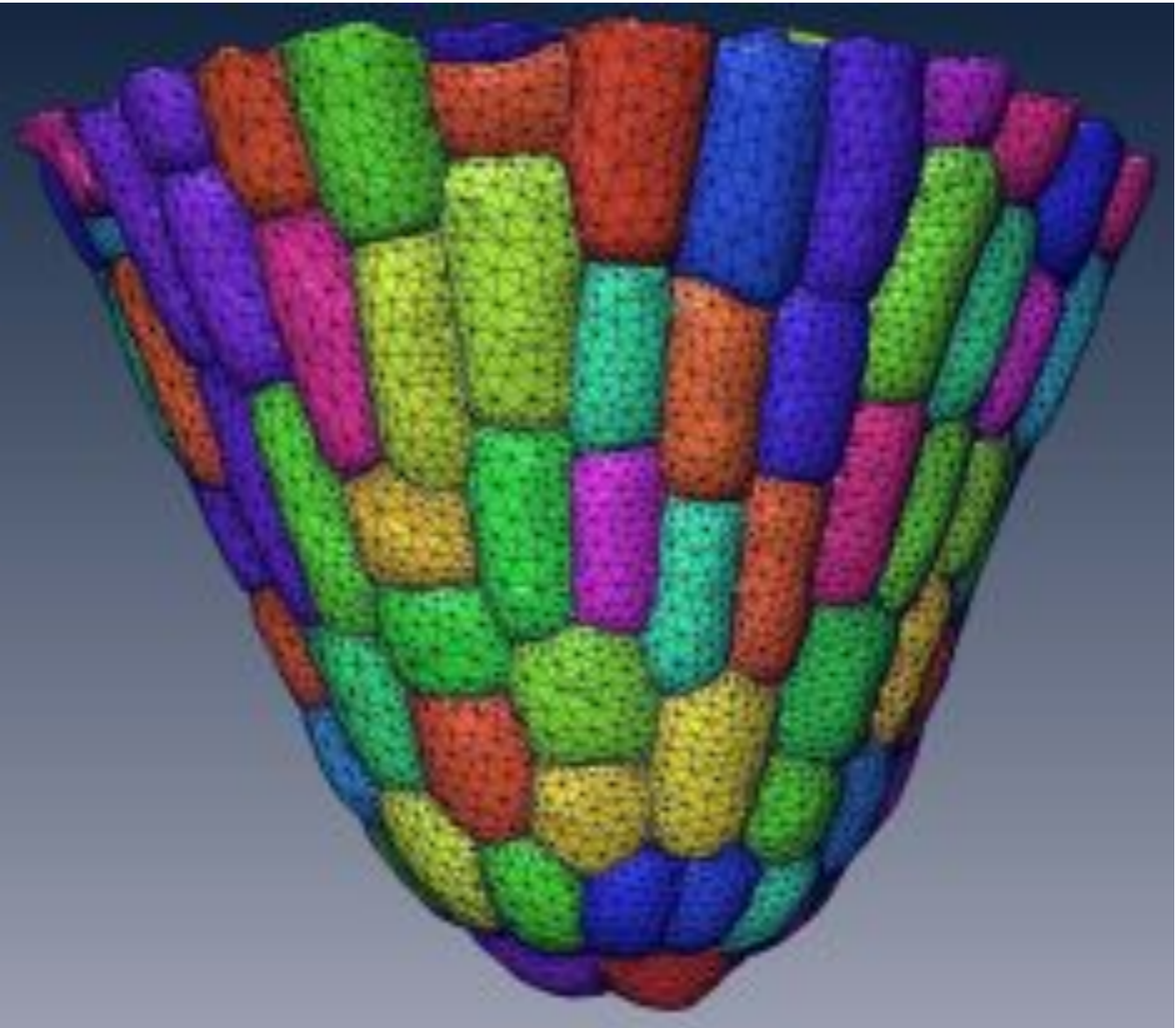
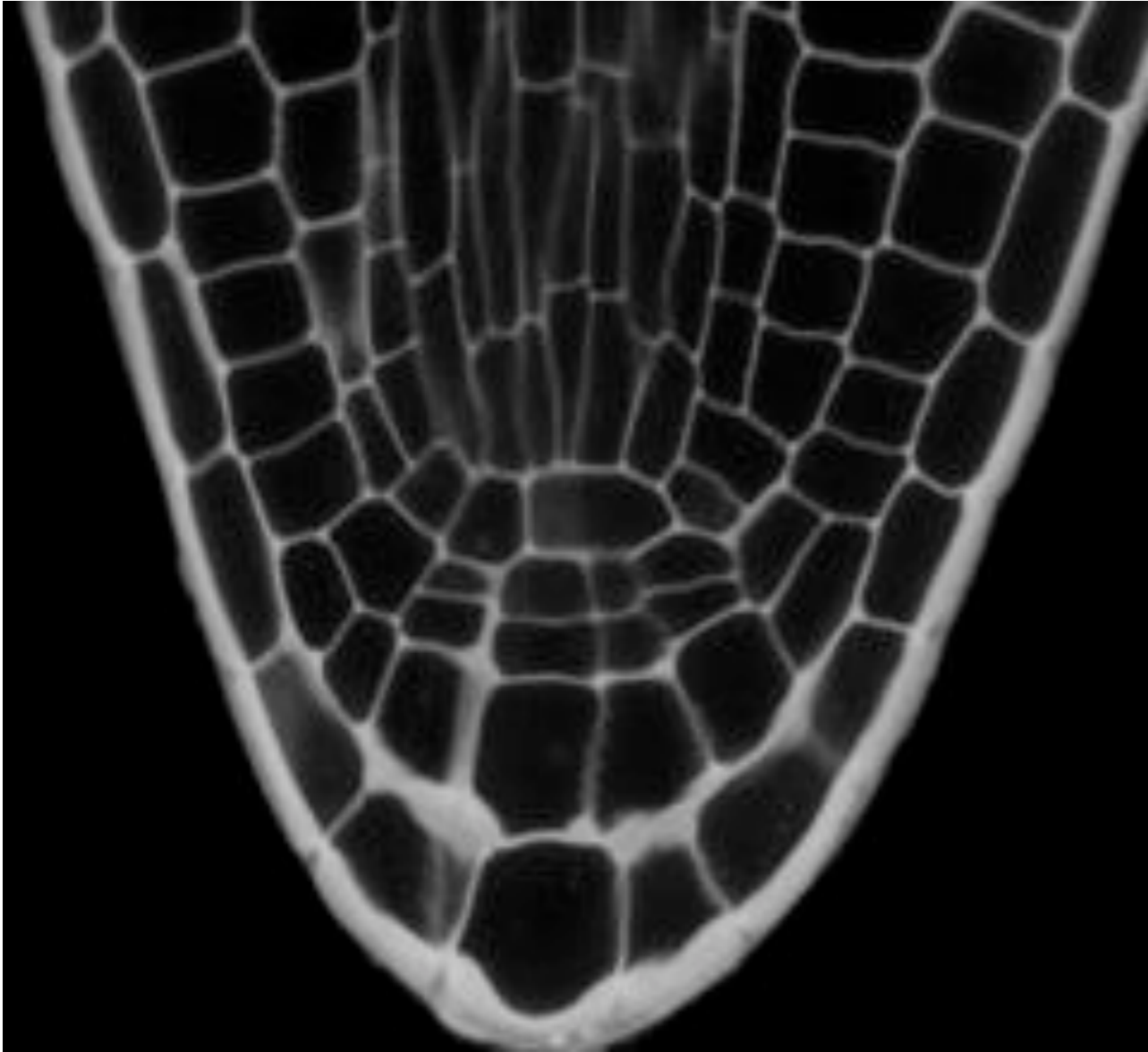


Design cycle for Synthetic Biology systems

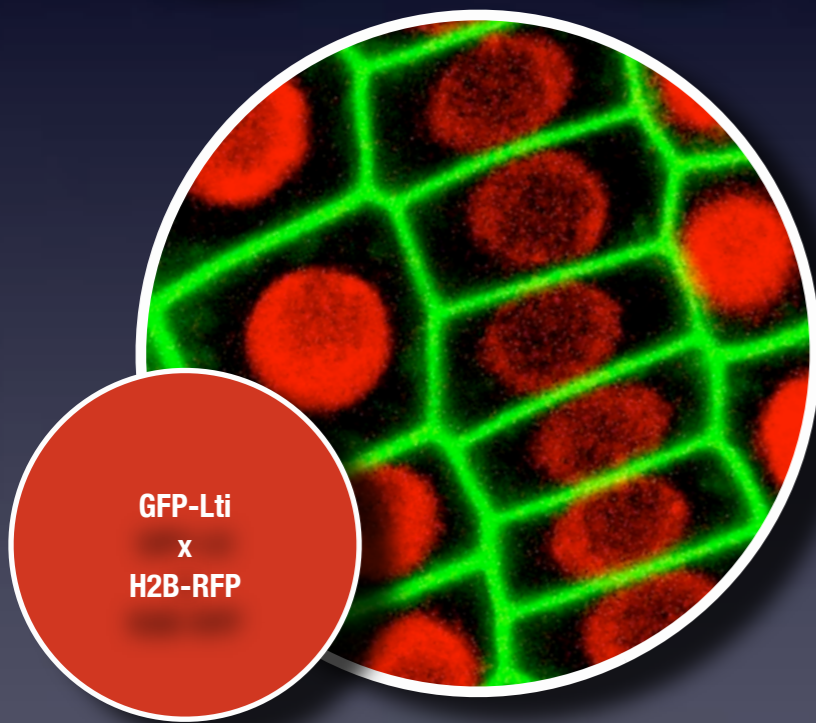
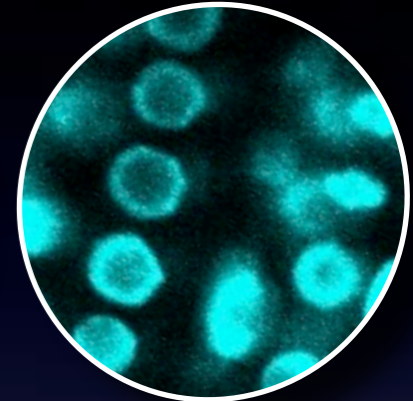
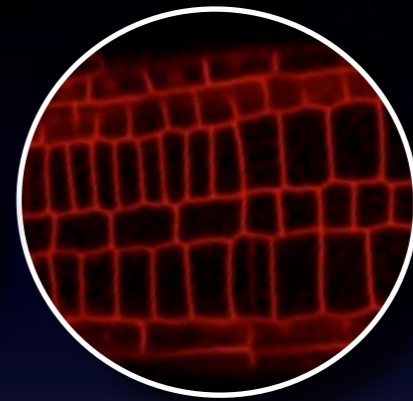
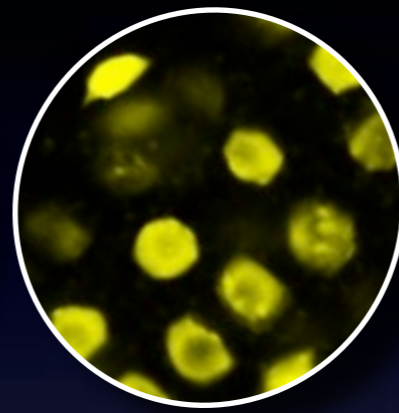
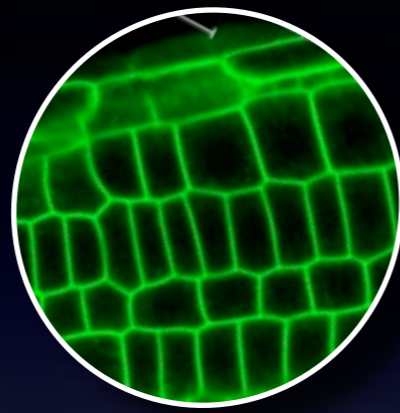
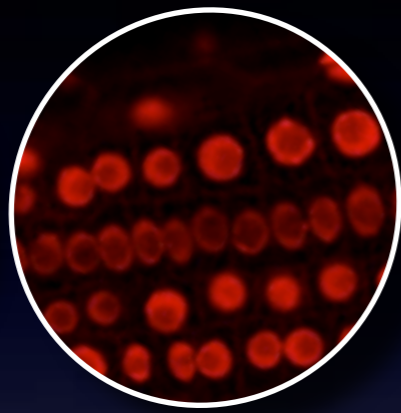
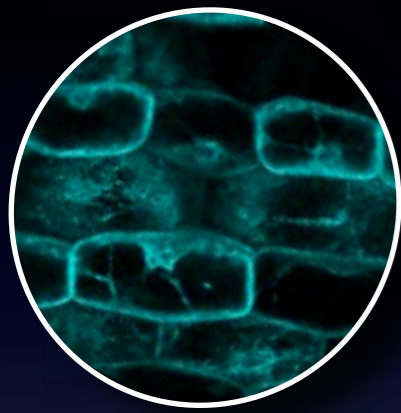
1. Specification and design of the system using computer models of the biological system
2. Construction of genetic circuits using standard DNA parts and high throughput assembly techniques
3. Transformation of chassis and visualisation of gene expression, cell states and phenotype



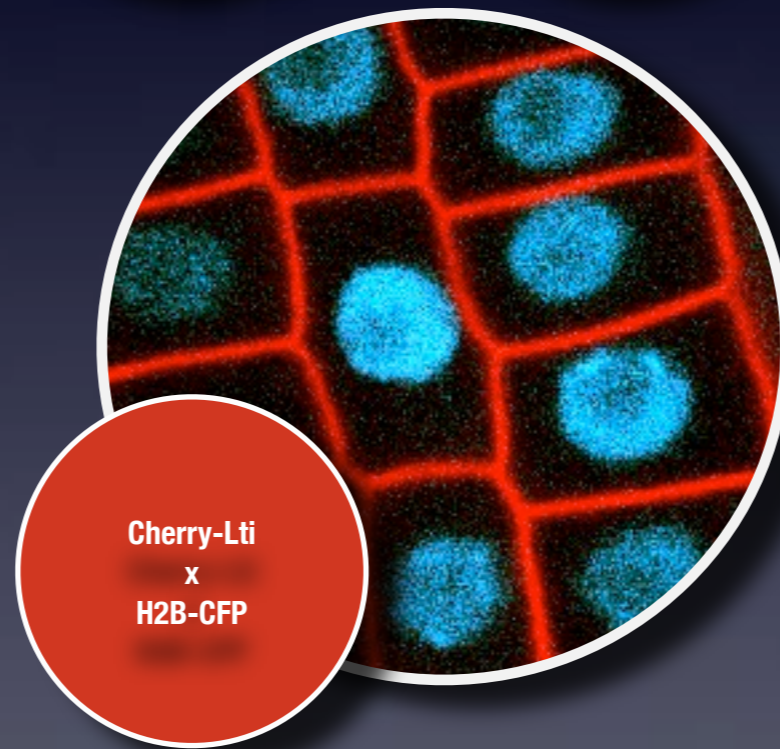
3D optical reconstruction of cellular features in plant tissues.



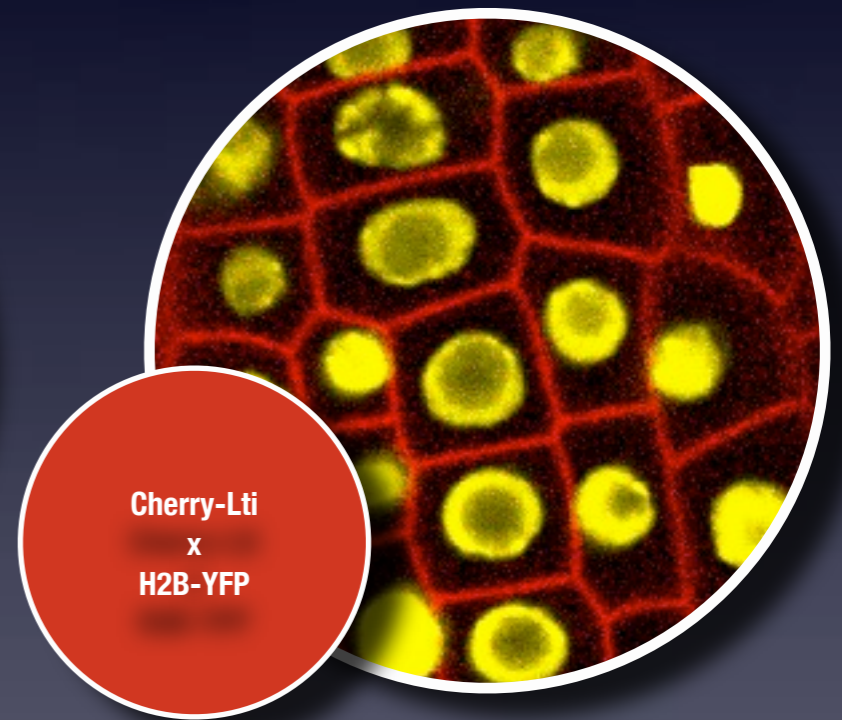
Live imaging of gene expression



GFP-Lti
x
H2B-RFP



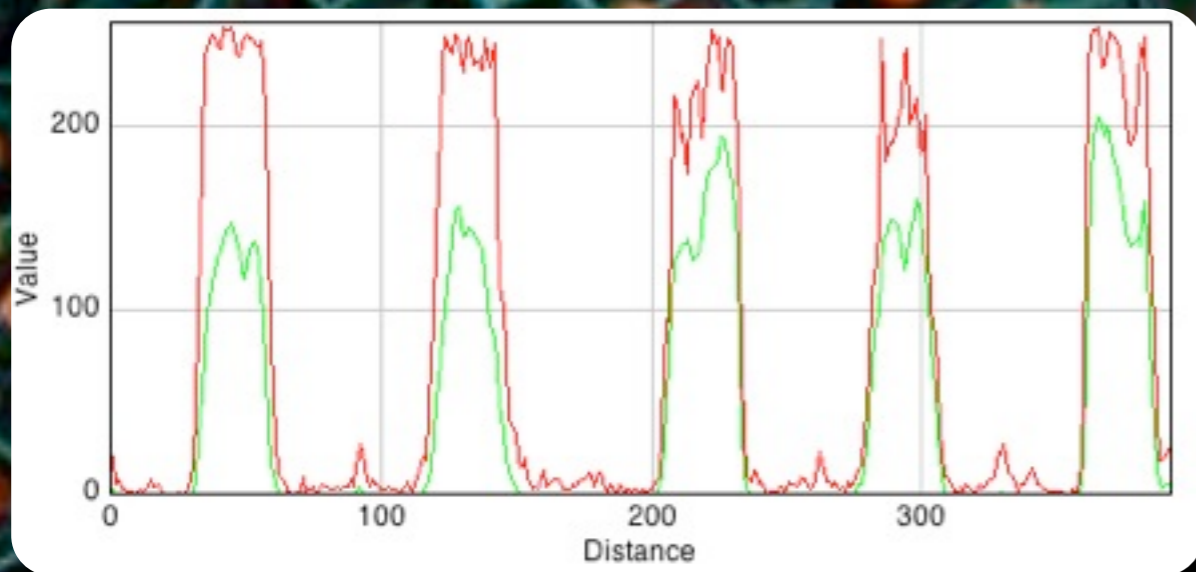
Cherry-Lti
x
H2B-CFP



Cherry-Lti
x
H2B-YFP

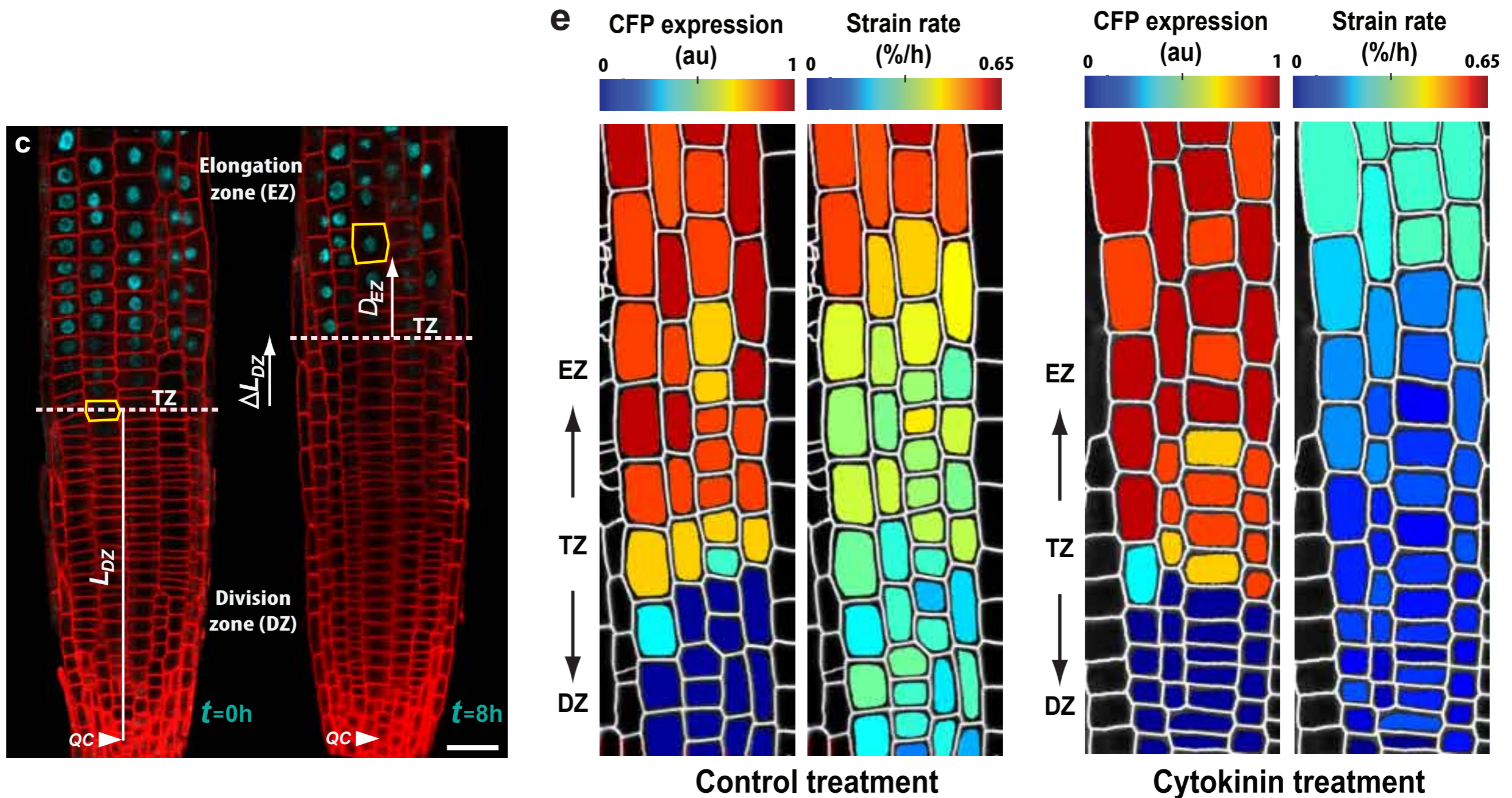
Ratiometric imaging of nuclear localised markers for standardised measurement of gene expression

Lionel Dupuy & Fernan Federici



In planta cytometry

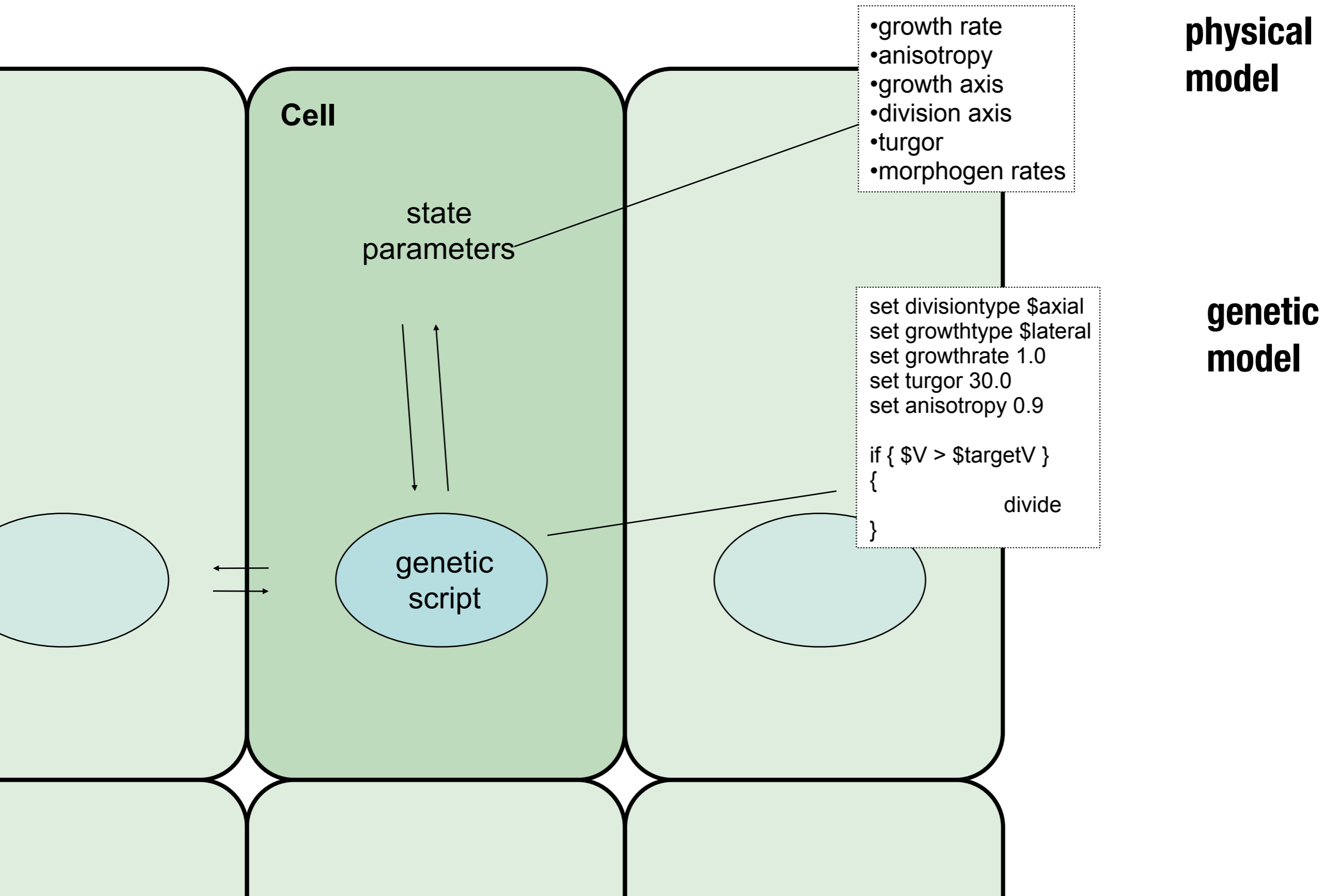
Provides a quantitative view of bacterial biofilms and plant microarchitecture and dynamics of cell shape and gene expression



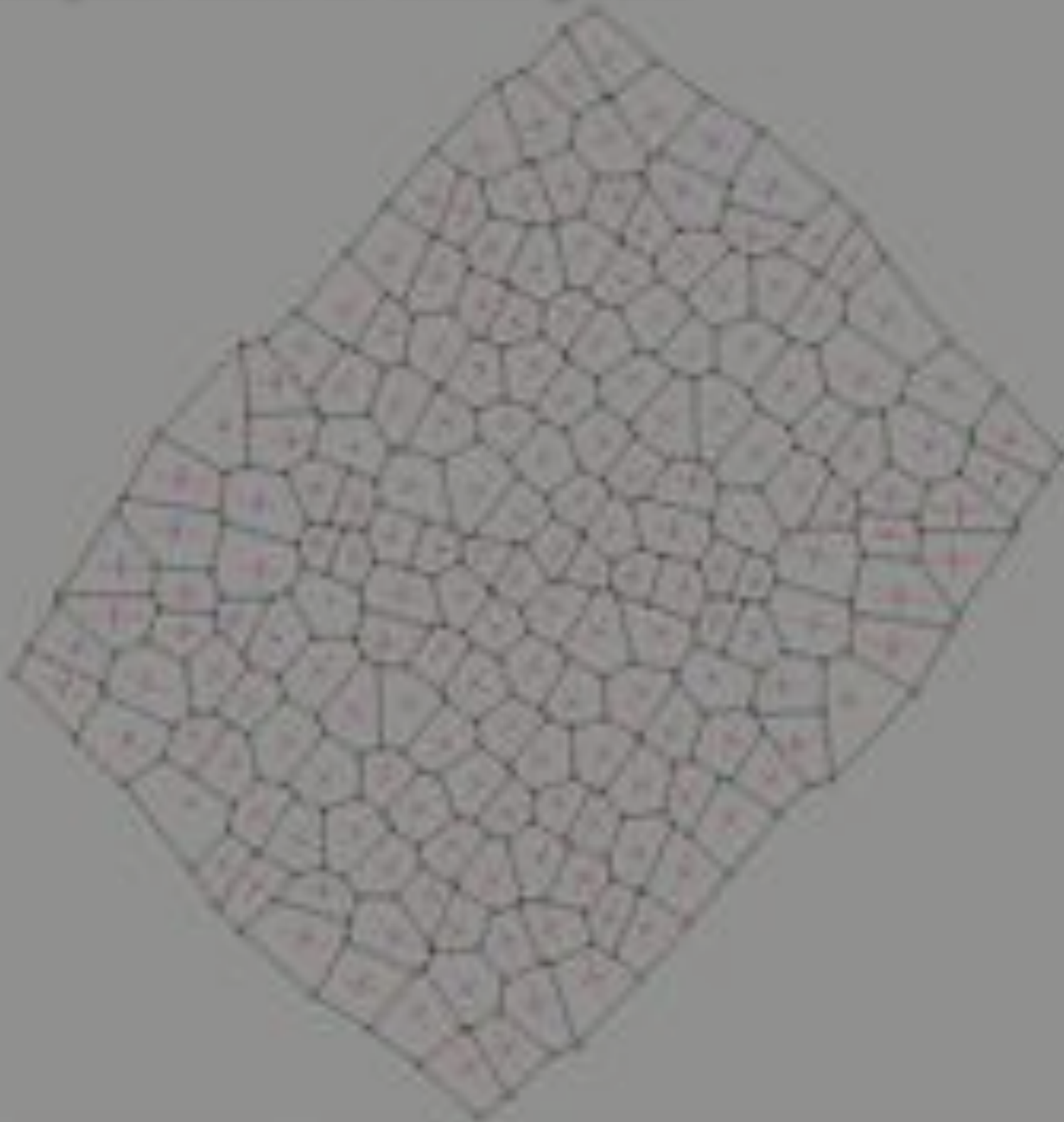
Integrated genetic and computation methods for in planta cytometry, **Nature Methods**, 2012

Fernán Federici, Lionel Dupuy, Laurent Laplaze, Marcus Heisler & Jim Haseloff

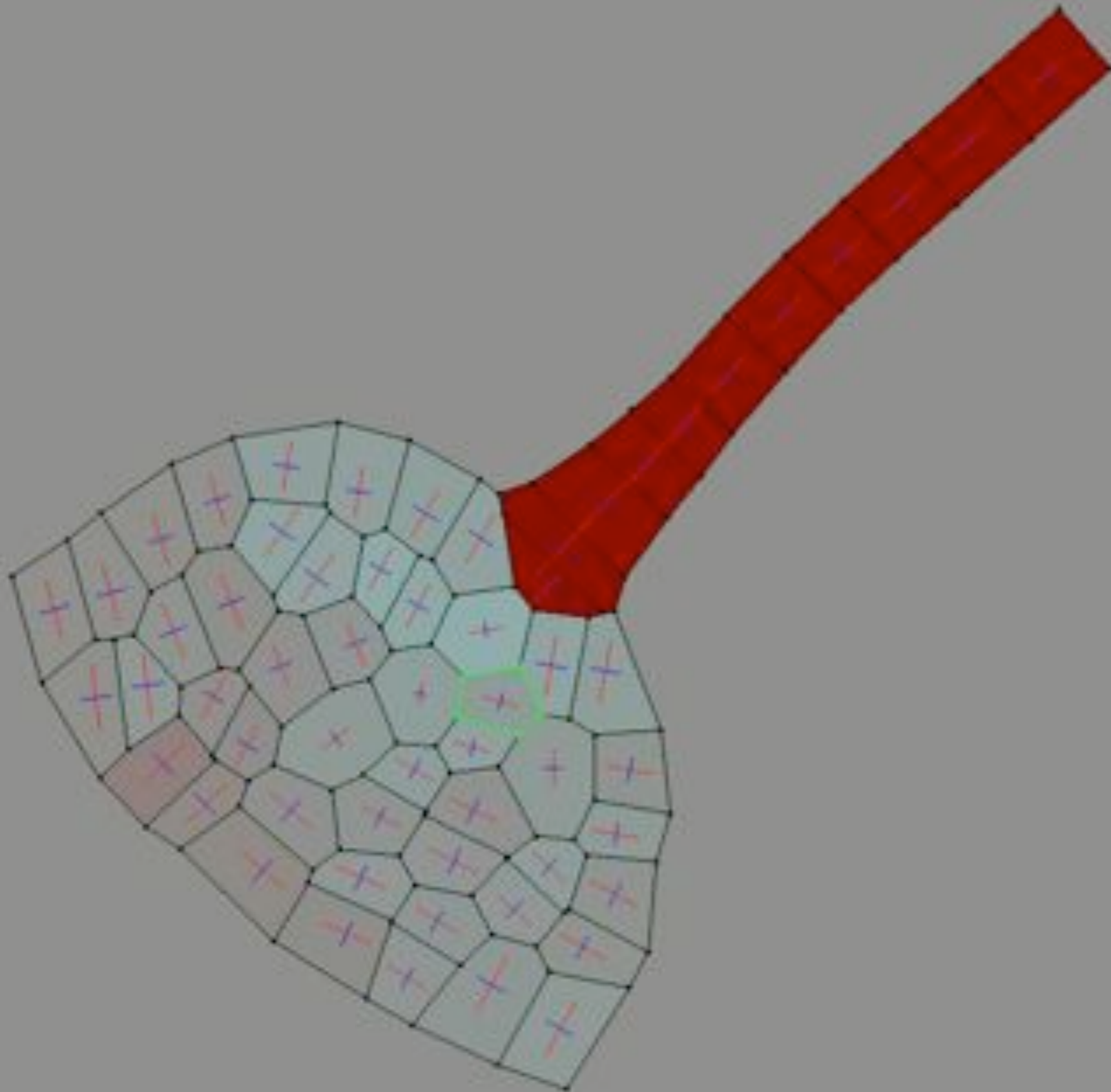
Cellular automata models for plant morphogenesis

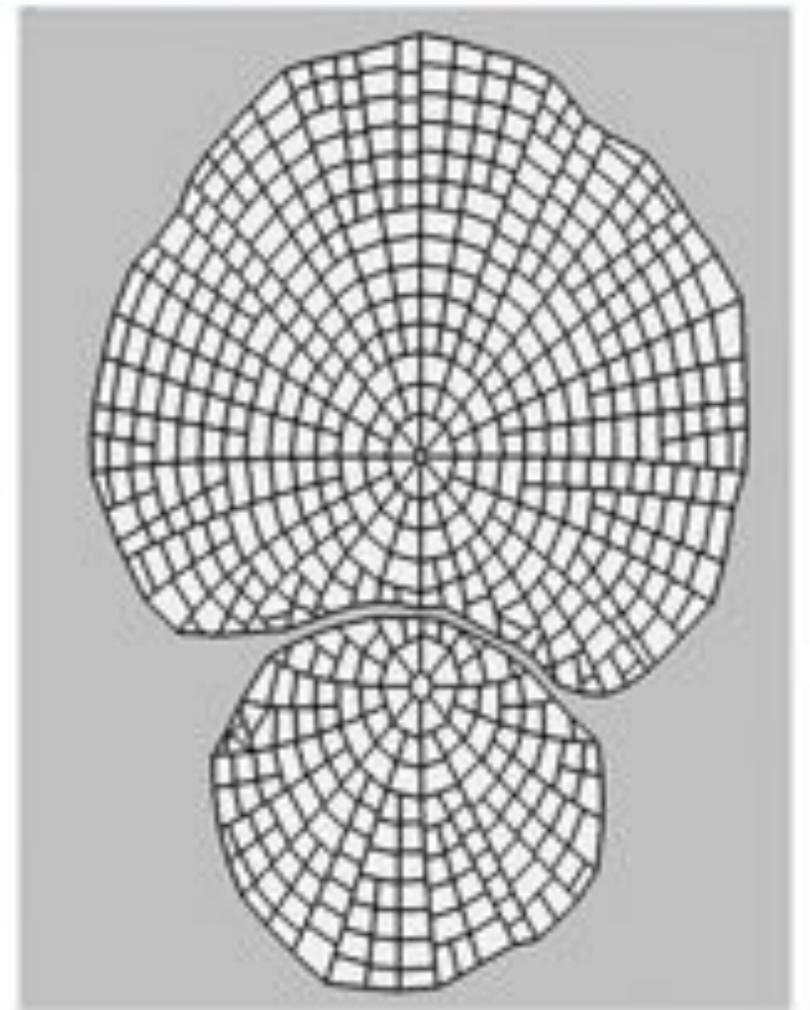
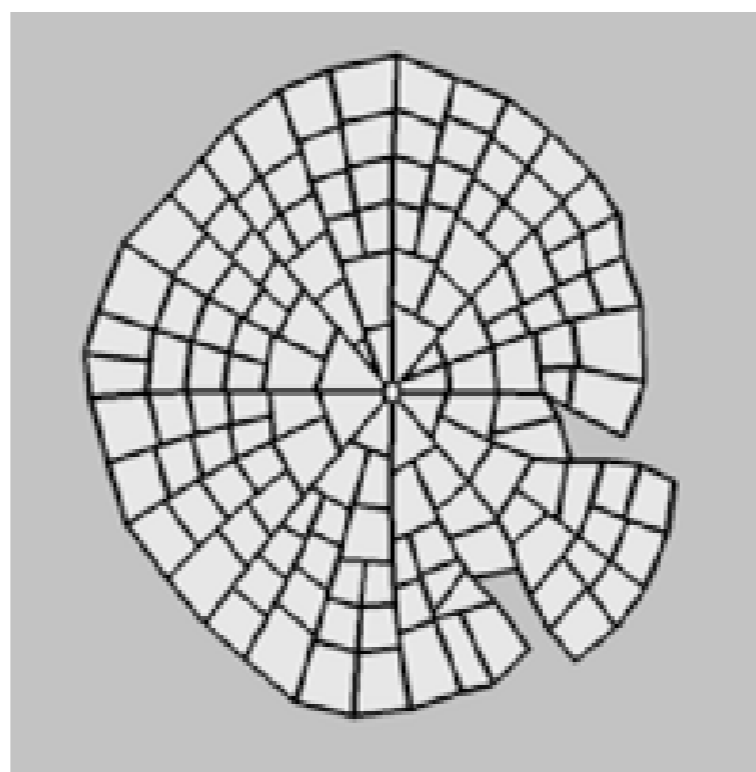
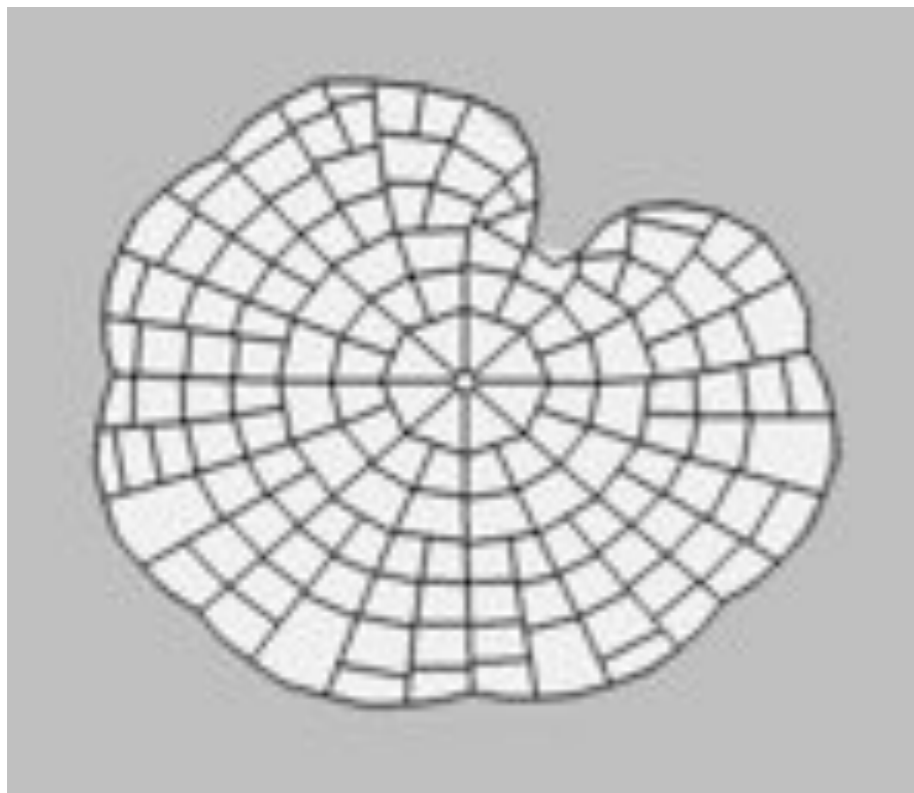
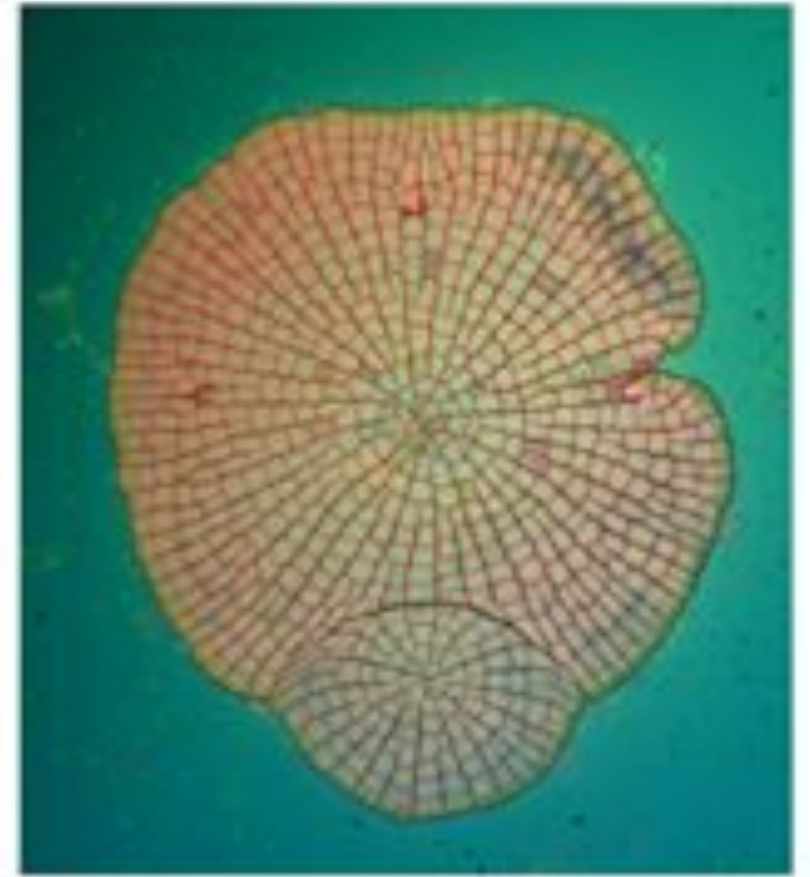
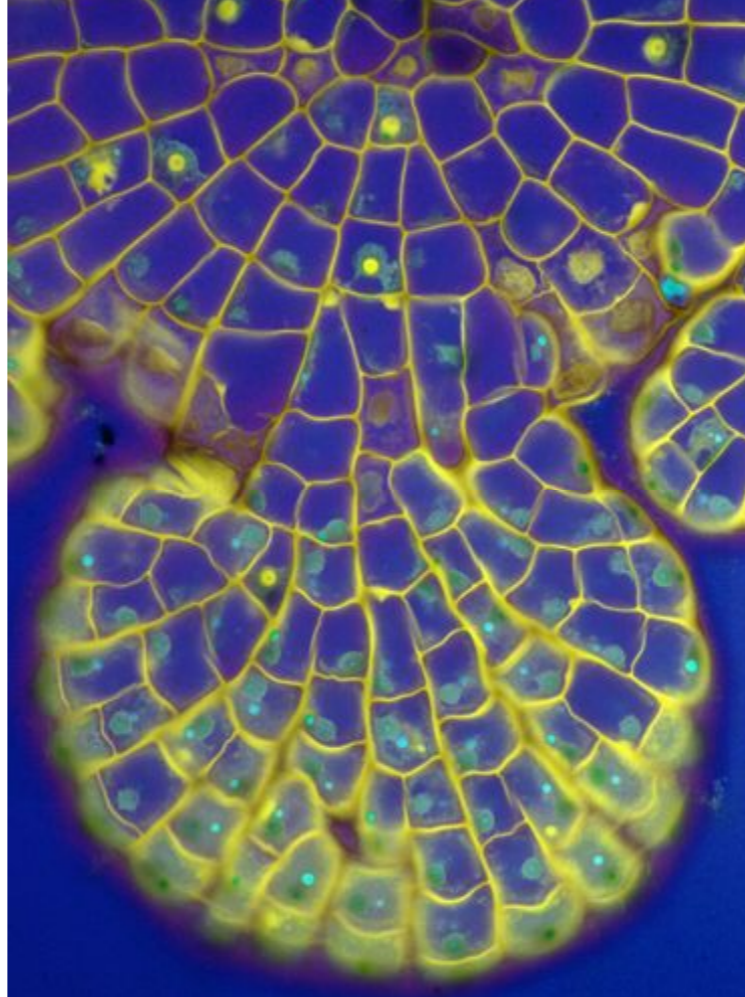


Computer model for cellular growth



Coupling a morphogen to cell proliferation





Modelling growth of Coleochaete

Testbed for synthetic biology in plants



Marchantia polymorpha

Testbed for synthetic biology in plants

Marchantia polymorpha

- Thalloid liverwort
- Descendant of earliest land plants
- Gametophyte dominant phase of lifecycle
- Haploid genetics
- Vegetative propagation by gemmae
- Easily propagated *in vitro*
- Gametes induced by far-red light
- Crossing easy
- Spores stable for >1 year
- EST collection available
- Easily regenerates in tissue culture
- High frequency transformation
- Y chromosome and plastid genomes seq'd
- 280 MB genome sequence due soon

www.marchantia.org



Marchantia has a highly simplified genome

Gene Family	<i>Arabidopsis</i>	<i>Selaginella</i>	<i>Physcomitrella</i>	<i>Marchantia*</i>
ARF	23	7	14	3
AUX/IAA	29	3	2	1
TIR1	6	2	4	1
TPL	5	3	2	1
TAA	3	1	4	1
GH3	19	17	2	1
Class I HD-Zip	17	4	17	1
Class II HD-Zip	10	2	7	1
Class III HD-Zip	5	3	5	1
Class IV HD-Zip	16	4	4	1
Class I KNOX	4	2	3	1
Class II KNOX	4	3	2	1
BELL homeobox	13	2	4	1
WOX	16	6	3	1
CLV3 (CLE)	28	6	4	1

*Based on EST data (>2,300,000) represents a minimum number

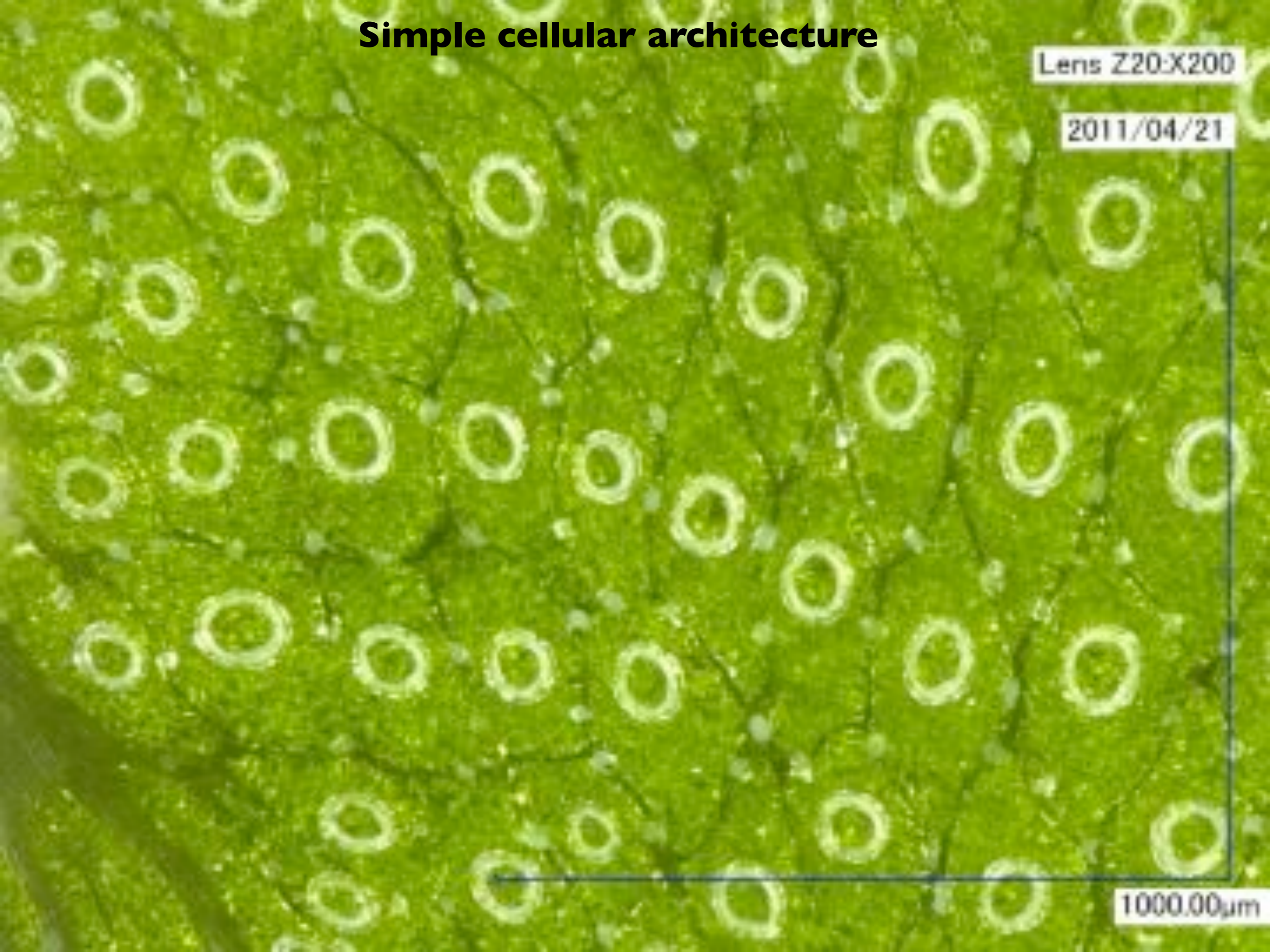
Hiroataka Kato, Kimitsune Ishizaki, Takayuki Kohchi
Grad. Sch. of Biostudies, Kyoto Univ.

In collaboration with J. Bowman's lab, Monash

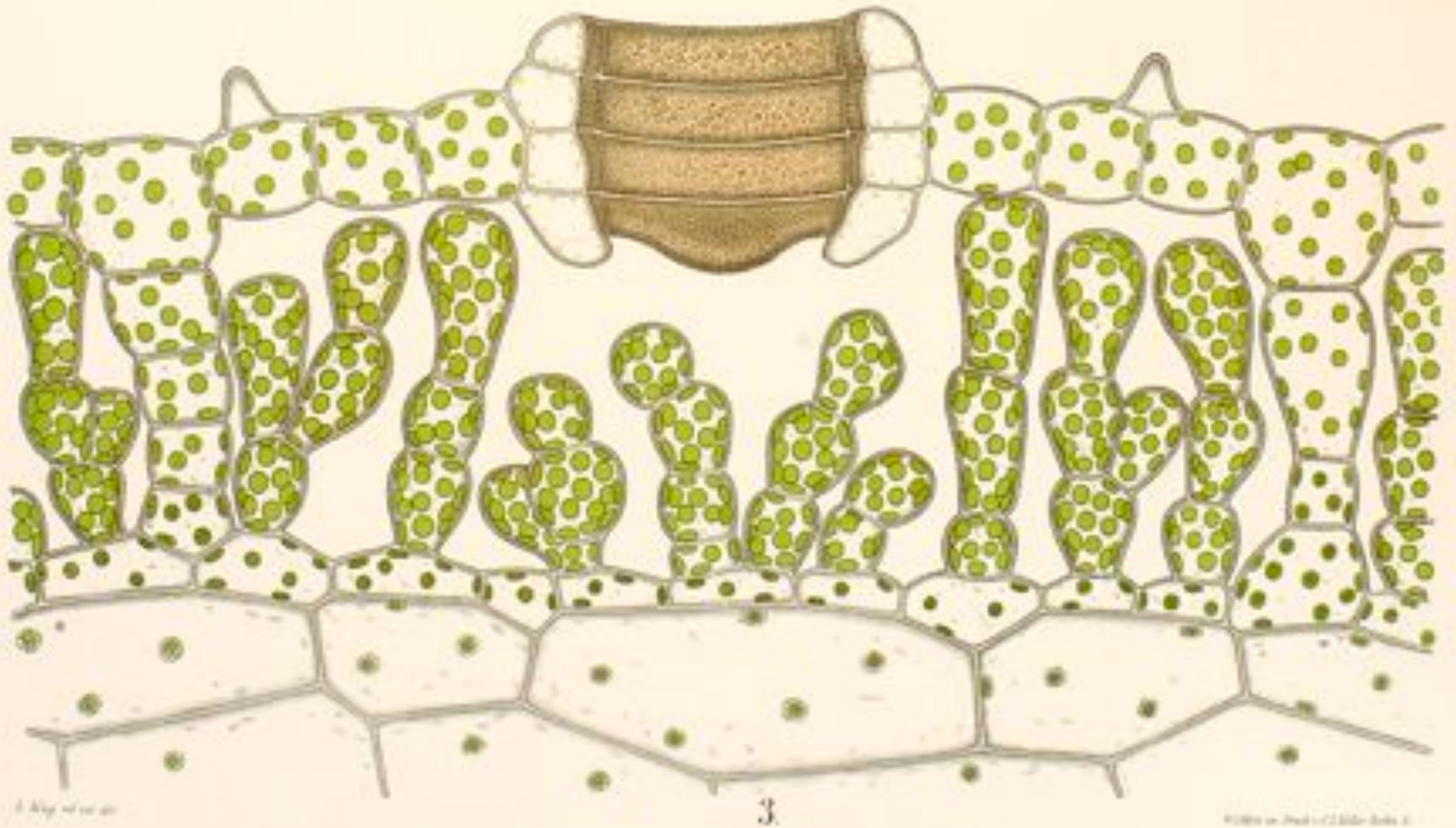
Simple cellular architecture

Lens Z20:X200

2011/04/21



1000.00µm



Transverse section of *Marchantia polymorpha* thallus (Leopold Kny)



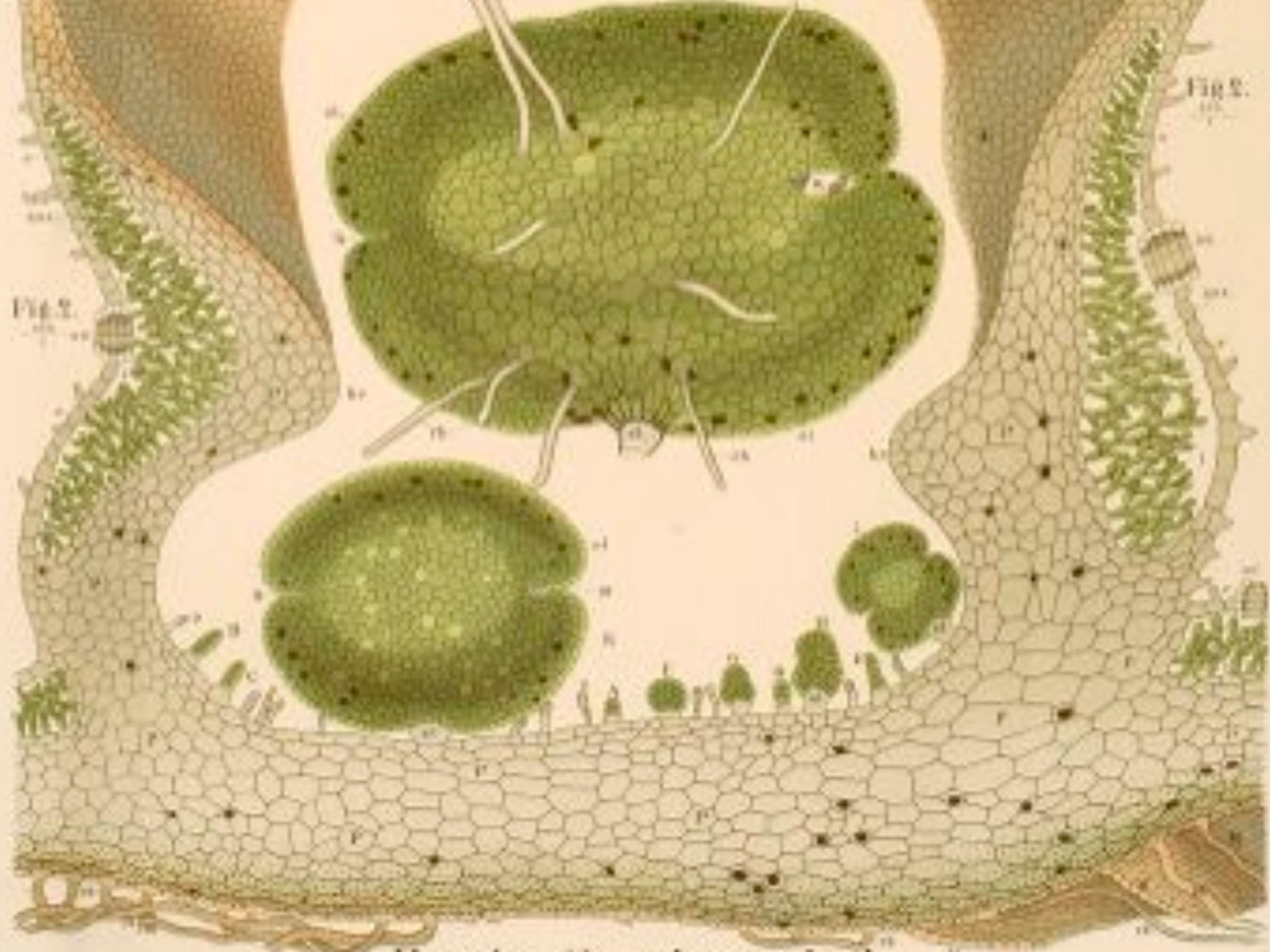
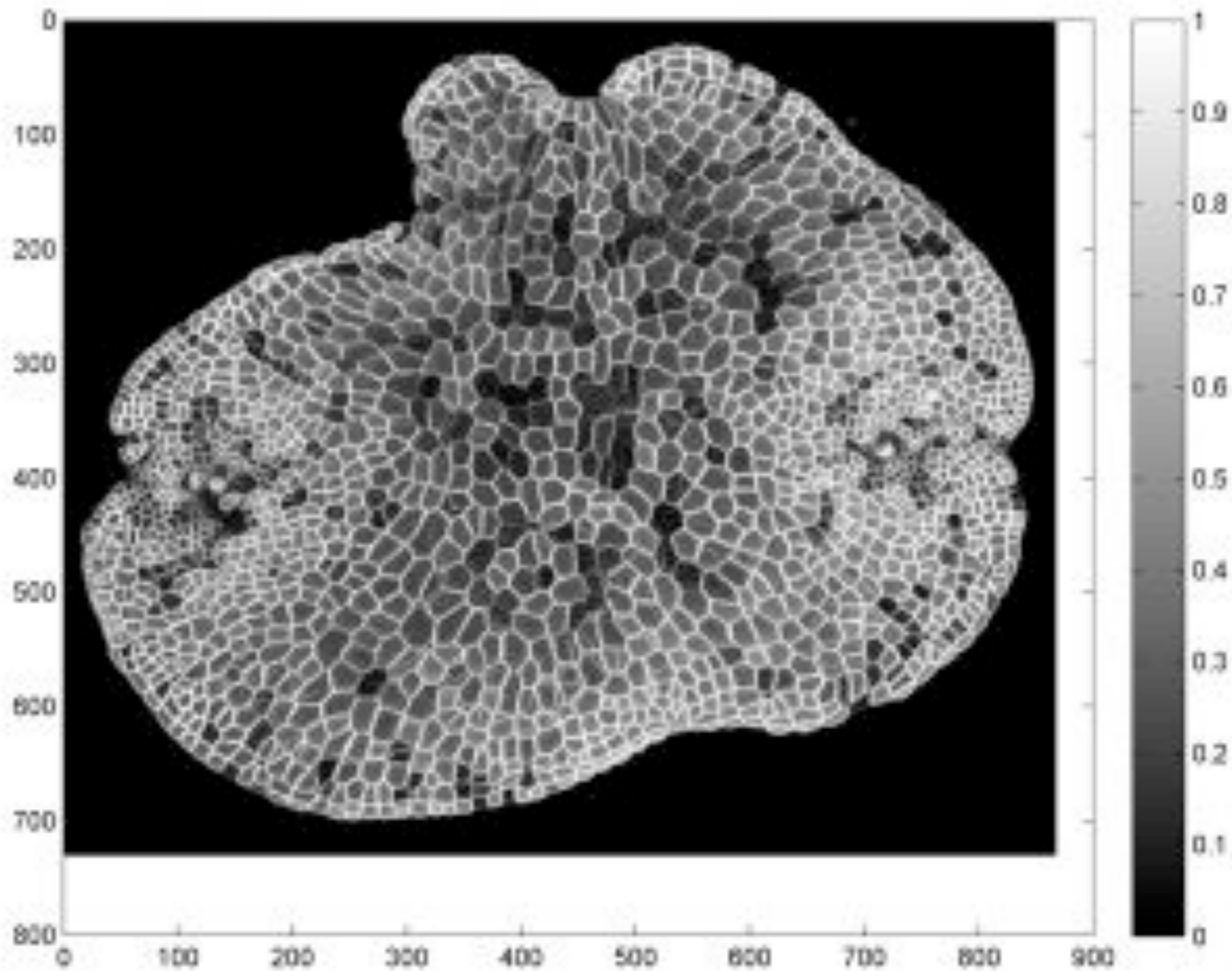


Fig. 2.

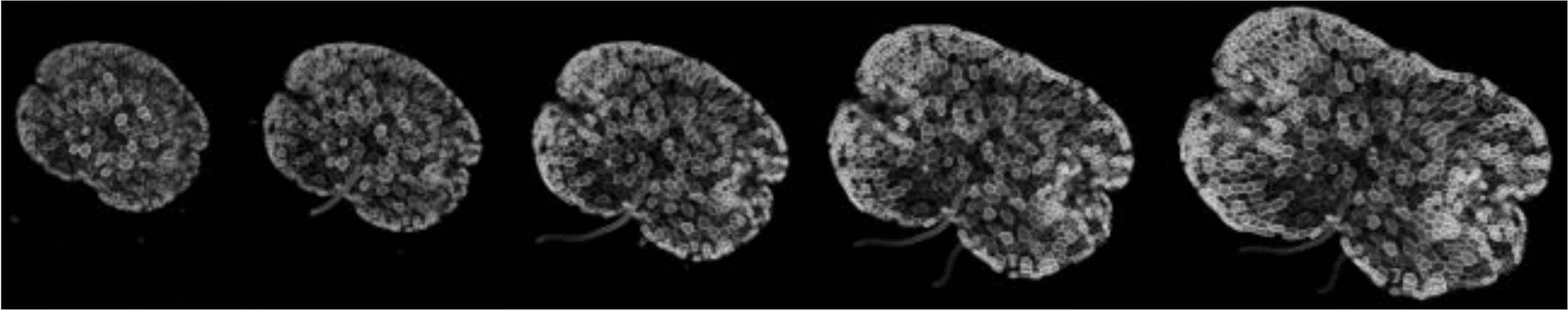
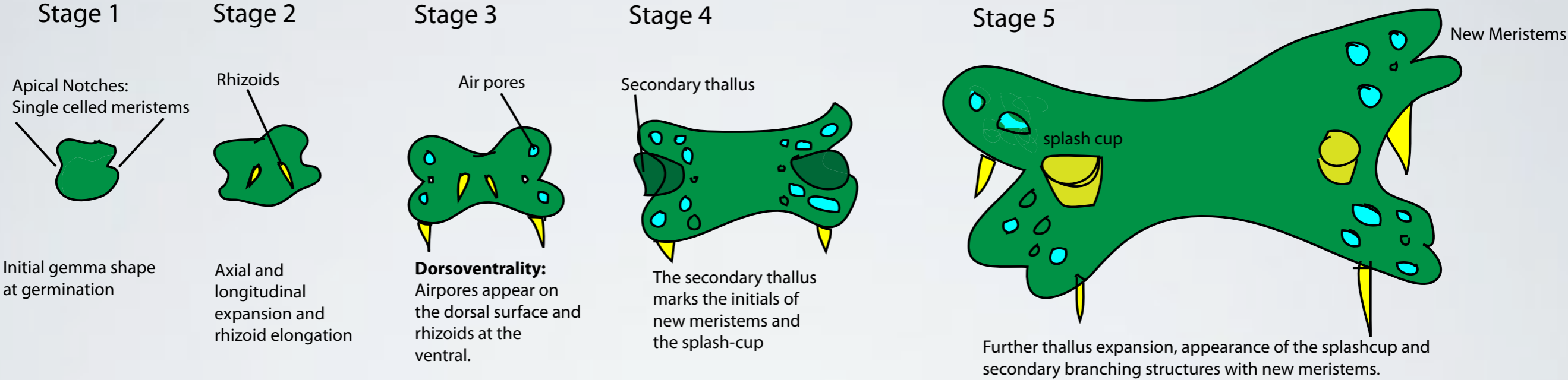
Fig. 3.

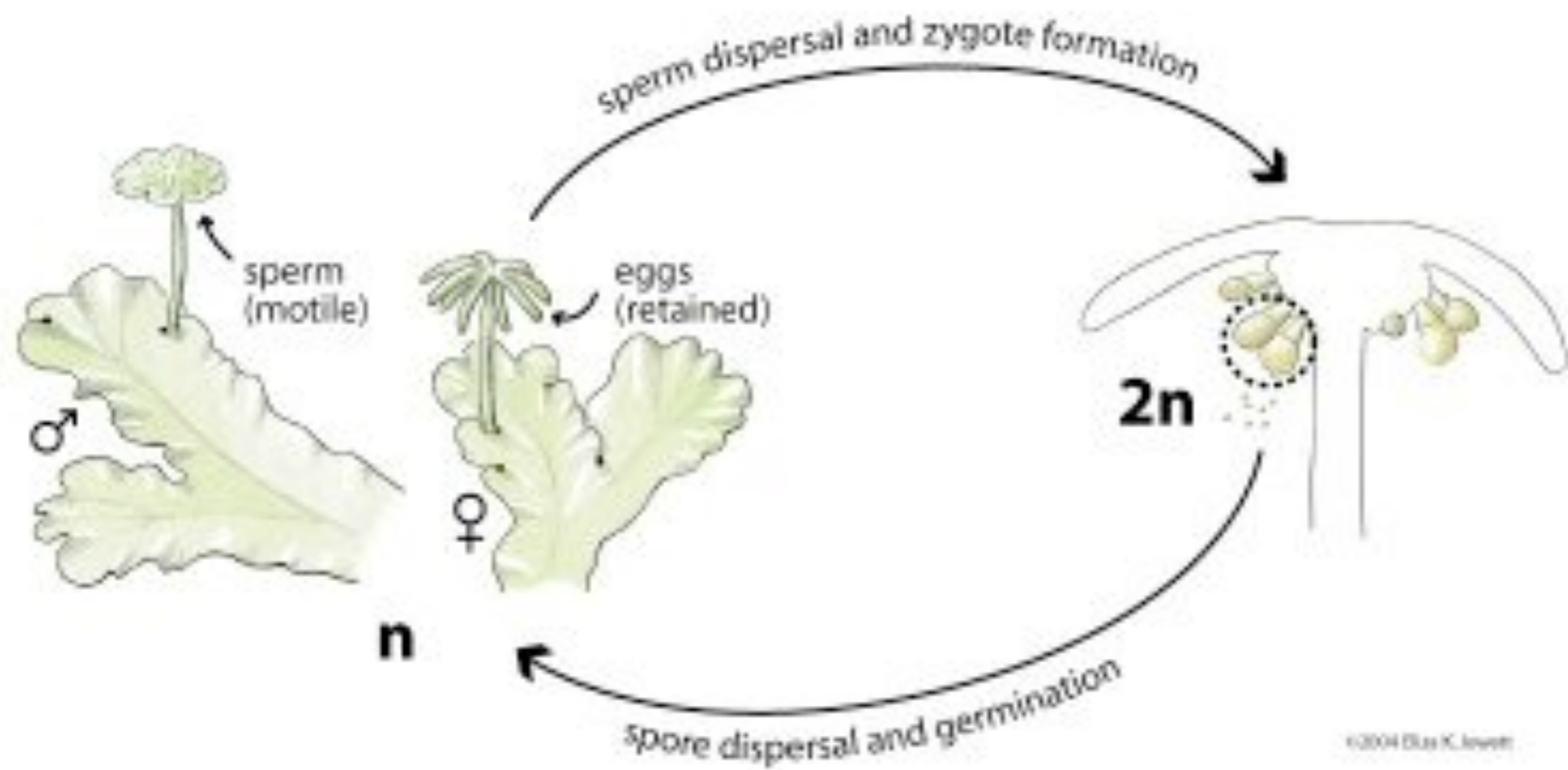
Marchantia polymorpha, L.

High resolution imaging of gemma



Staging of Marchantia gemma development

















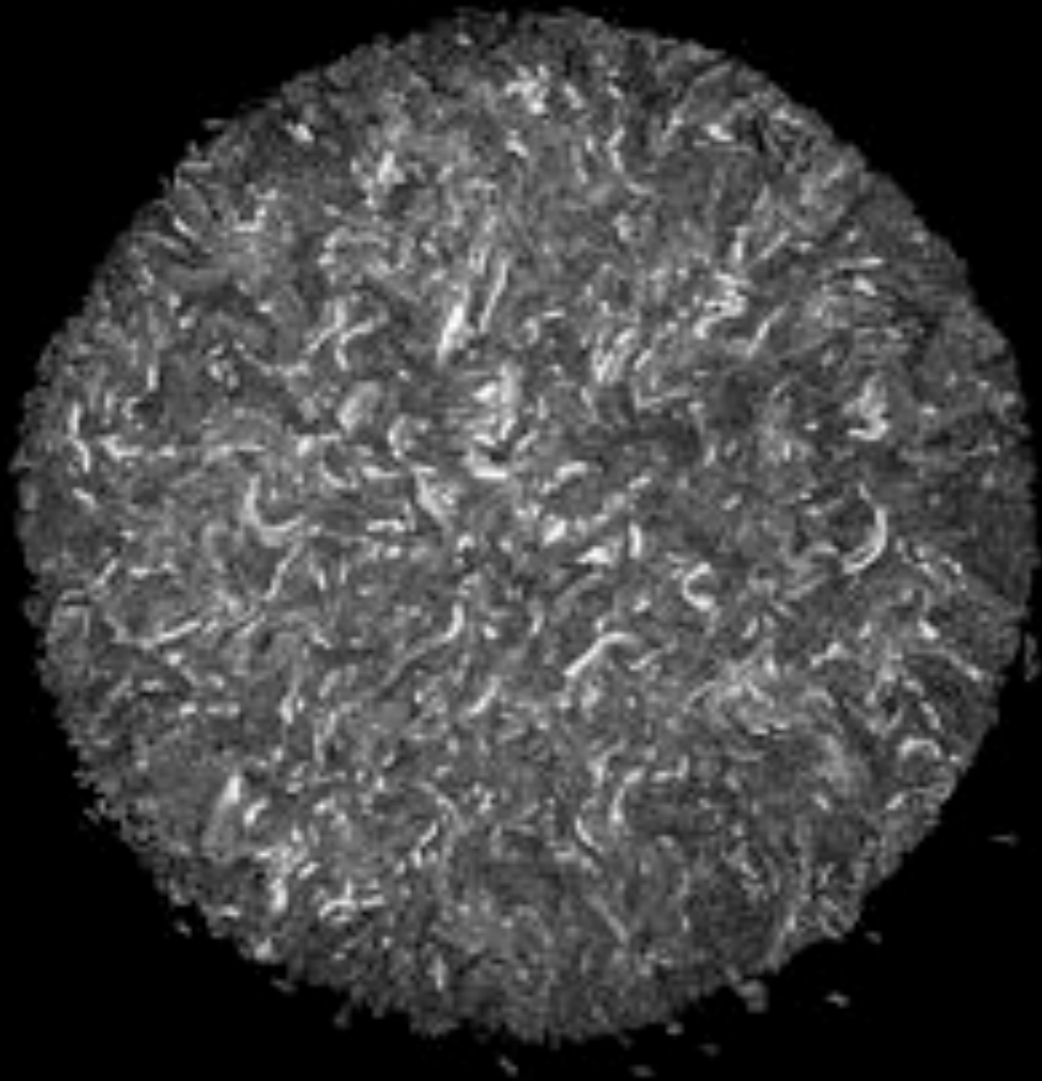




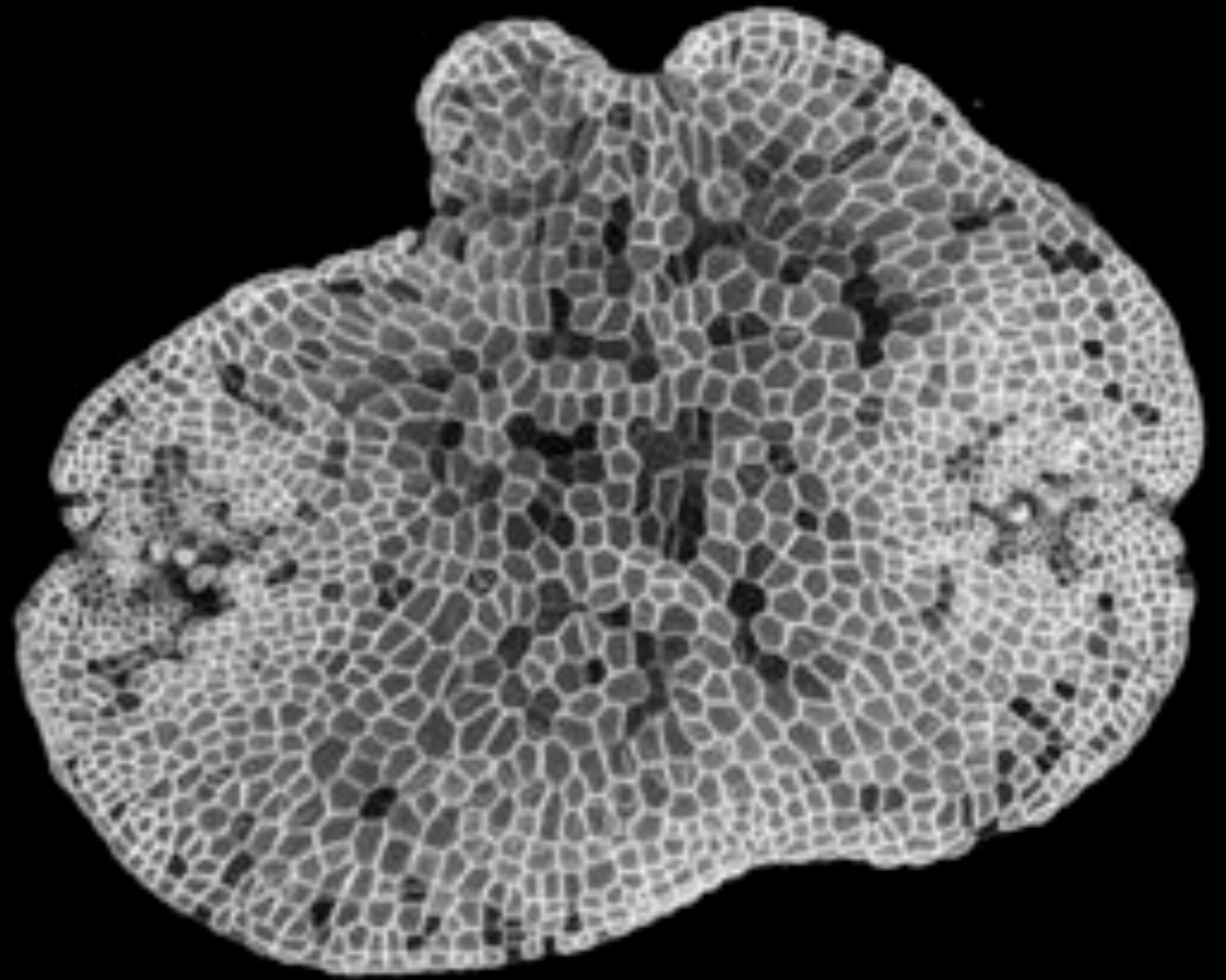


Model systems for engineering patterns in cell populations

microbes



plants



Rules and systems for engineering cell populations in plants

We need a new generation of feedback regulated genetic circuits with the ability to organise and coordinate cell cohorts.

GENERATION 1

promoter::gene

GENERATION 2

GAL4 > gene array

GENERATION 3

Turing-type population switch

Modern crop plants are derived from their natural ancestors by thousands of generations of selection and breeding.

What if we could reprogram the distribution of existing cell types in living systems?



James Brown



microbial engineering

PJ Steiner



Paul Grant



Nuri Purswani



Fernan Federici



Li Hua Robertson



plant morphogenesis

Tim Rudge



Michael Pedersen



computational biology

plus:

**Judy Savitskaya
Anton Kan
Robyn Cooper**

**Prof Chris Voigt
MIT**

**Dr Lionel Dupuy
SCRI, Dundee**

**Dr Jim Ajioka
Department of Pathology**

**Dr Andrew Phillips
Microsoft Research
Cambridge**

**Dr Neil Wipat
University of Newcastle**