

Controlling the lymphocyte - empirical rules for a calculus of signal integration

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KITP Santa Barbara, 2012



Walter+Eliza Hall
Institute of Medical Research

Adaptive immunity: Complexity at multiple scales

Movies from WEHI-TV - see Fighting infection
by Clonal Selection - [http://www.youtube.com/
watch?v=HUSDvSknlgl](http://www.youtube.com/watch?v=HUSDvSknlgl)

Etsuko Uno & Drew Berry WEHI-TV

Macfarlane Burnet



1965

*“The principles of immunity
are now known*

*Immunology is now just
working out details”*

Macfarlane Burnet



1965

*“The principles of immunity
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*Immunology is now just
working out details”*

The details were hard!

Specificity

Features of antibody

Class regulation

Self tolerance

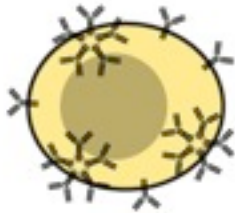
Memory

B cells play a key role in the humoral immune response

Antibody-producers

- Neutralisation
- Opsonisation

- Activation of complement
- Induction of phagocytosis



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Differentiate by

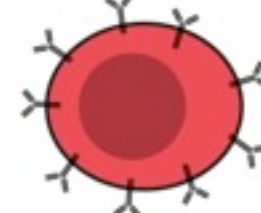
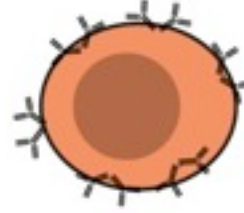
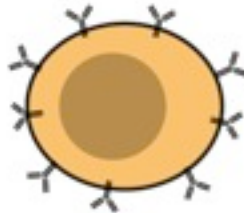
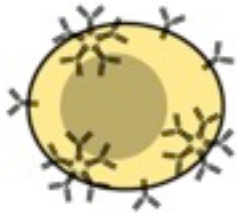
IgM, IgD

IgG

IgA

IgE

isotype
switching



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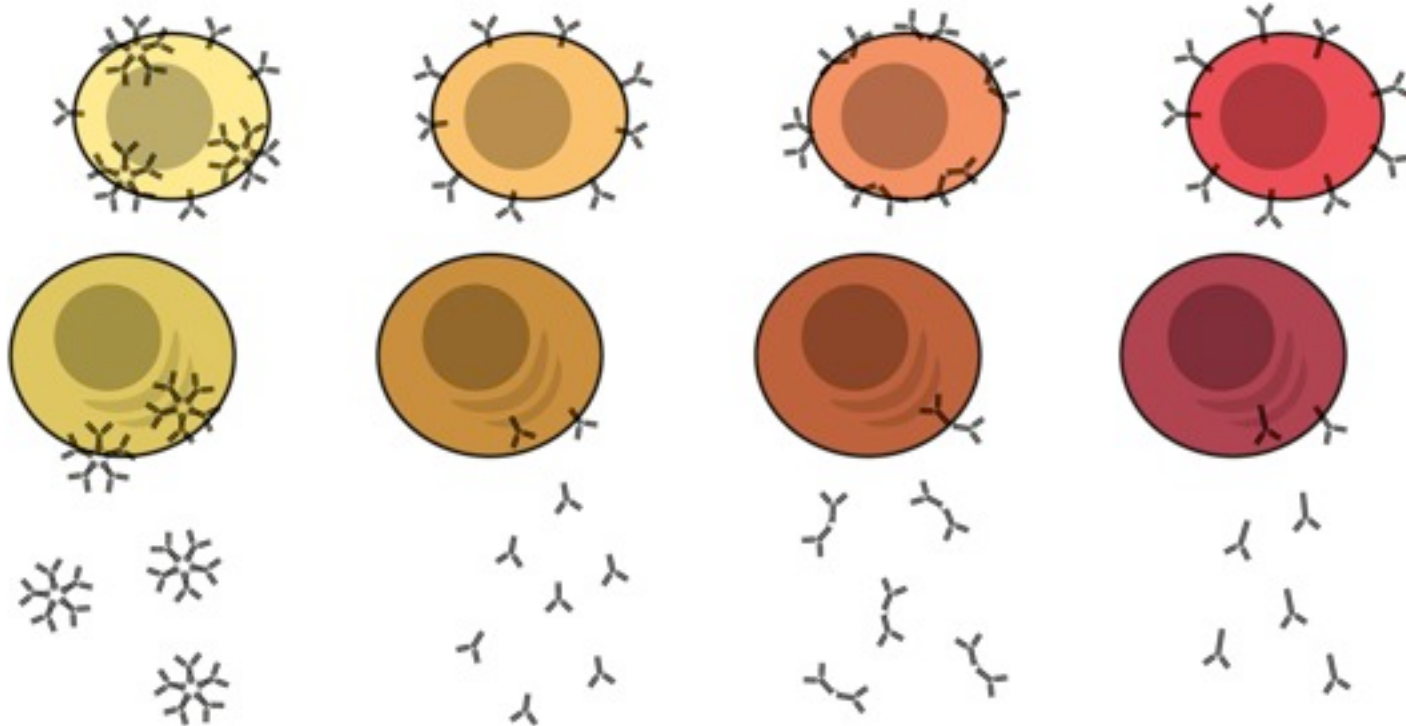
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becoming antibody secreting cells (ASC)

B cells play a key role in the humoral immune response

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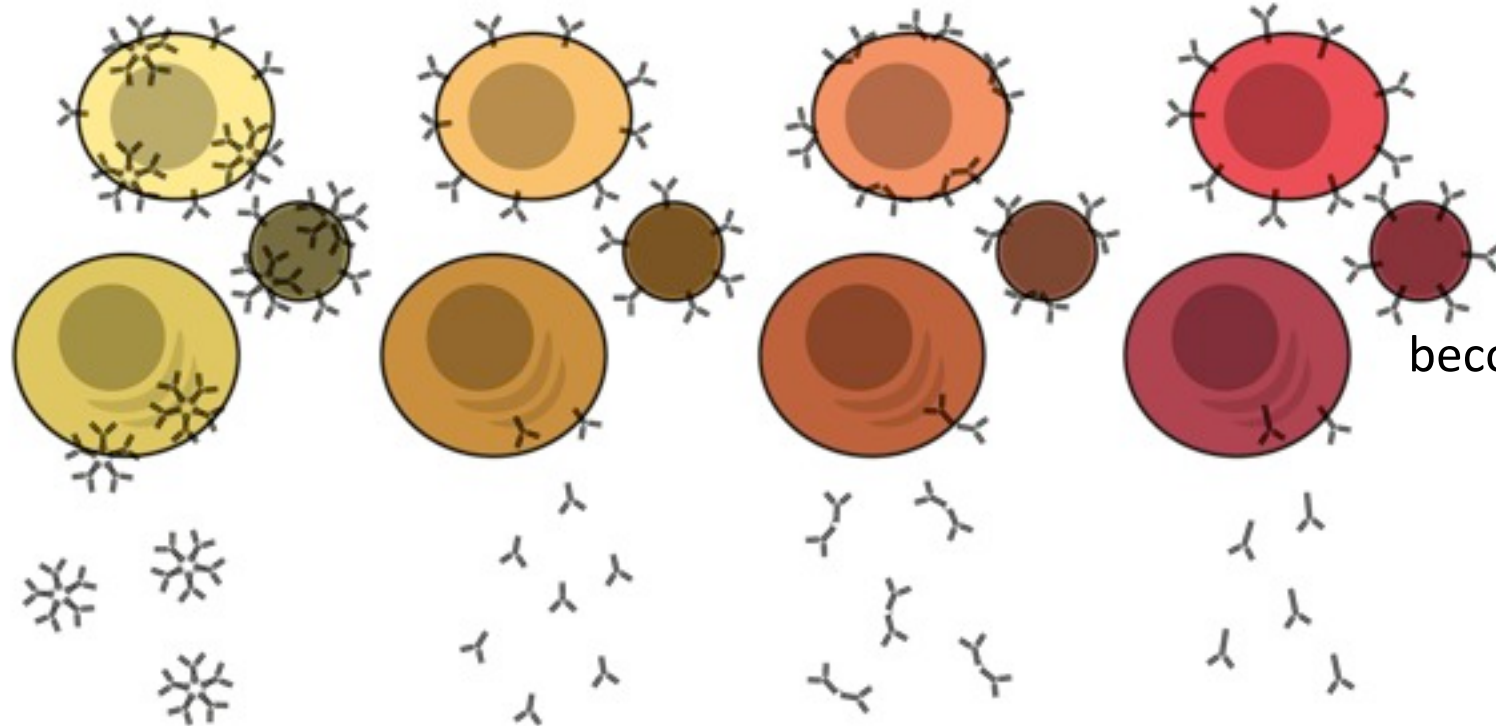
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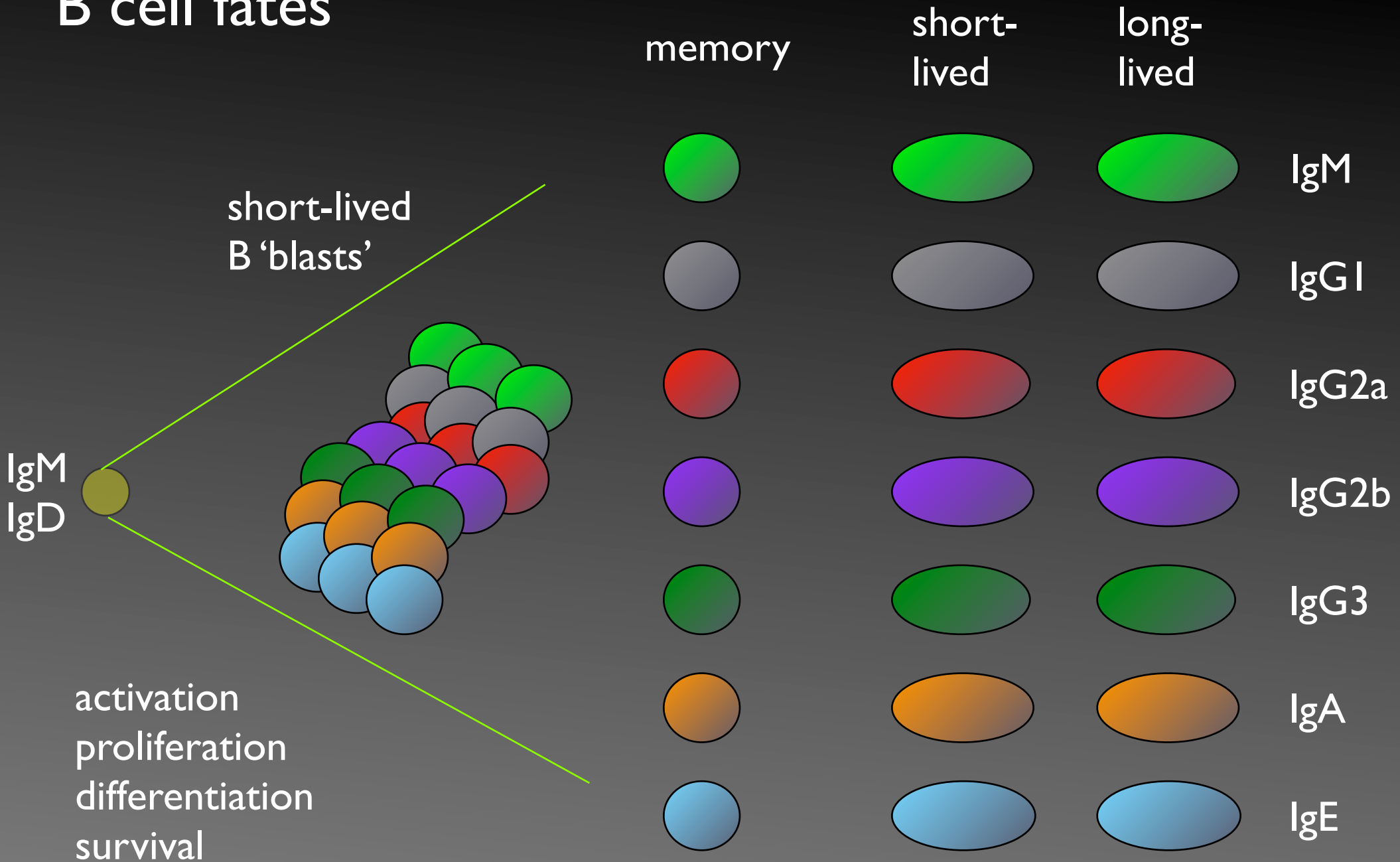
isotype switching



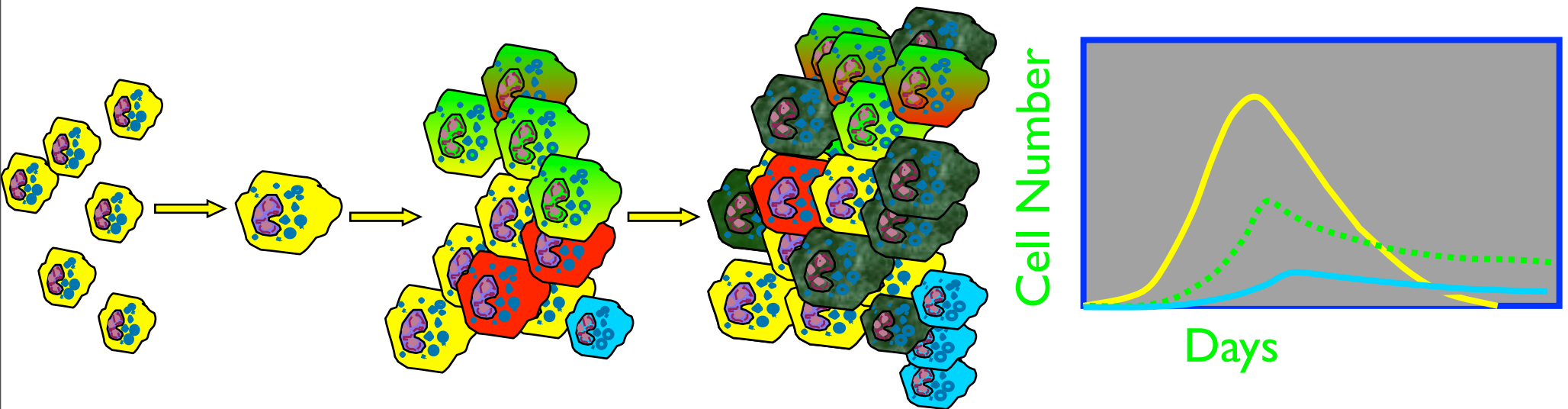
becoming antibody secreting cells (ASC)

becoming memory cells

B cell fates

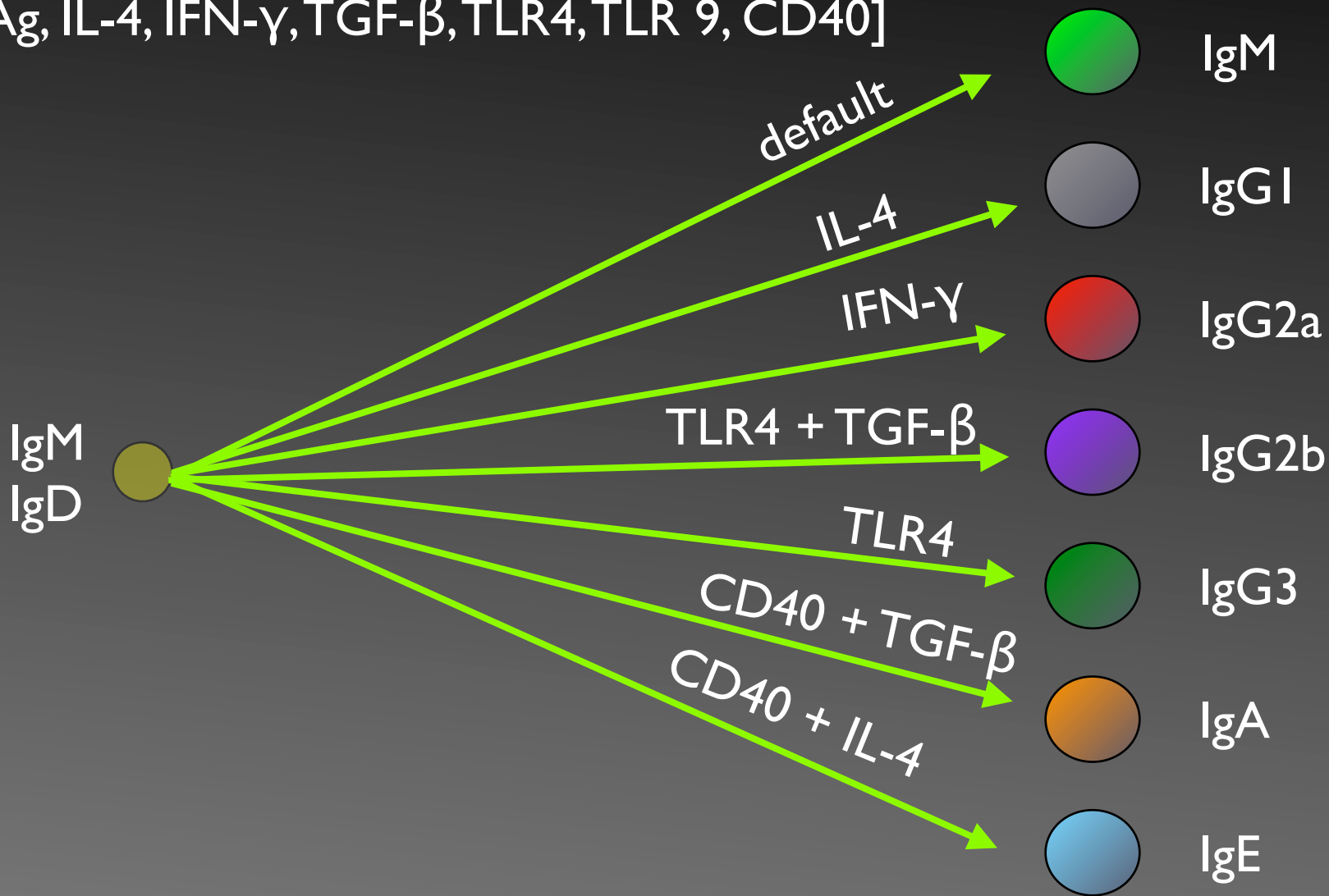


- Solve the logic puzzle - which signals and in what combinations lead to which outcomes?

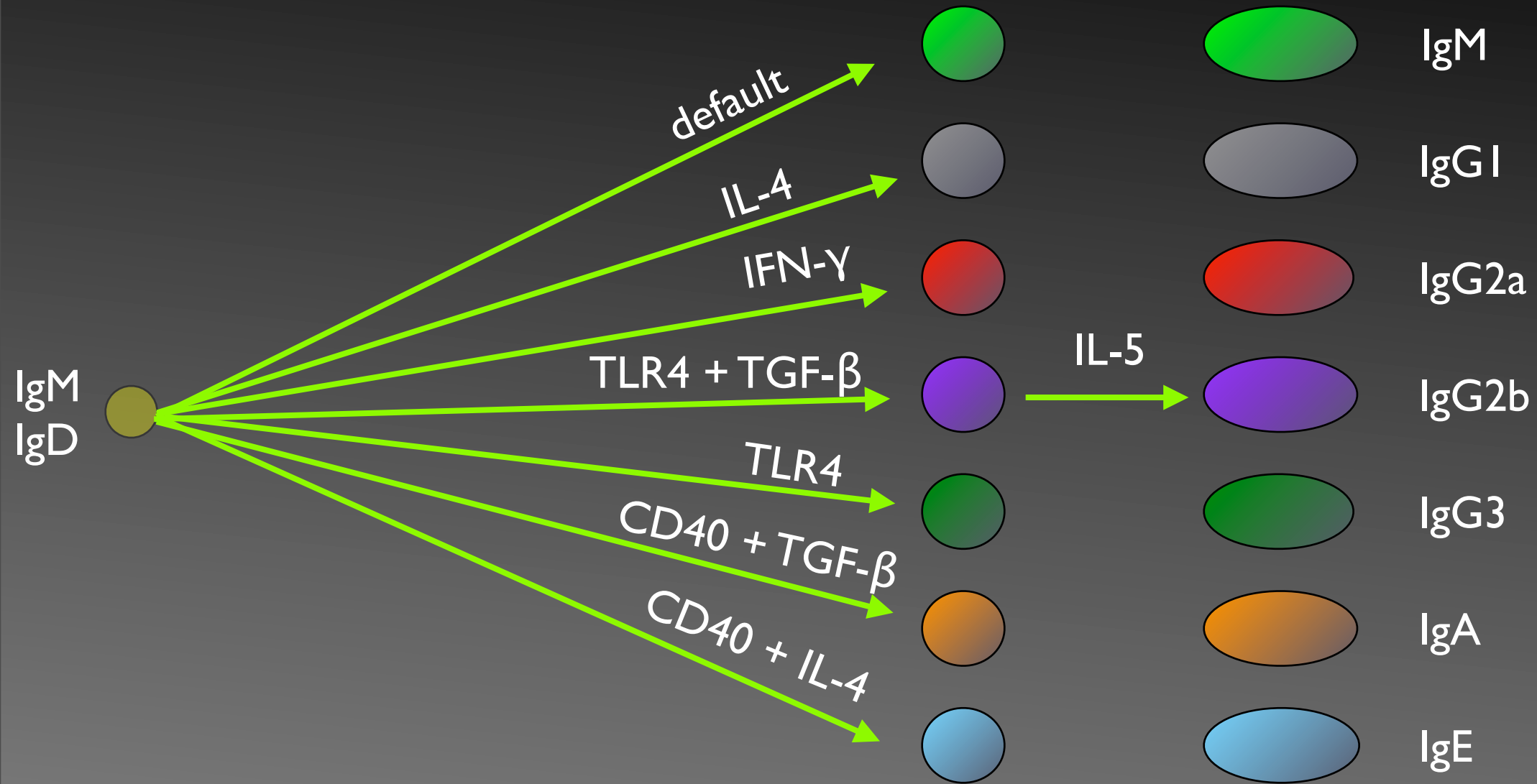


Generation of diversity as a Cellular journey: A logic problem

activation [Ag, IL-4, IFN- γ , TGF- β , TLR4, TLR 9, CD40]



ASC



Complex problem

Typically cytokines have varied effects on proliferation, survival and differentiation

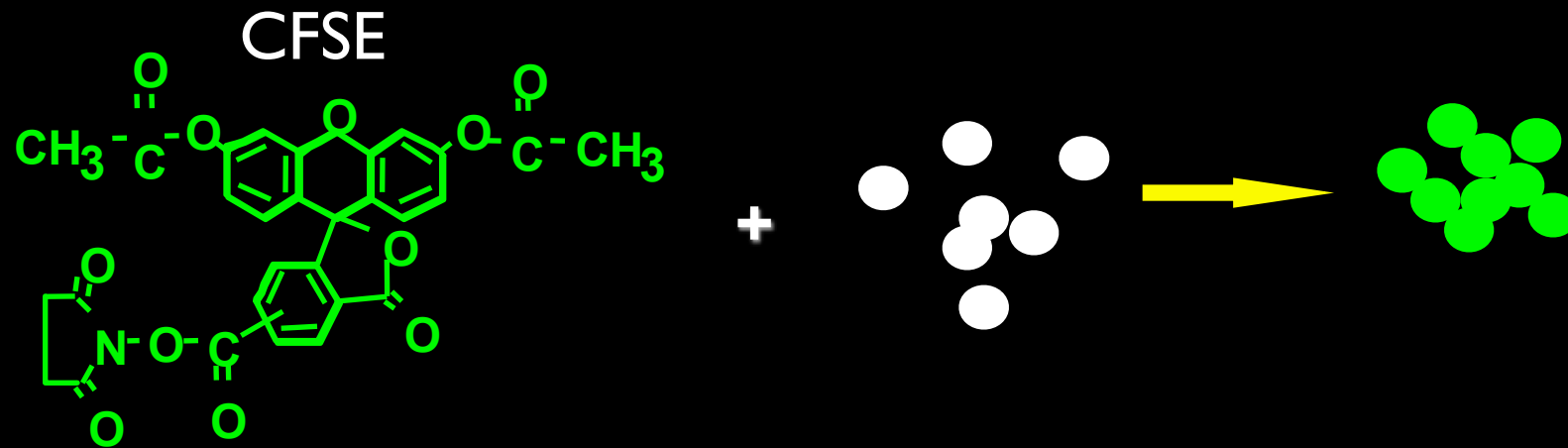
When exposed to combinations of different signals - how do cells calculate an outcome?

(search for principles of a 'Cellular Calculus' - governing signal 'integration' and cell 'differentiation') -

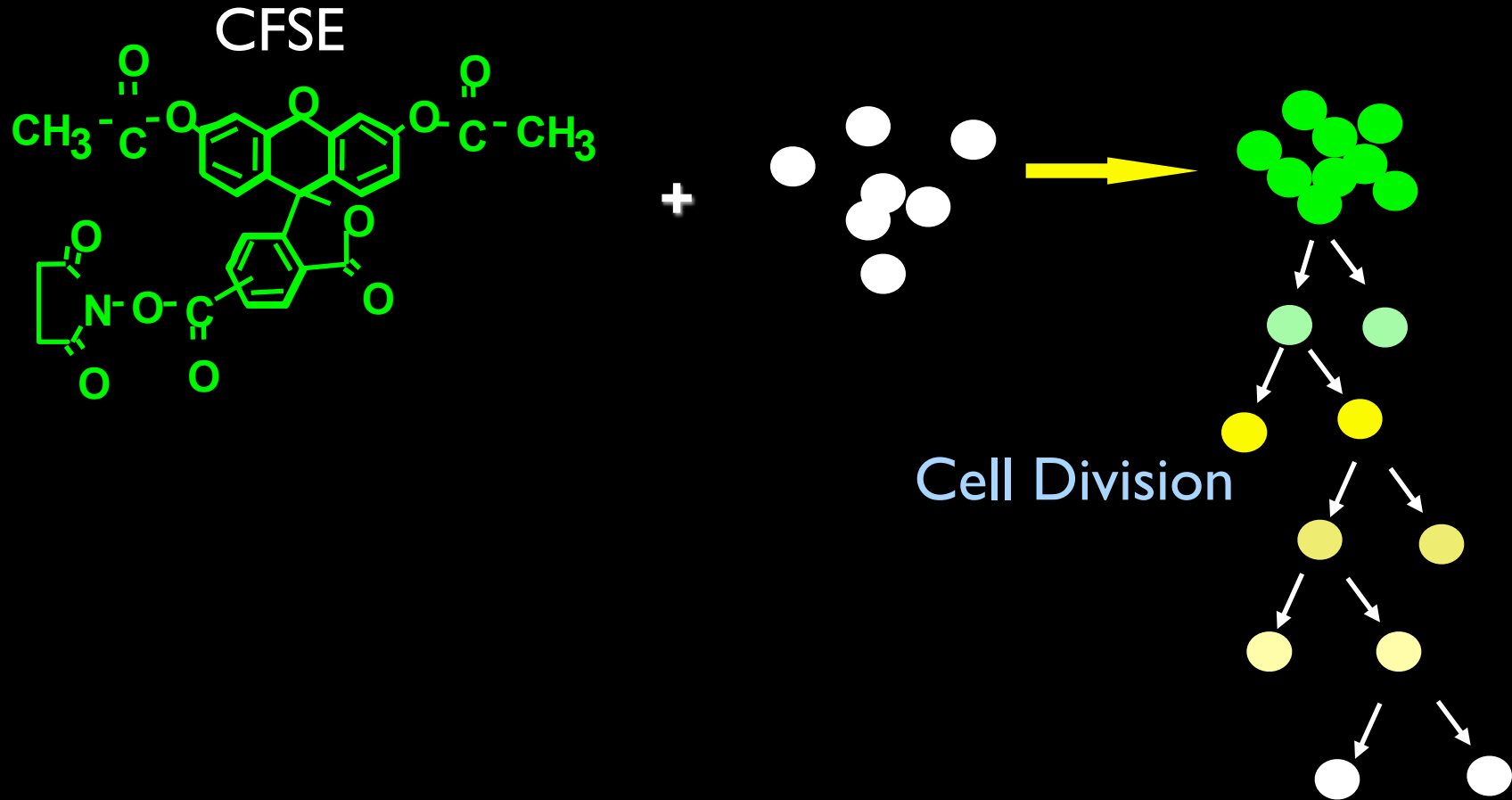
[note - strategy FIRST ORDER (no other interactions) first]

Cell Division number as a hidden variable in differentiation outcomes

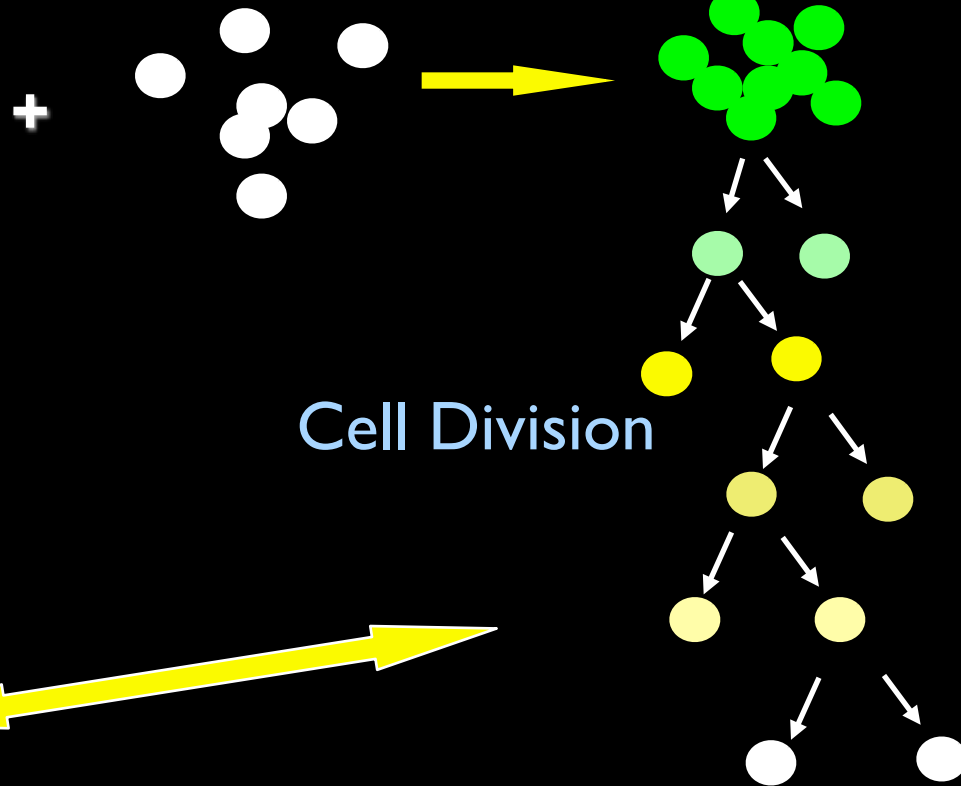
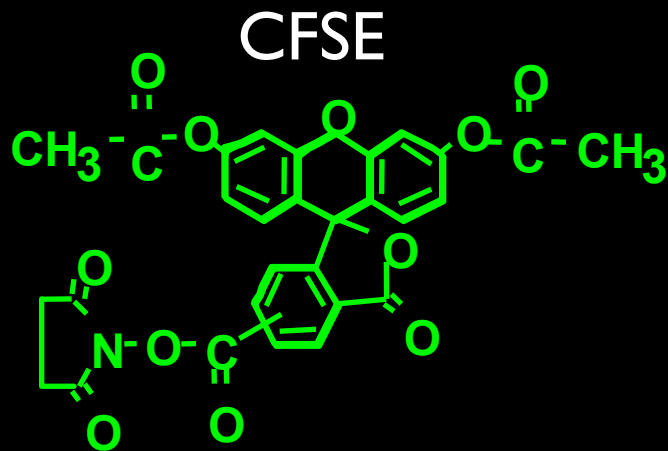
Division tracking with Carboxyfluorescein succinimidyl ester (CFSE) :Lyons and Parish



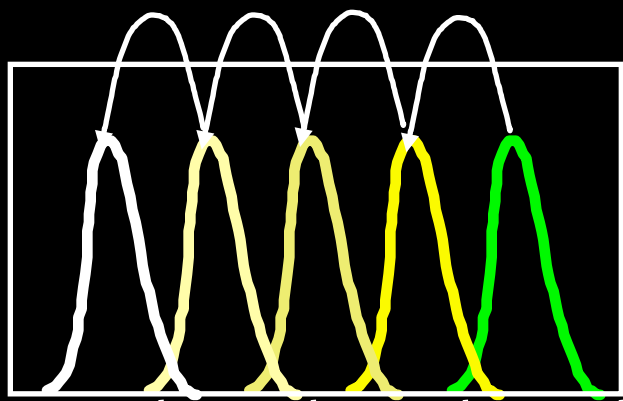
Division tracking with Carboxyfluorescein succinimidyl ester (CFSE) :Lyons and Parish



Division tracking with Carboxyfluorescein succinimidyl ester (CFSE) :Lyons and Parish



Cell Division



CFSE

Flow cytometer

This method allows proliferation, survival & differentiation of 1000s of cells to be monitored

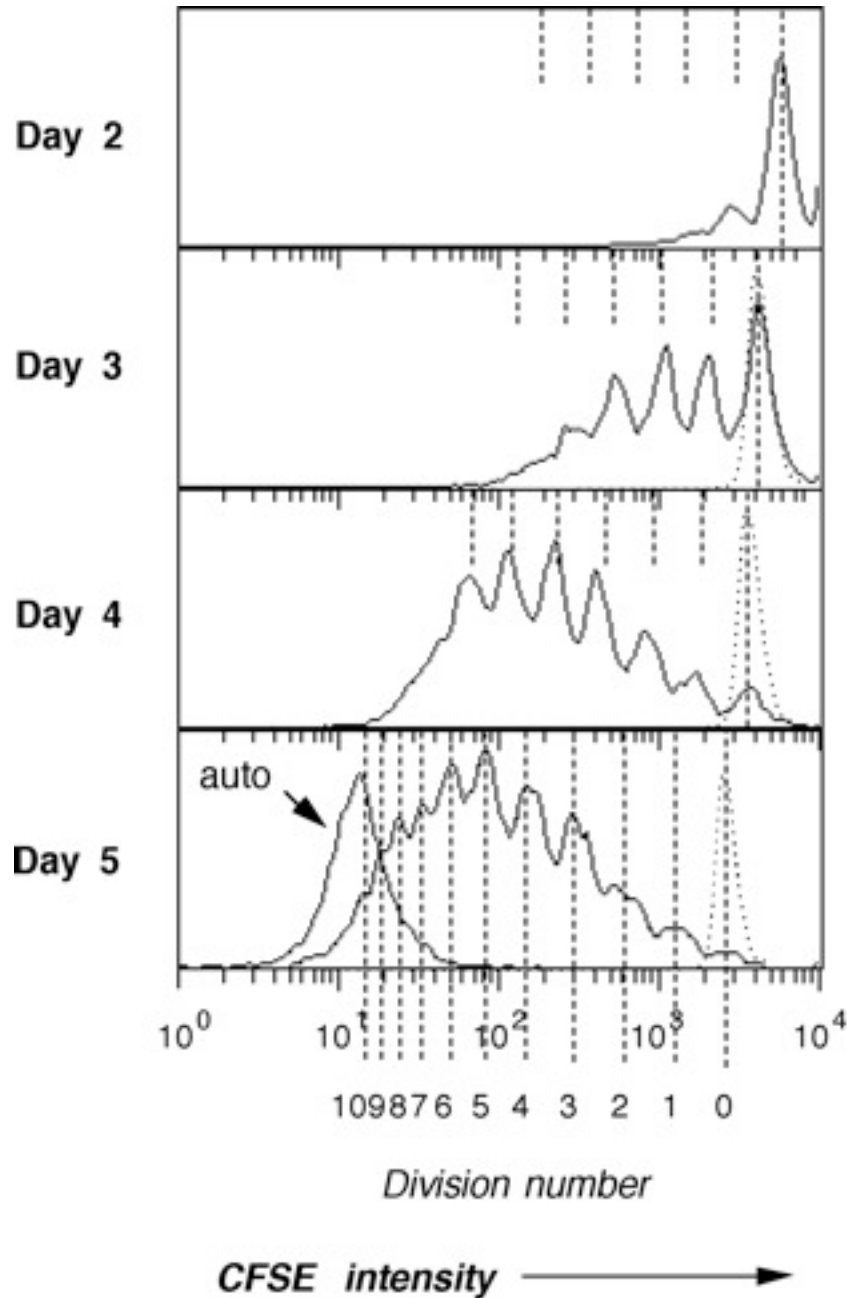
B cells stimulated in vitro

Purified mouse spleen resting B cells + CD40Ligand and Interleukin 4

- Division is asynchronous

- The peaks are limited to the autofluorescence by formula

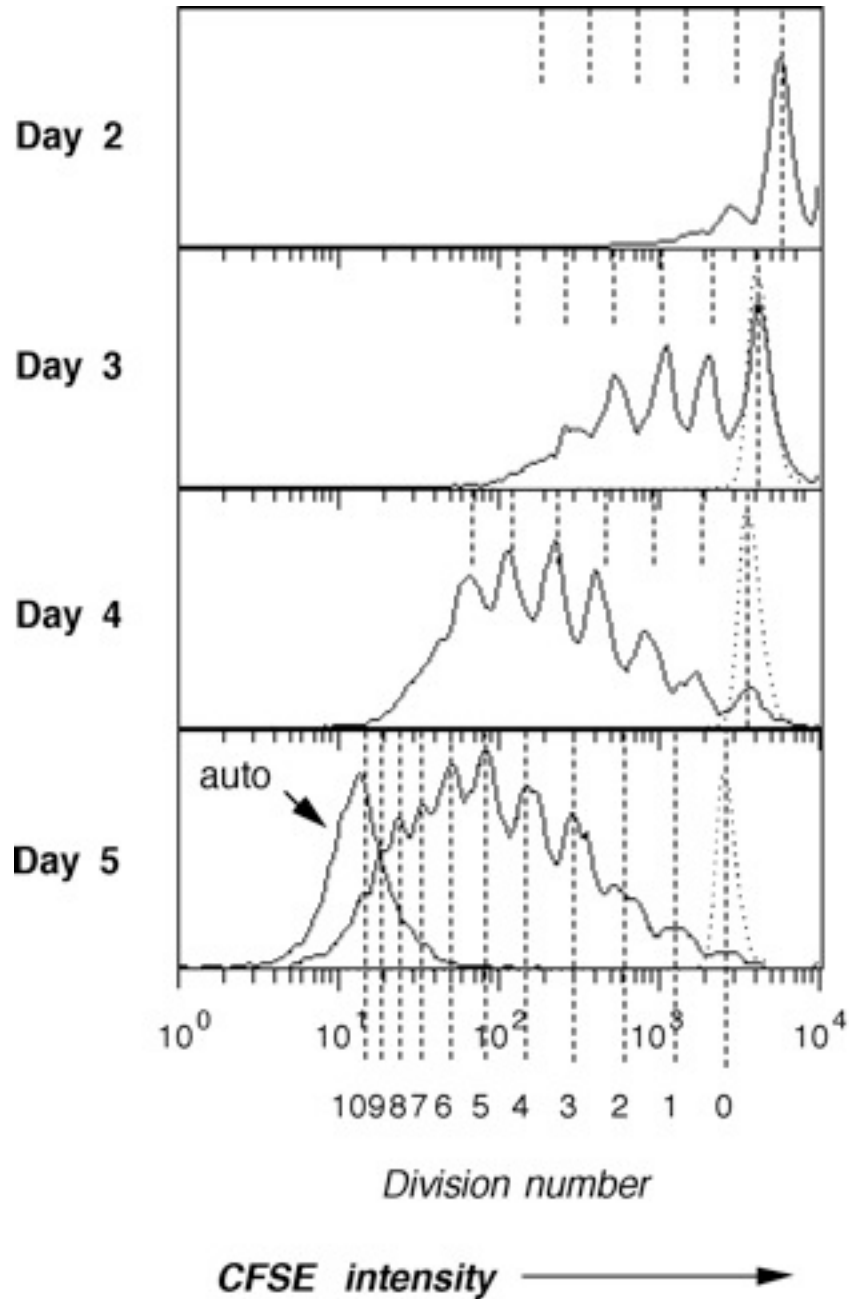
$$((\text{Start FL} - \text{AF}) / 2^{\text{div number}}) + \text{AF}$$



B cells stimulated in vitro

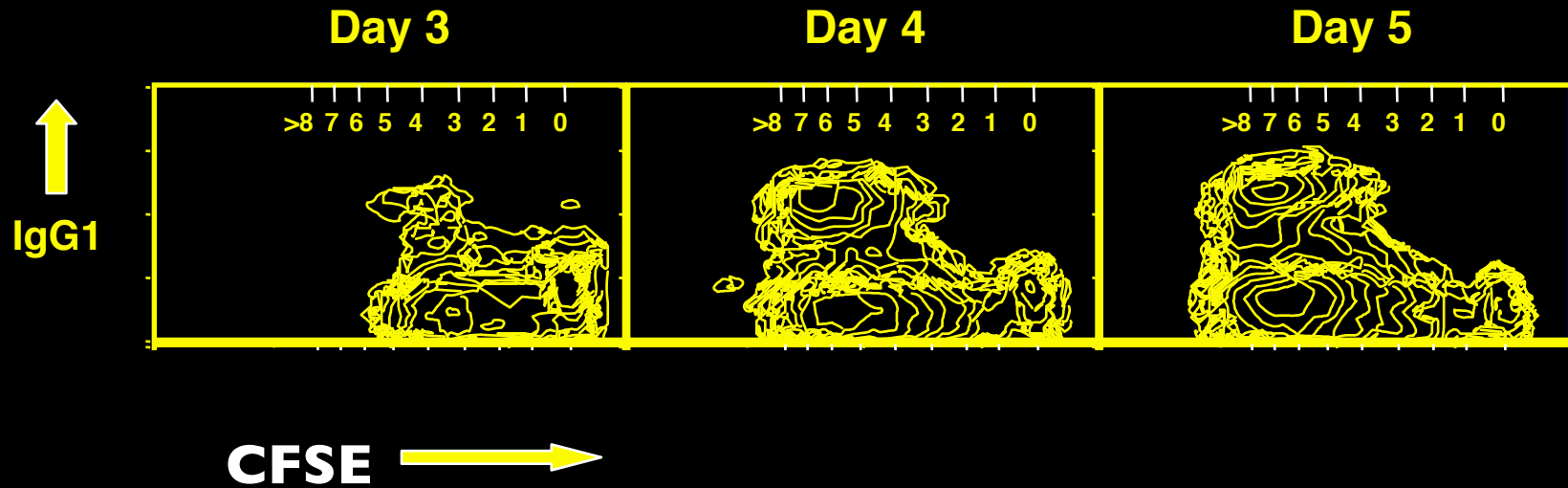
All B cells start as IgM+ and switch to IgG1 due to IL-4

When do they switch?

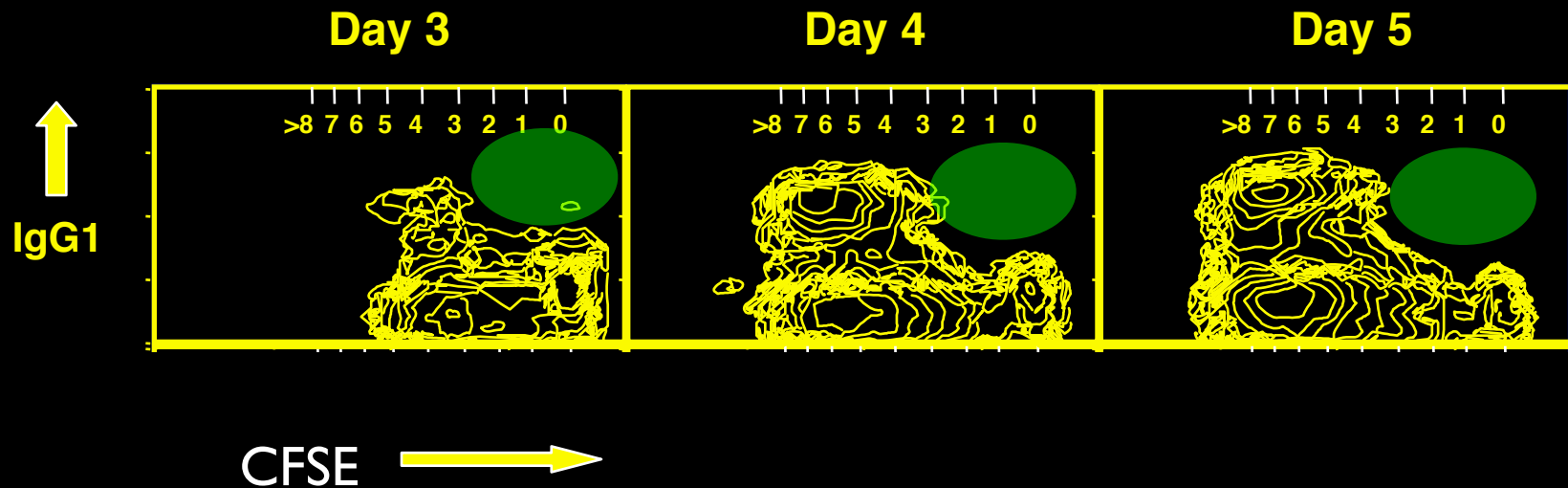


(Work of J Hasbold)

Switch from IgM to IgG1 comes after 3 divisions

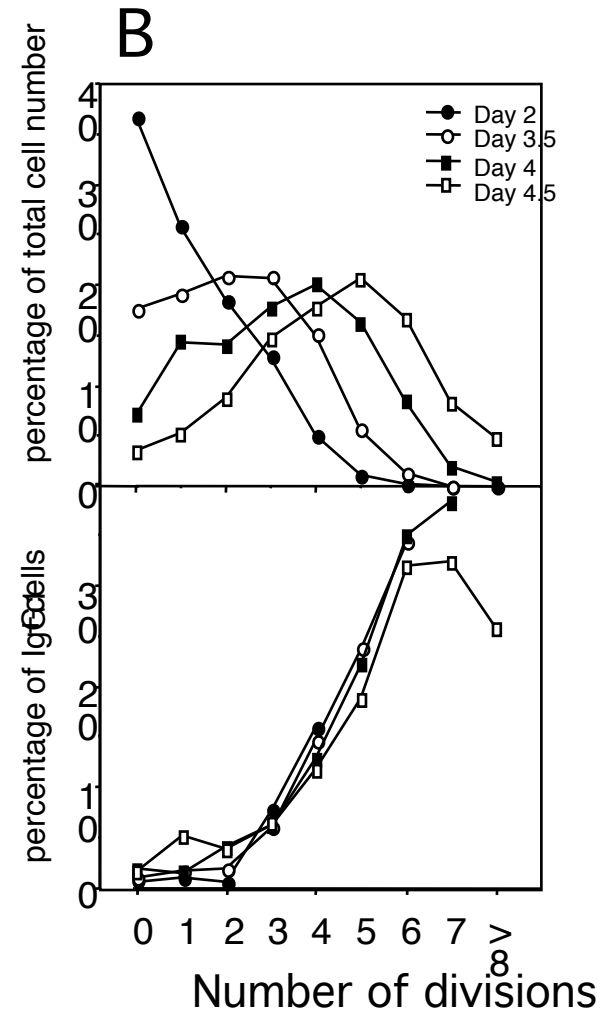
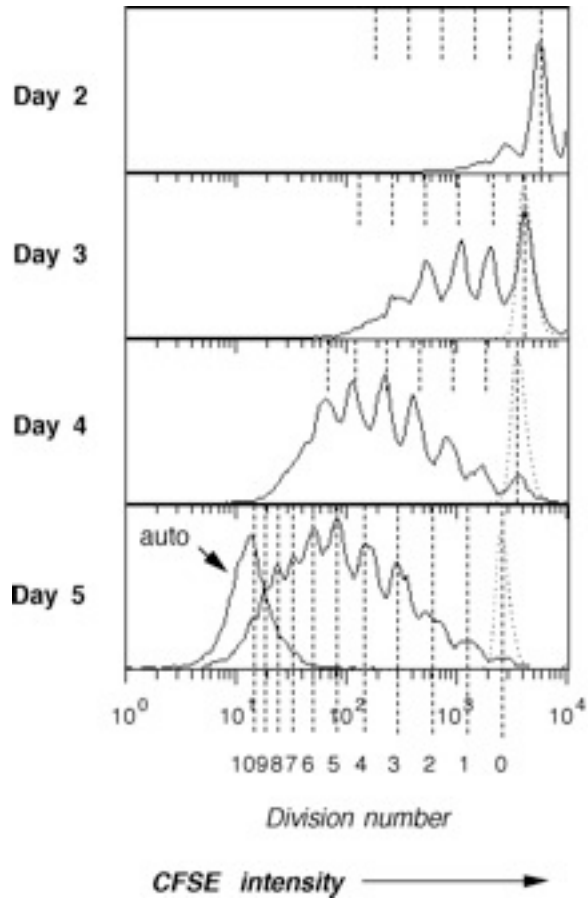


Switch from IgM to IgG1 comes after 3 divisions



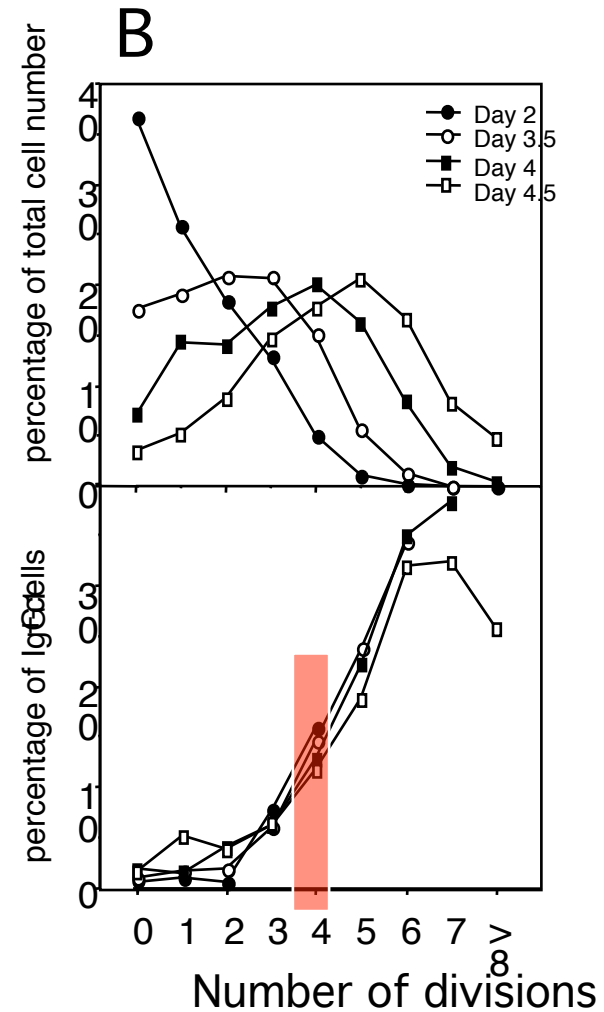
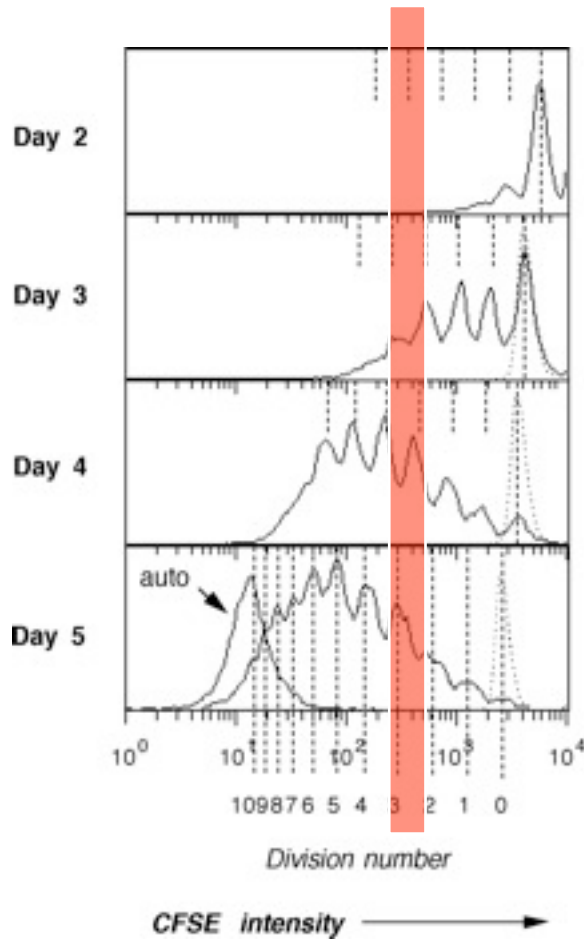
No IgG1 cells appears before 3 divisions

Isotype switching is 'Division-linked'



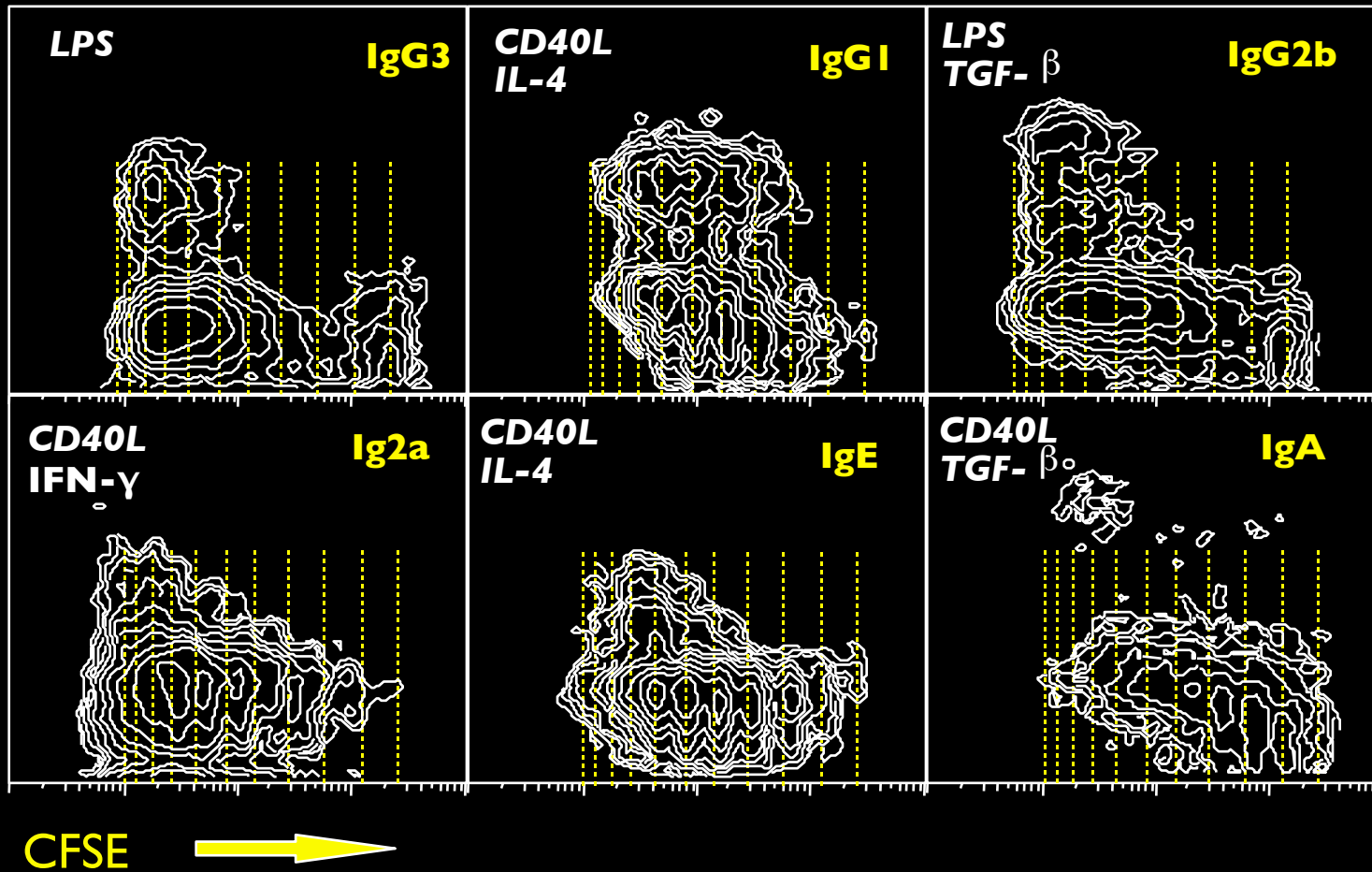
Not time-linked

Isotype switching is 'Division-linked'

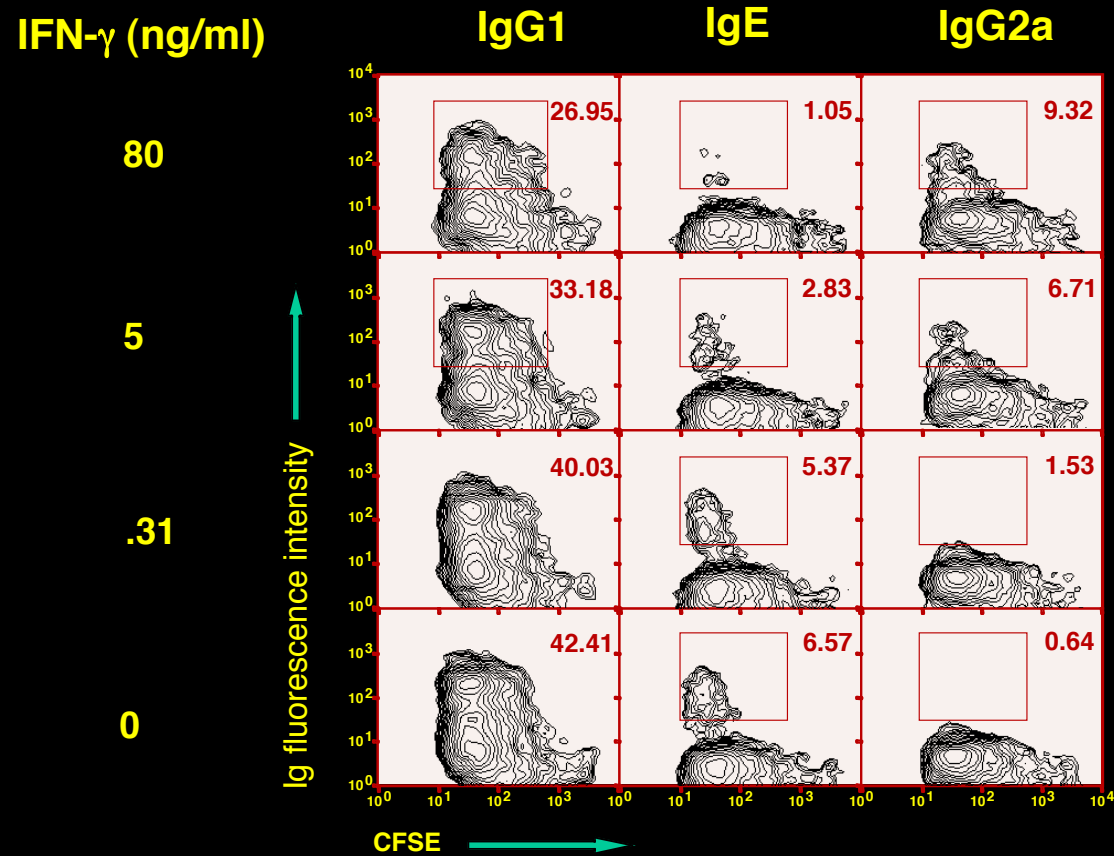


Not time-linked

All Isotypes show Division-linked switching



IFN- γ induces IgG2a expression, but down-regulates IgE



B cell rules:

Progression through division changes probability of switching

Cytokines/signals change relation with division

Rules can be found for combinations of signals - indicating cross talk or independence

2. Division linked-differentiation separate from regulation of -

Proliferation and survival

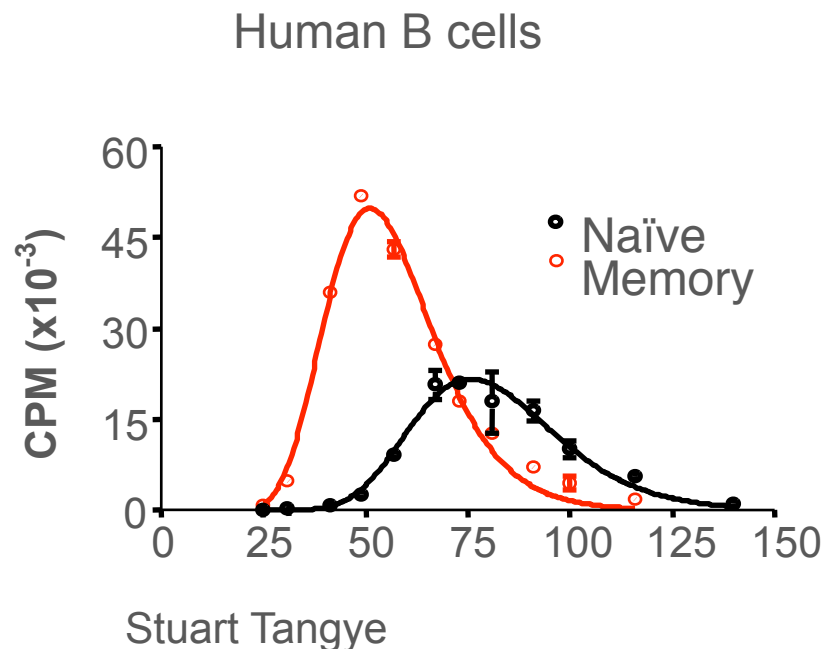
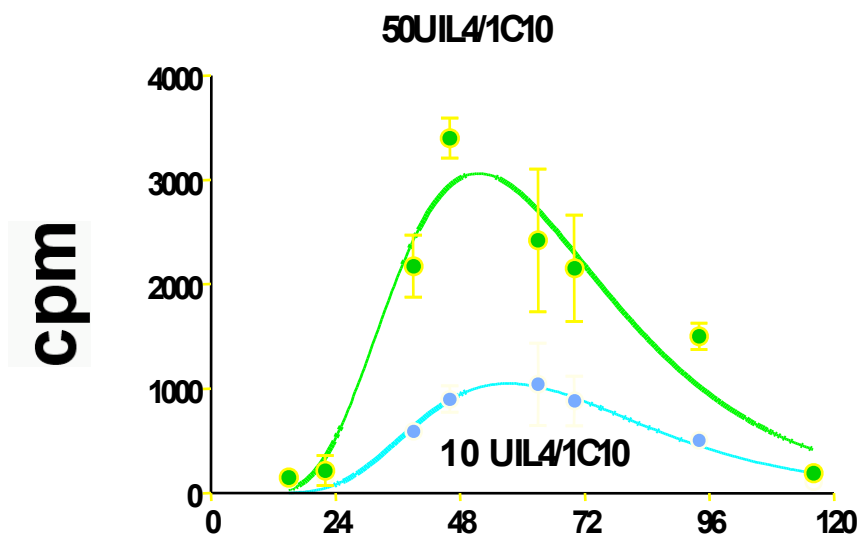
Model differentiation - combine

Models of CFSE proliferation patterns: insights into regulation of growth

- *Separation of differentiation/proliferation*
- 4-parameter model of proliferation
 - (Amanda Gett: Excel models) 2000
- 6-parameter model of proliferation and survival — (Elissa Deenick: Excel and Cellular Calculator) 2002
- 13-parameter model: Cyton model — (Edwin Hawkins, Carel van Gend: java and matlab) 2007
- Multi-parameter Cyton model: Branching process formulation (Vijay Subramanian, Ken Duffy) 2008

Proliferation empirical law -

Lognormal variation in time to first division



Variables - Mean time to divide, variance and area

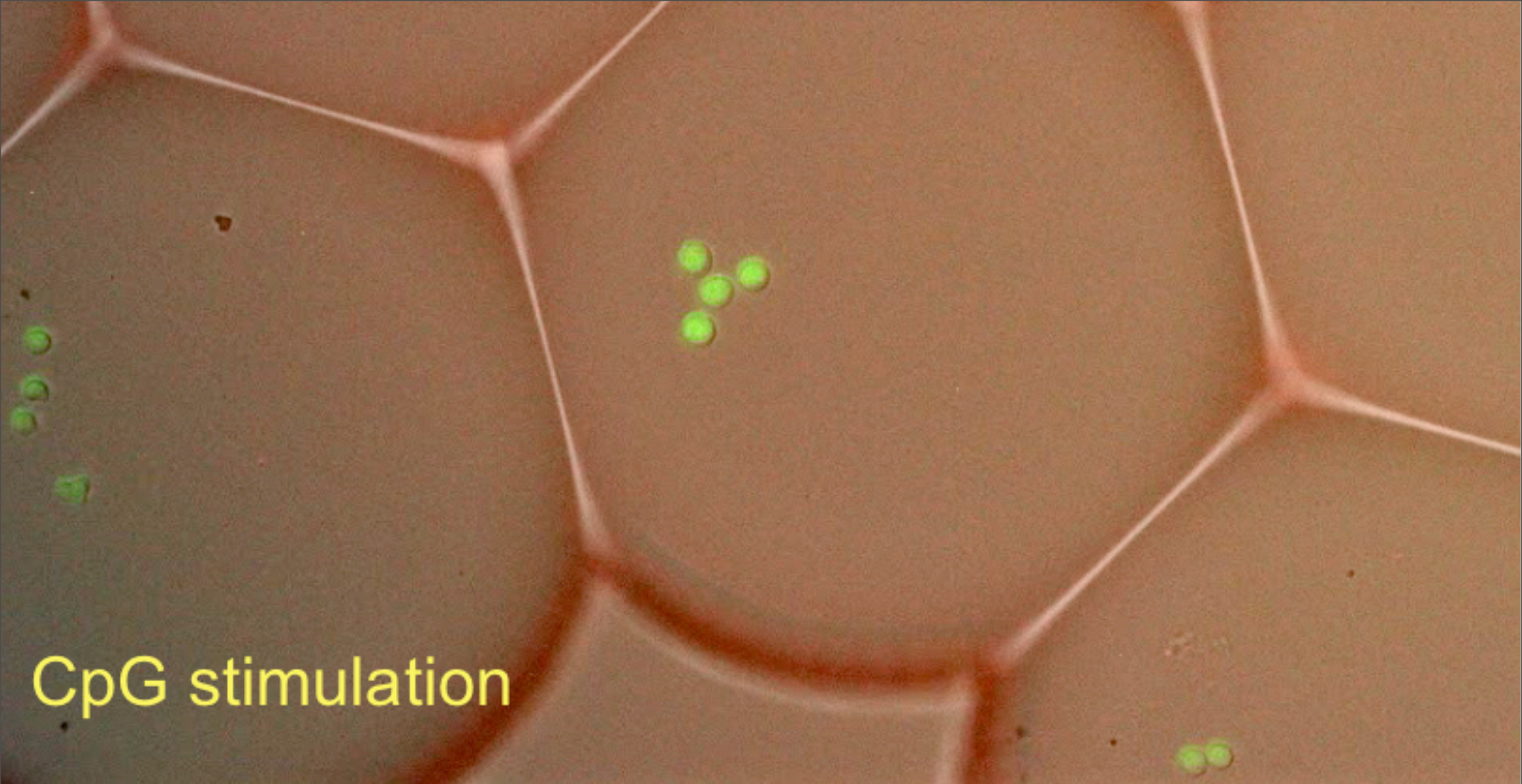
This is major source of division heterogeneity

All examples - mouse T and B - human T and B - all stimuli - one model fits all!

Subsequent divisions and inheritance of times?

Hypothesis

**Filming*



CpG stimulation

A single-cell pedigree analysis of alternative stochastic lymphocyte fates

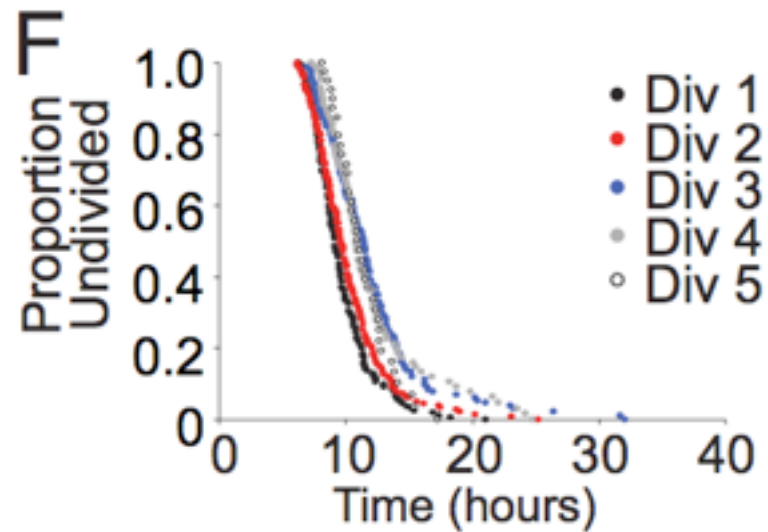
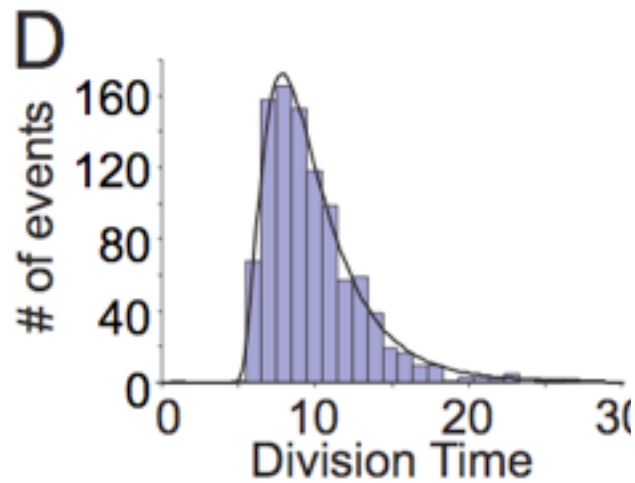
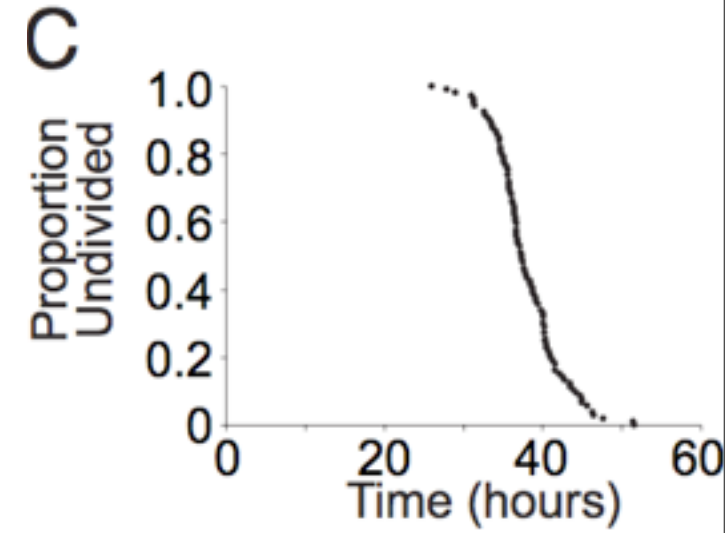
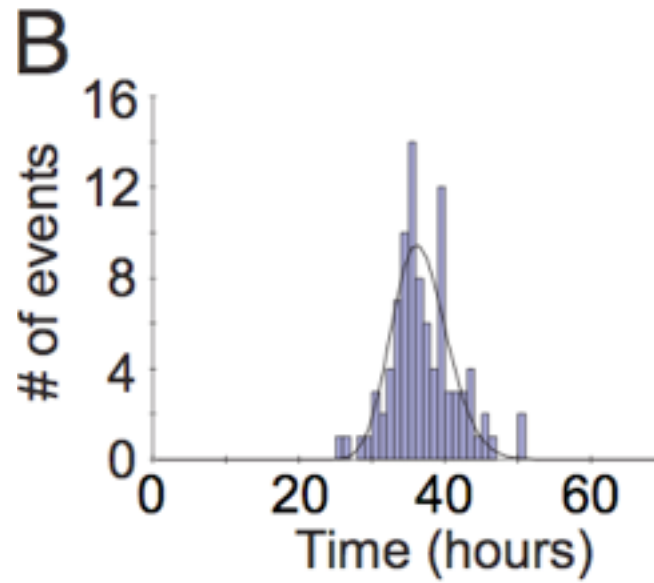
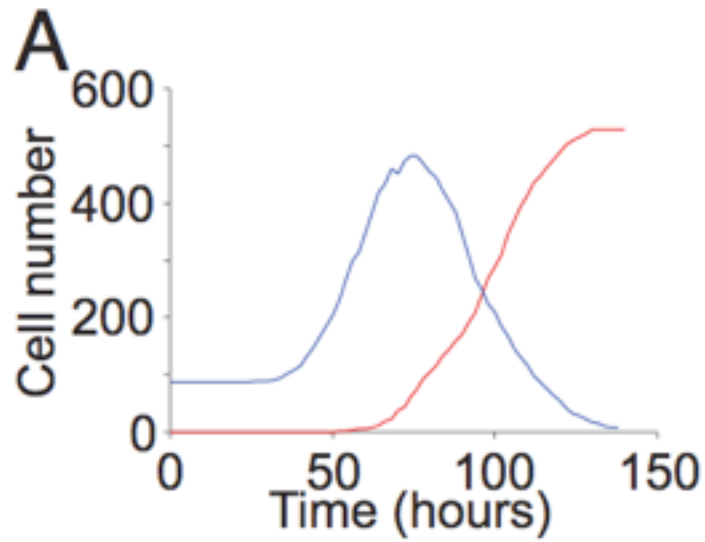
E. D. Hawkins^a, J. F. Markham^{a,b}, L. P. McGuinness^a, and P. D. Hodgkin^{a,1}

^aImmunology Division, The Walter and Eliza Hall Institute of Medical Research, 1G Royal Parade, Victoria 3050, Australia; and ¹National Information and Communications Technology Australia, and Department of Electrical Engineering, University of Melbourne, Lvl2/Building 193, Victoria 3010, Australia

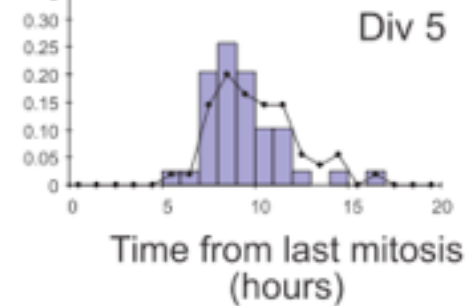
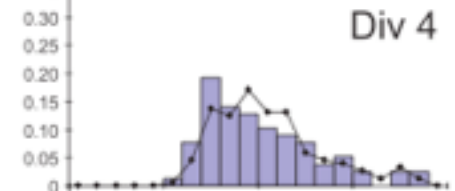
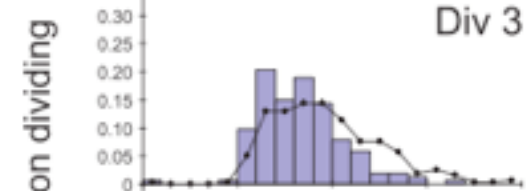
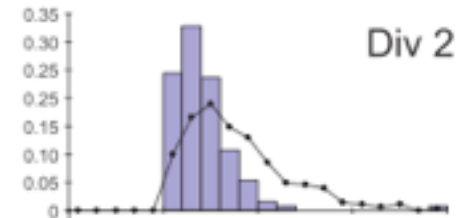
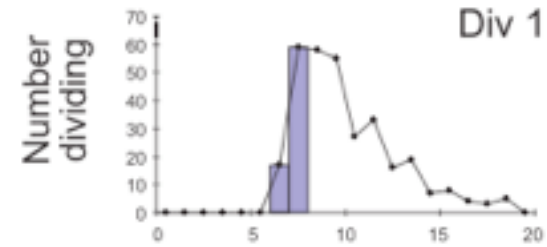
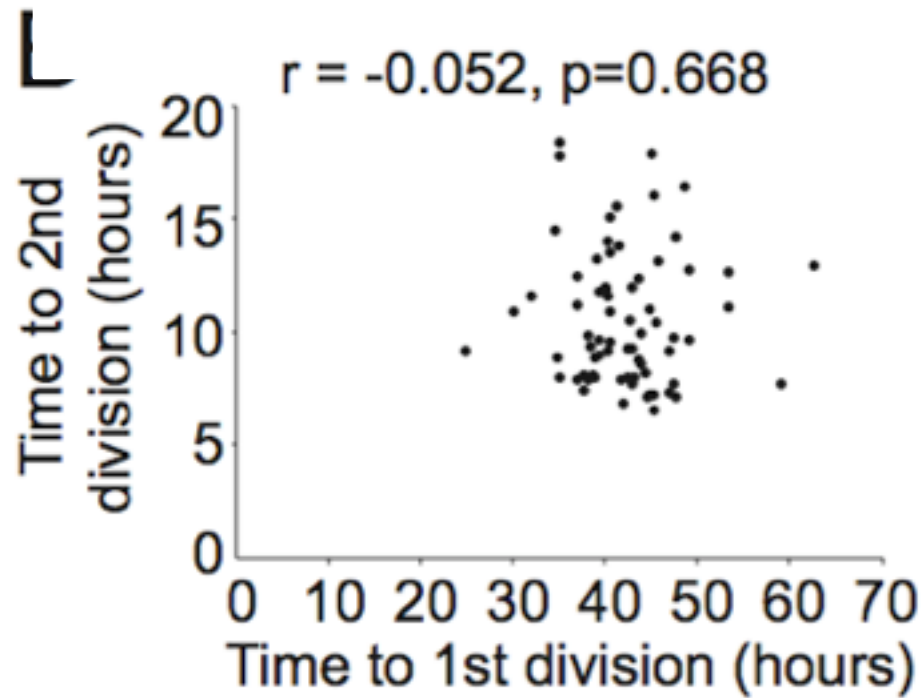
Communicated by Gustav J. Nossal, University of Melbourne, Victoria, Australia, June 8, 2009 (received for review February 2, 2009)

In contrast to most stimulated lymphocytes, B cells exposed to Toll-like receptor 9 ligands are nonself-adherent, allowing individual lymphocyte responses, and to date there is no quantitative information on how lymphocyte survival is regulated and altered

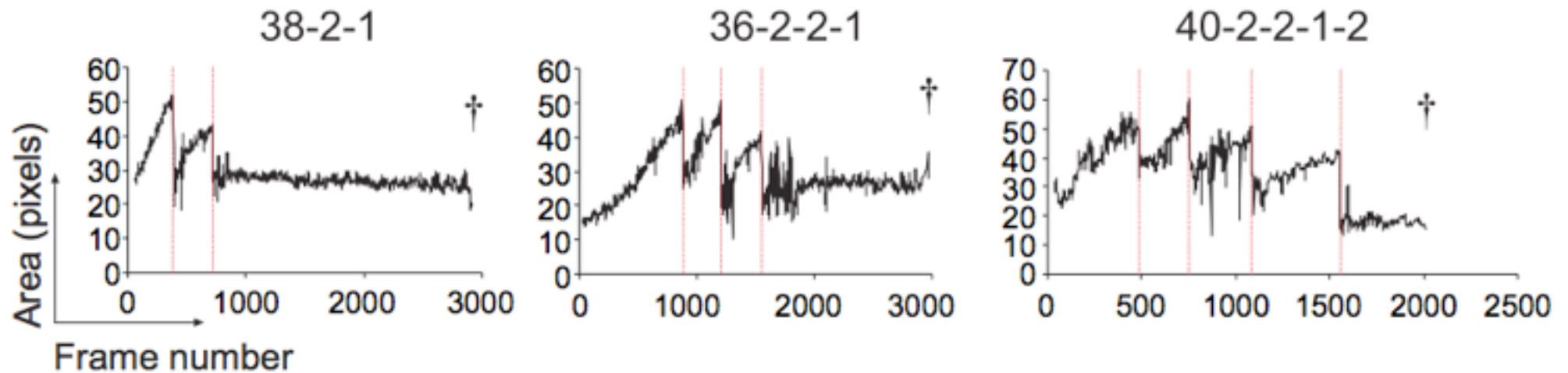
Proliferation, cessation and death



Fast does not beget fast...

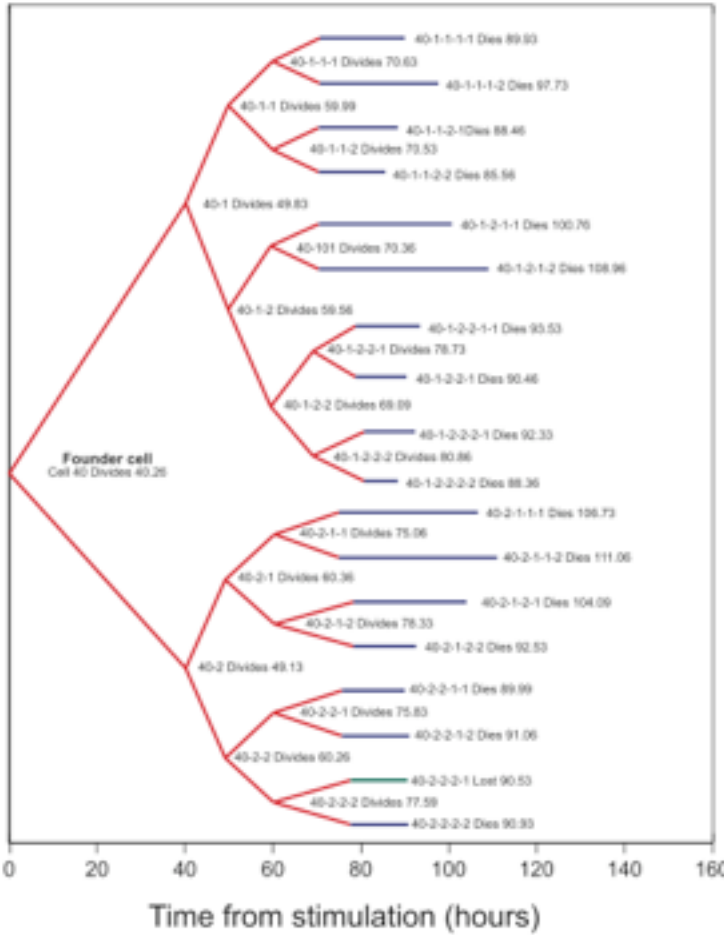
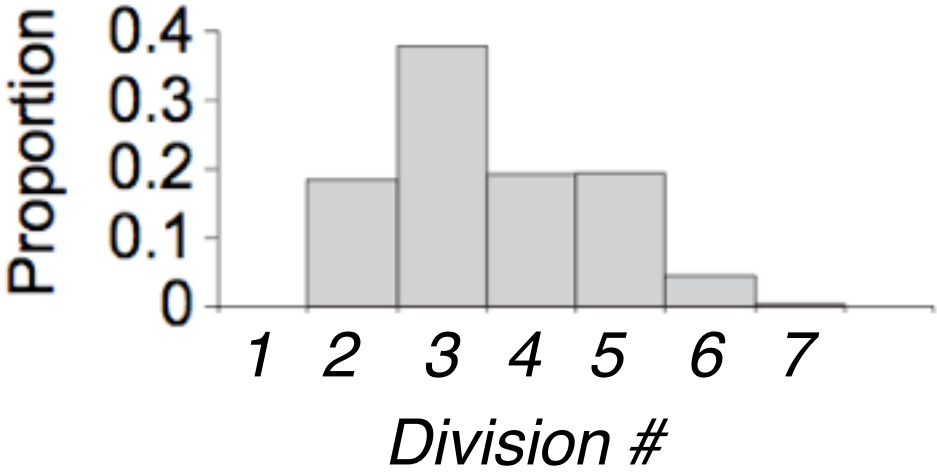


Growth, division, cessation and death



Founder effects on division 'destiny'

4 symmetric divisions



Key experimental observations -

1. Lognormal times to divide
2. Resetting of times after division - lack of inheritance
- 3- Division 'counting' can alter parameters
eg. division progression
- 4 - Independent regulation of division and death

A model of immune regulation as a consequence of randomized lymphocyte division and death times

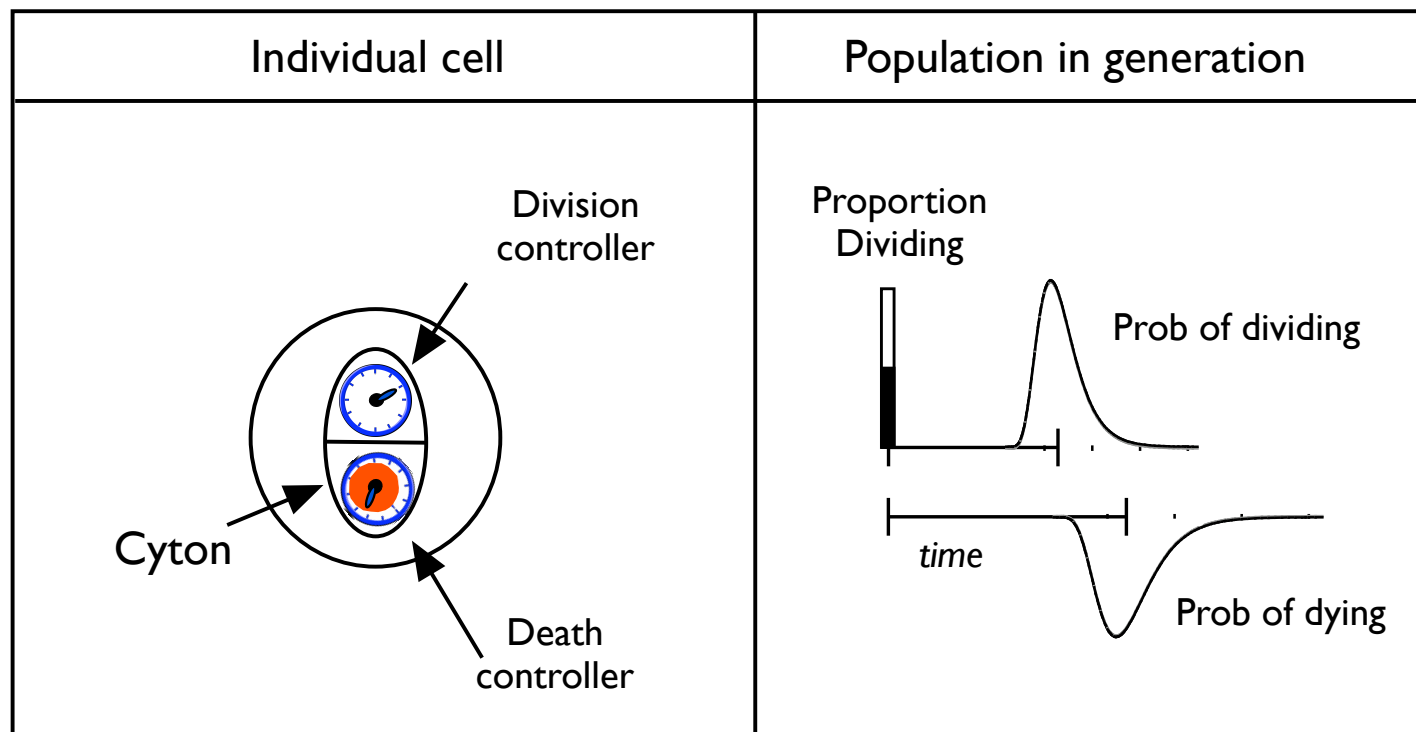
E. D. Hawkins^{*†}, M. L. Turner^{*†}, M. R. Dowling^{**‡}, C. van Gend^{*}, and P. D. Hodgkin^{*5}

^{*}Immunology Division, The Walter and Eliza Hall Institute of Medical Research, 1G Royal Parade, Parkville, Victoria 3050, Australia; [†]Department of Medical Biology, University of Melbourne, Parkville, Victoria 3010, Australia; and [‡]School of Physical Sciences, University of Queensland, Queensland 4072, Australia

Communicated by Gustav J. Nossal, University of Melbourne, Victoria, Australia, January 6, 2007 (received for review September 19, 2006)

The magnitude of an adaptive immune response is controlled by the interplay of lymphocyte quiescence, proliferation, and apoptosis. How lymphocytes integrate receptor-mediated signals influencing

When first examined, cell loss seemed to follow an exponential decay function consistent with a constant probability of dying over time (Fig. 1A and refs. 7, 9, and 15). However, we noted deviations



Use to measure genetic effects

and 'Calculation of signal integration'

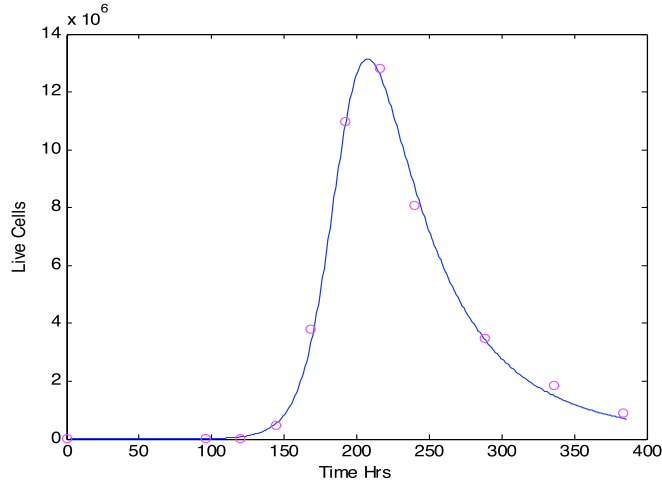
Fits T and B cells - allows sensitive regulation
relies on the randomising features for control

The model can provide a quantitative description of *in vivo* cellular immune responses

Proliferation of LCMV-specific T cells post infection

GP276: CD8

specific T cells per spleen



Starting cell number

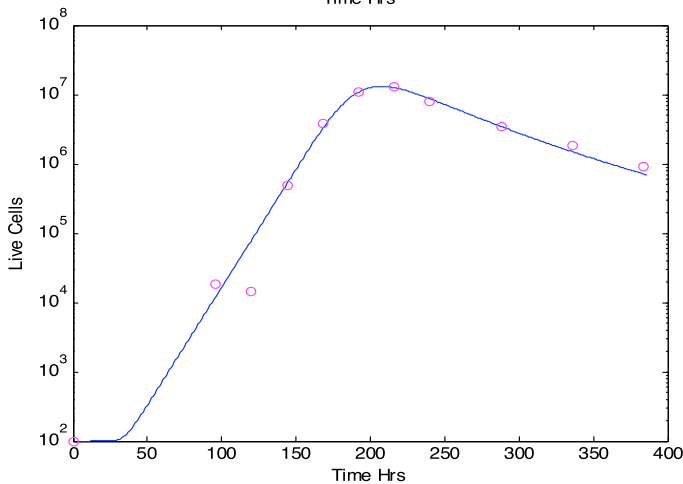
GP61: 100

GP276: 100

Subsequent division time

GP61: med = 9.9 h, s = 0.09

GP276: med = 9.4 h, s = 0.06



Division destiny

GP61: med = 19.3, s = 0.9

GP 276: med = 17.5, s = 1.1

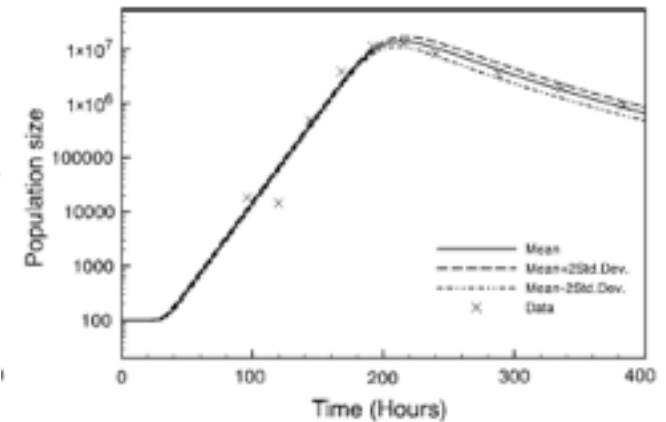
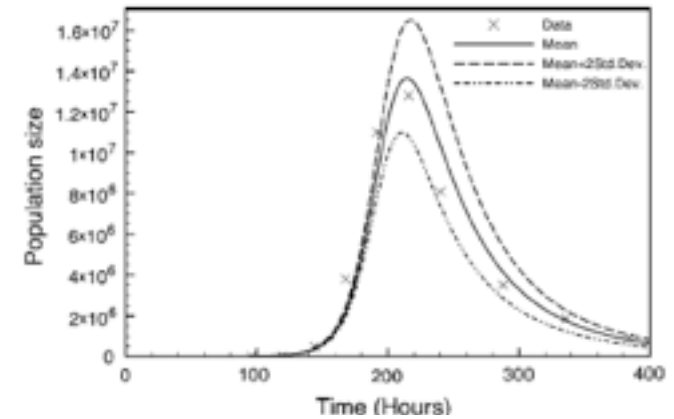
Subsequent division death rat

GP61: med = 46.1 h, s = 1.3

GP276: med = 56.7 h, s = 0.8

Branch process -

V. Subramanian, K Duffy



Data provided by Dirk Homann, (Homann et al., Nature Medicine (2003))

Cyton Model

Exploits evidence for heritable stochastic processes governing division and death times

Individual cells exhibit extreme heterogeneity

Population highly predictable

External signals alter parameters of stochastic process

Differentiation

How do division, death, differentiation all interleave?

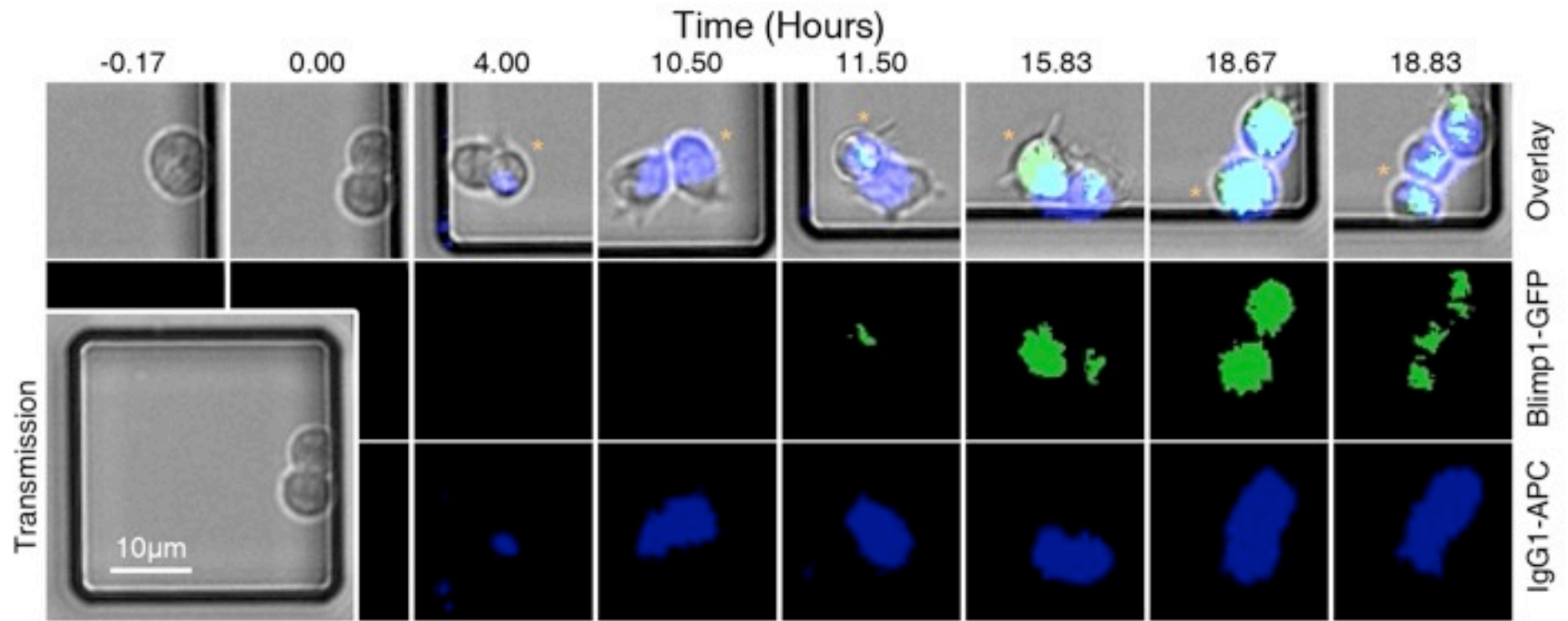
B cells - Switch antibody type, develop to ASC

Can we extend internal competition in cells to other fates?

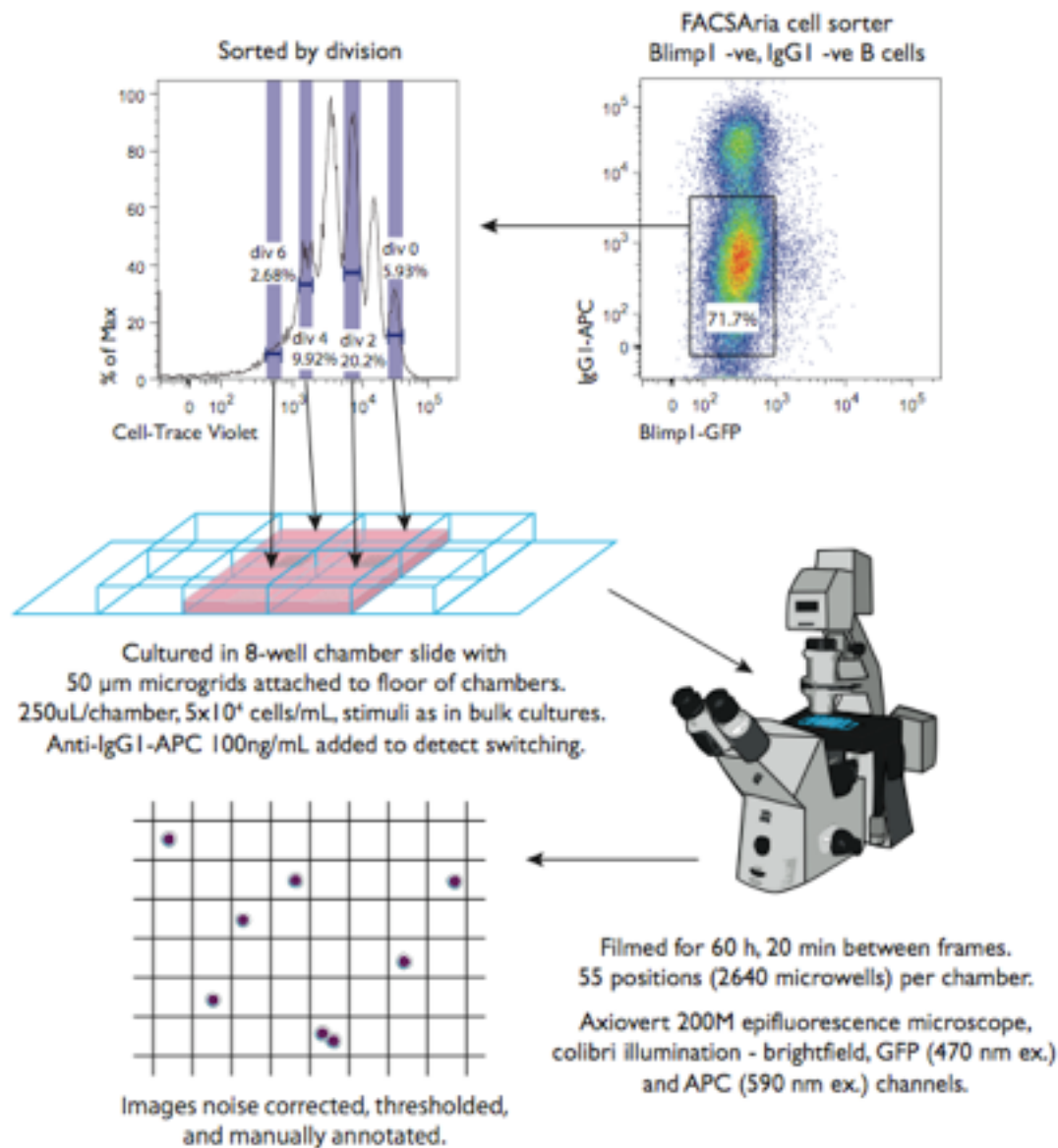
Microscopy allows the study of individual cell fates over time

(Hawkins, E.D *et al.*, *PNAS* 2009;
Duffy, K.R. *et al.*, *Science* 2012)

- Blimp-1-GFP reporter
- IgG1-APC stain



Generation filming



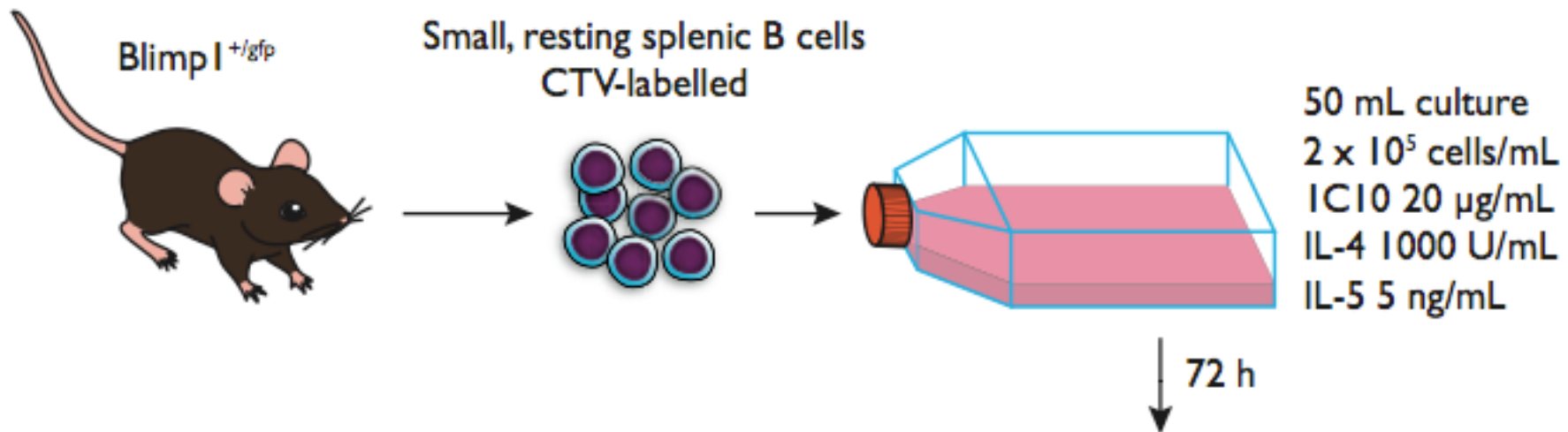
Experiment - Observe 4 fates operating simultaneously

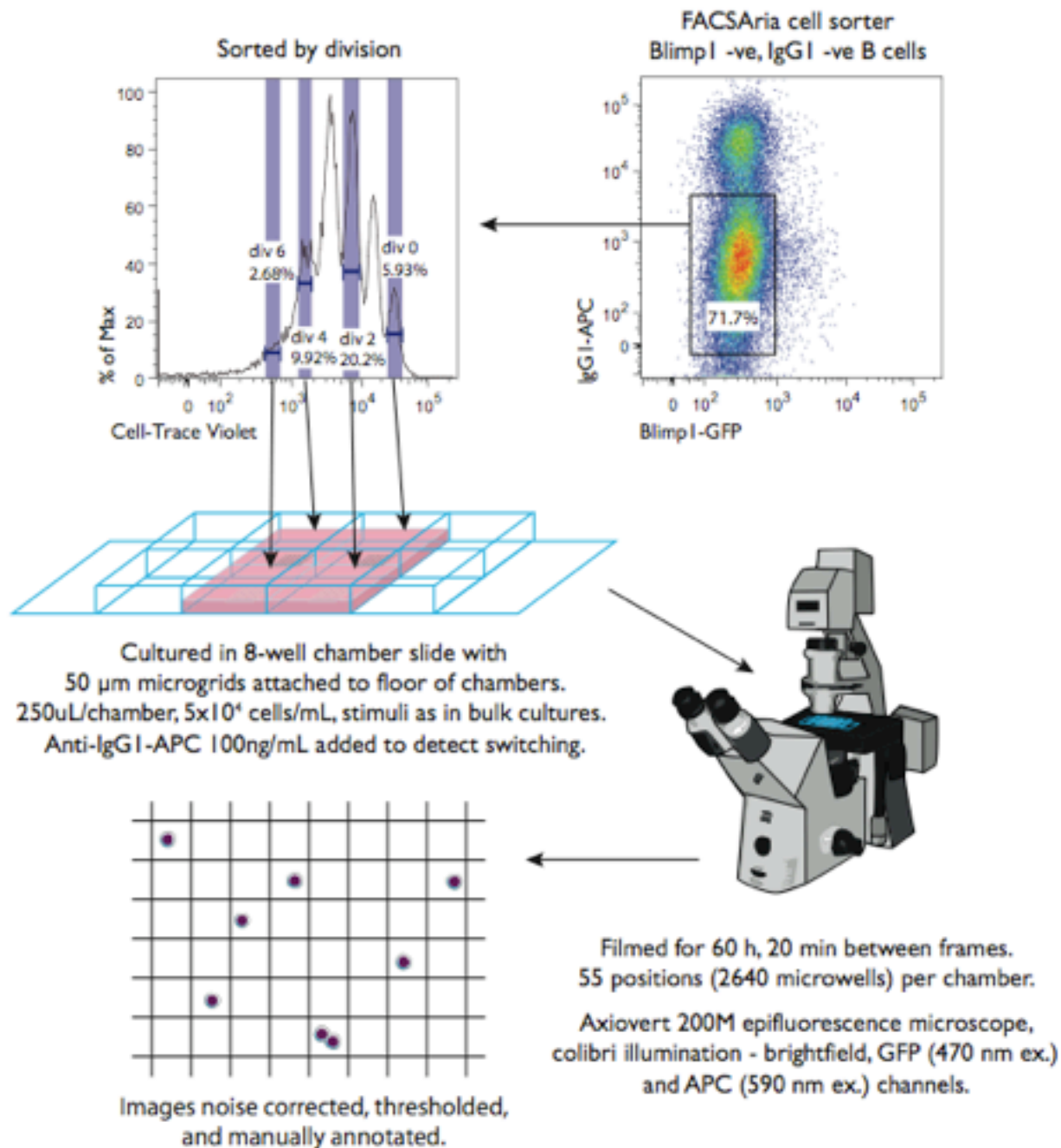
Test for competition

Mark Dowling, John Markham, Hasbold,
Ross Holmberg, Jie Zhou, Cam Wellard, Ken Duffy

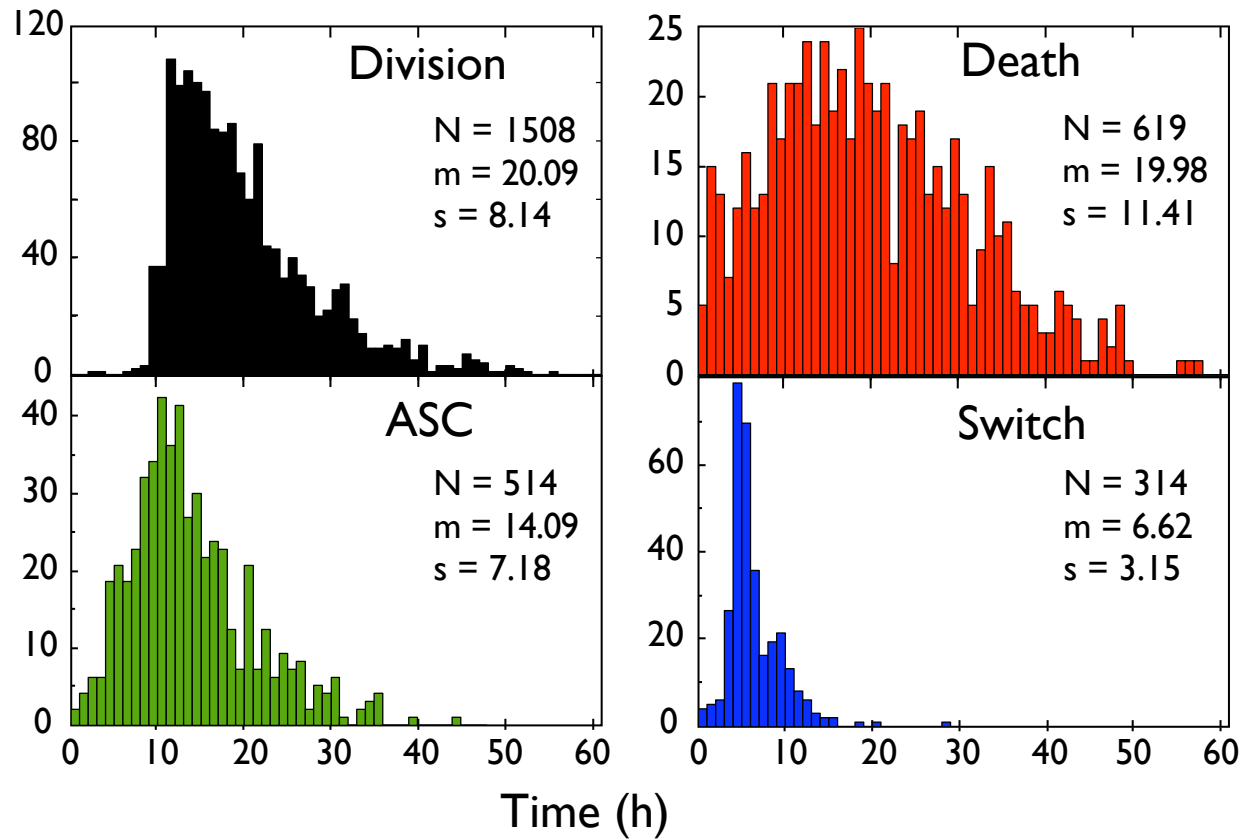
Conditions that induce division, death, switch to IgG1 and development of ASC

Examine for statistical hallmarks of competition*





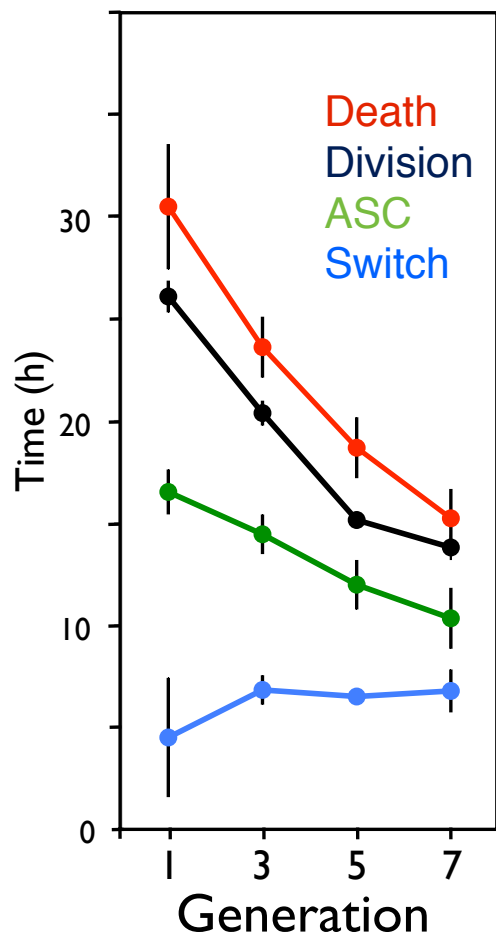
All Divisions - summary



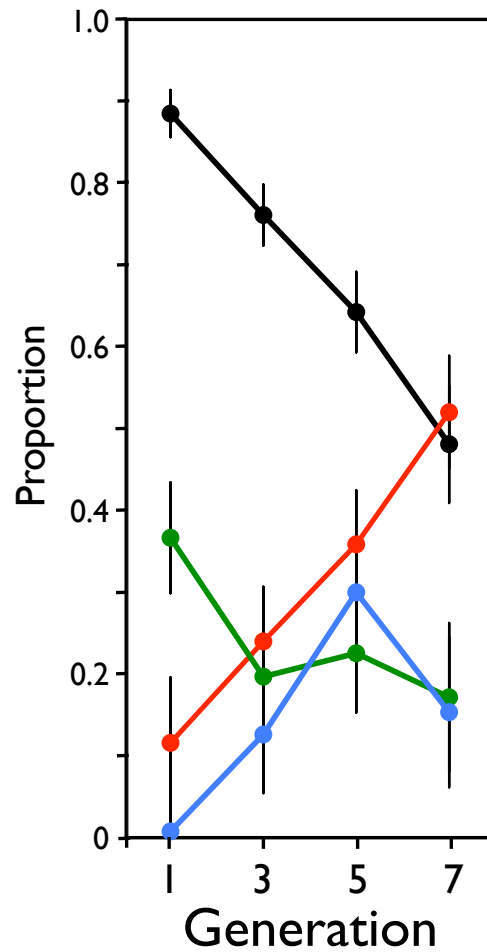
Times - long tailed distributions

Different means/variances/frequencies

Average time

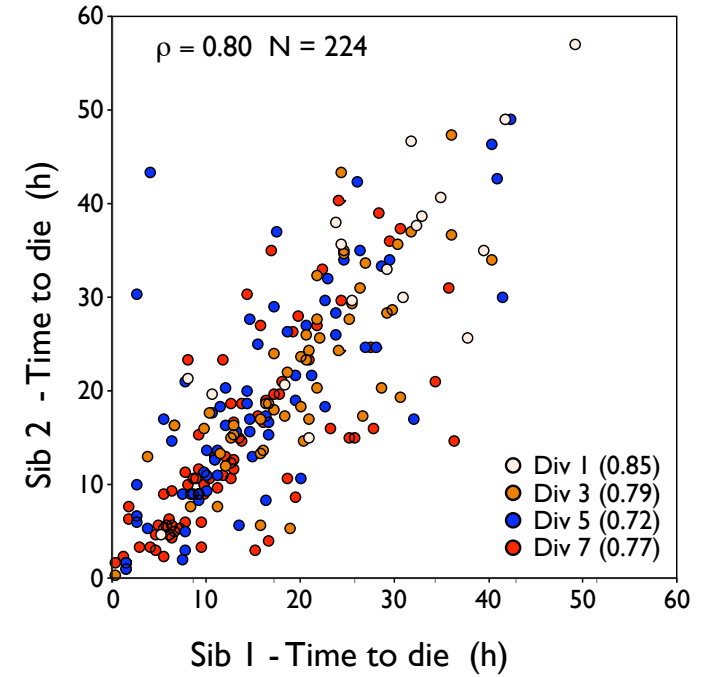
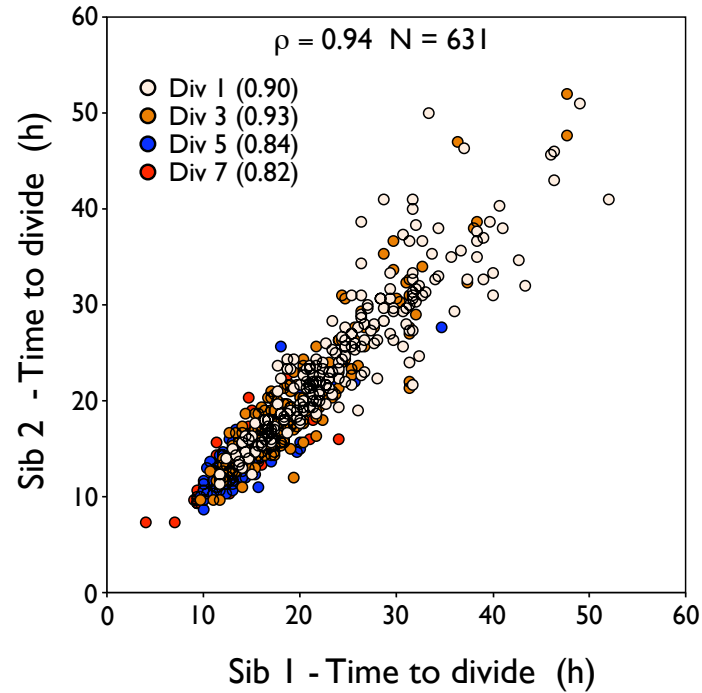
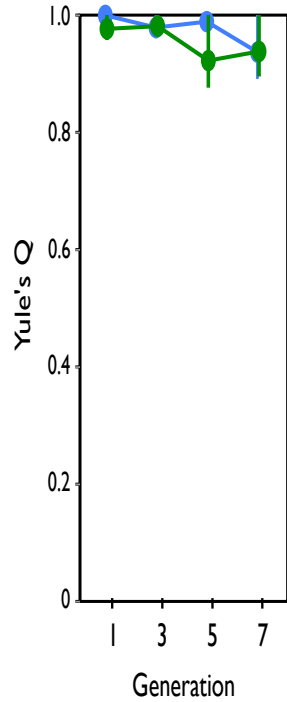
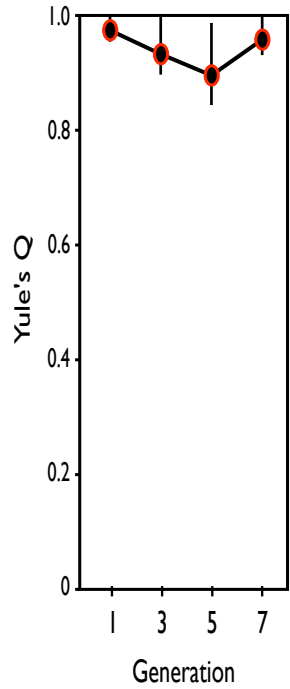


Frequency



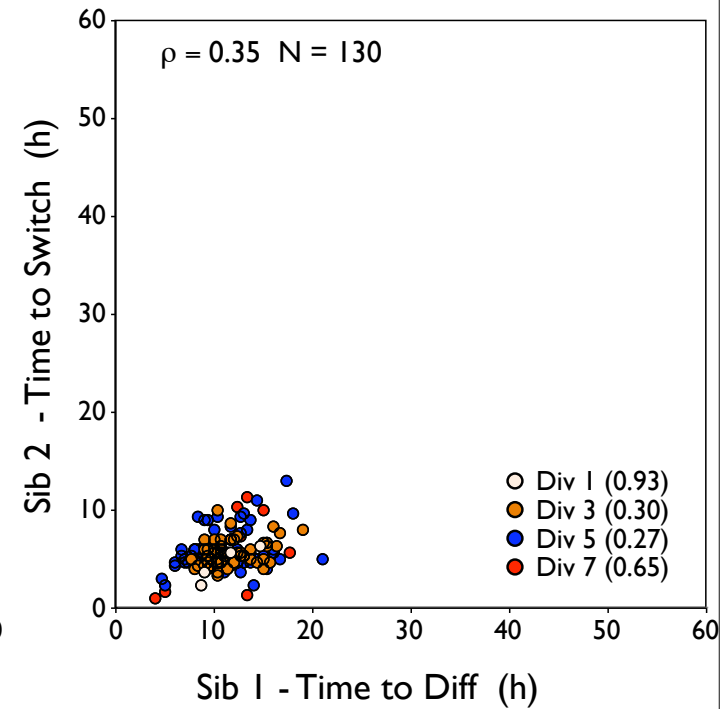
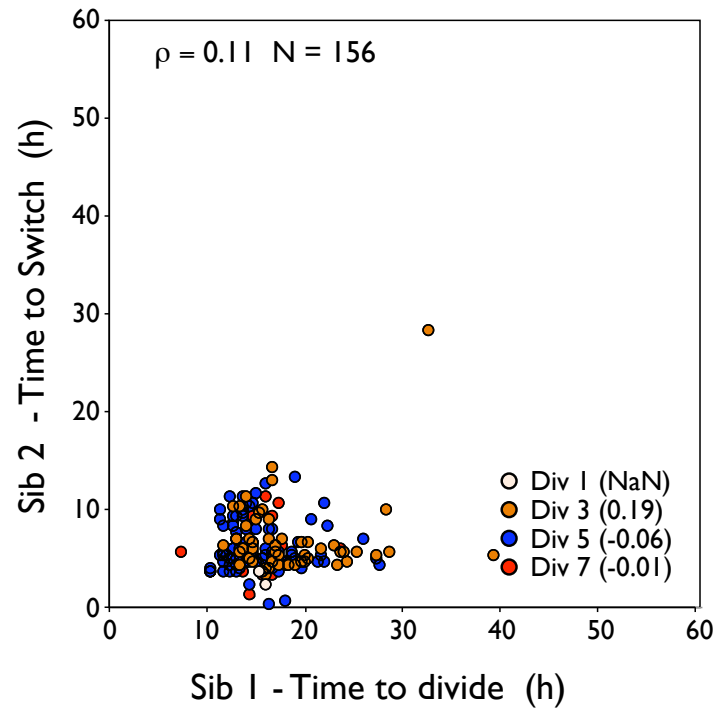
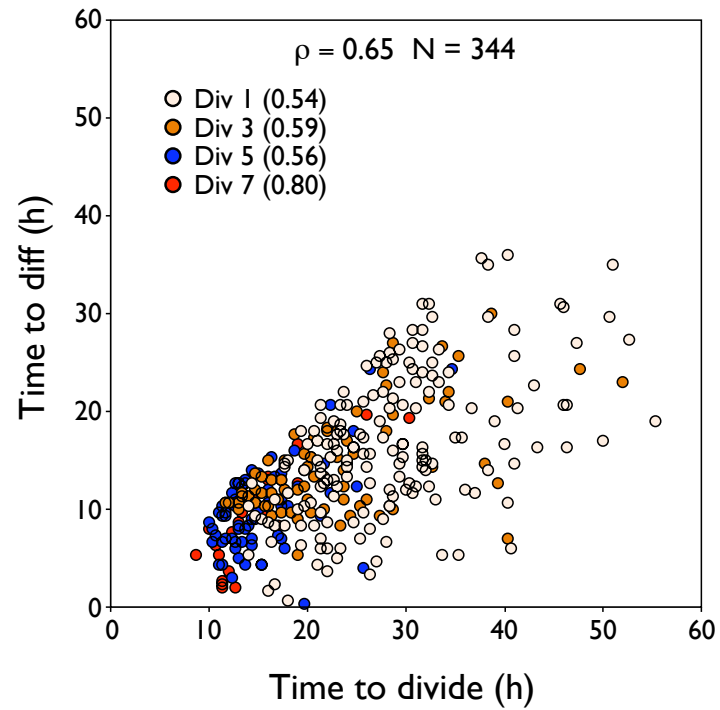
Sibling correlations

Intercellular concordant fates

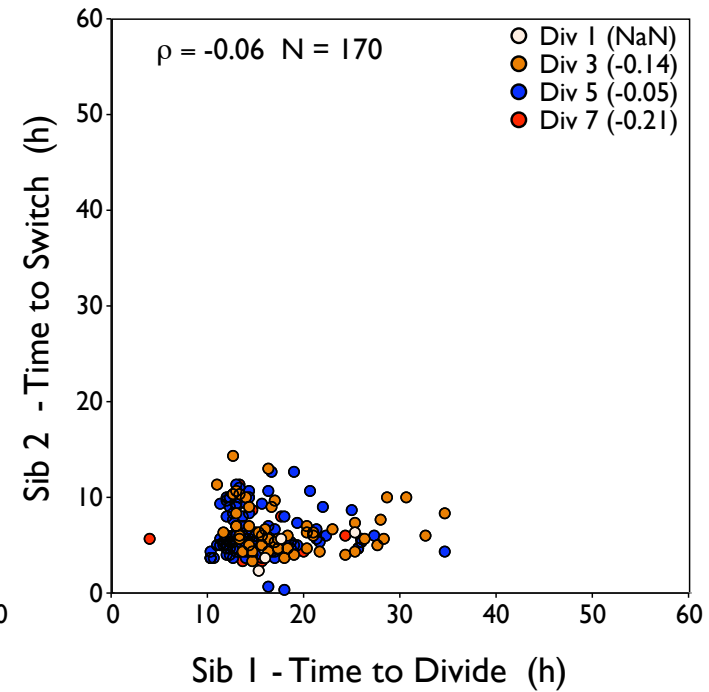
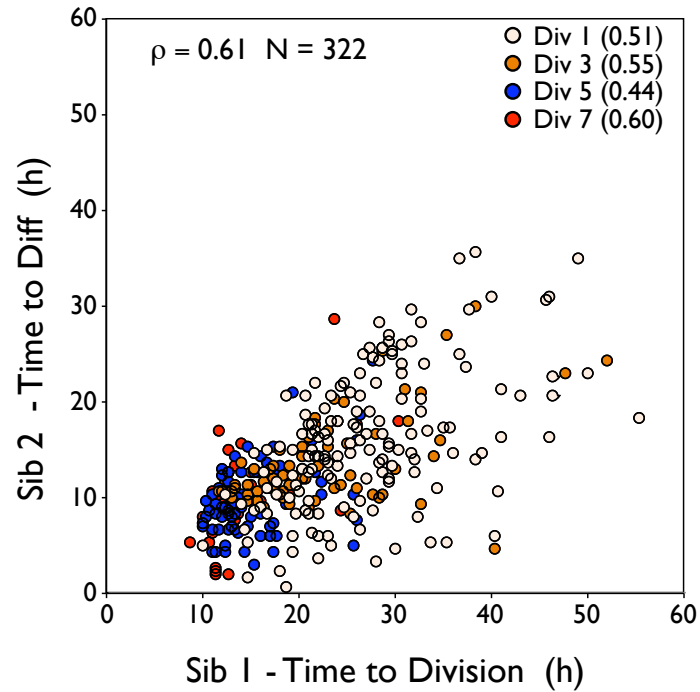
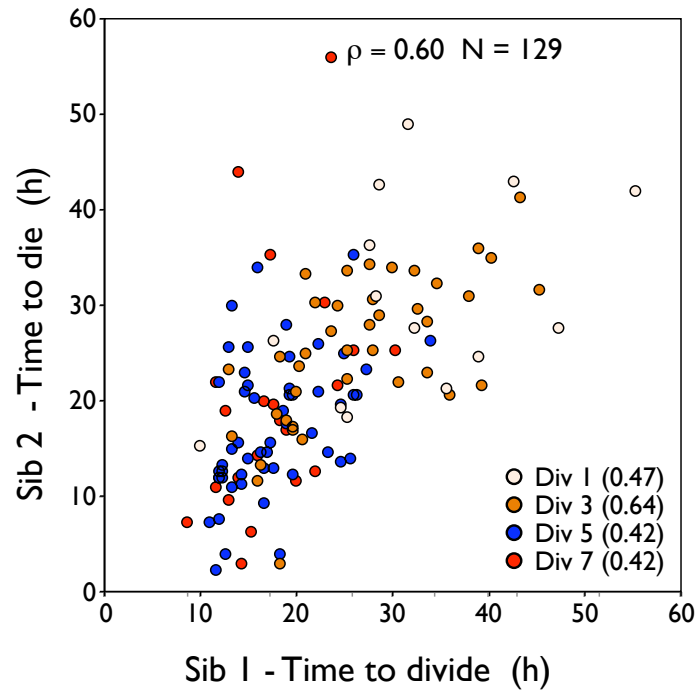


Intracellular correlations

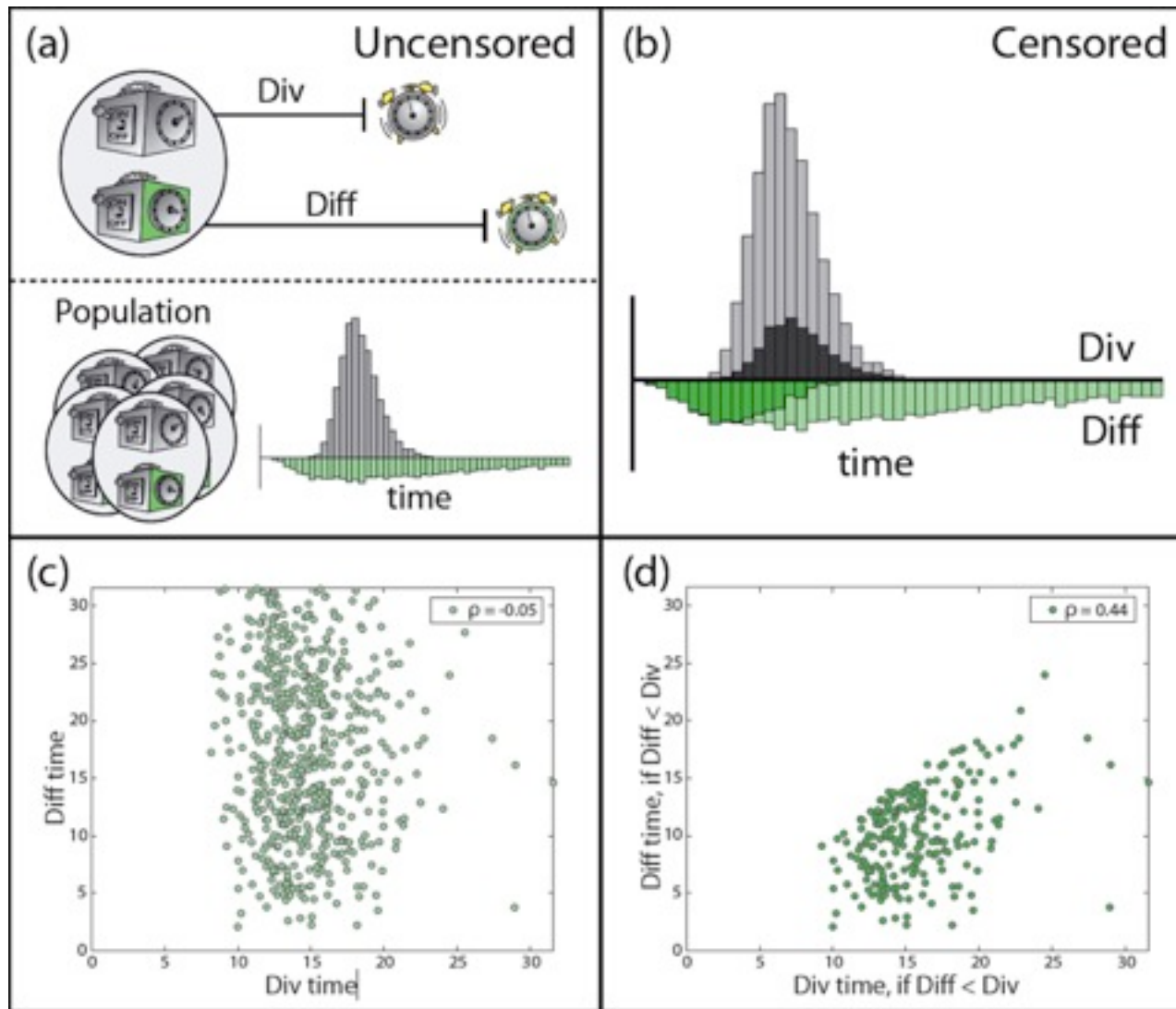
Intracellular fates



Intercellular correlations non-concordants

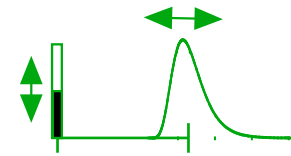
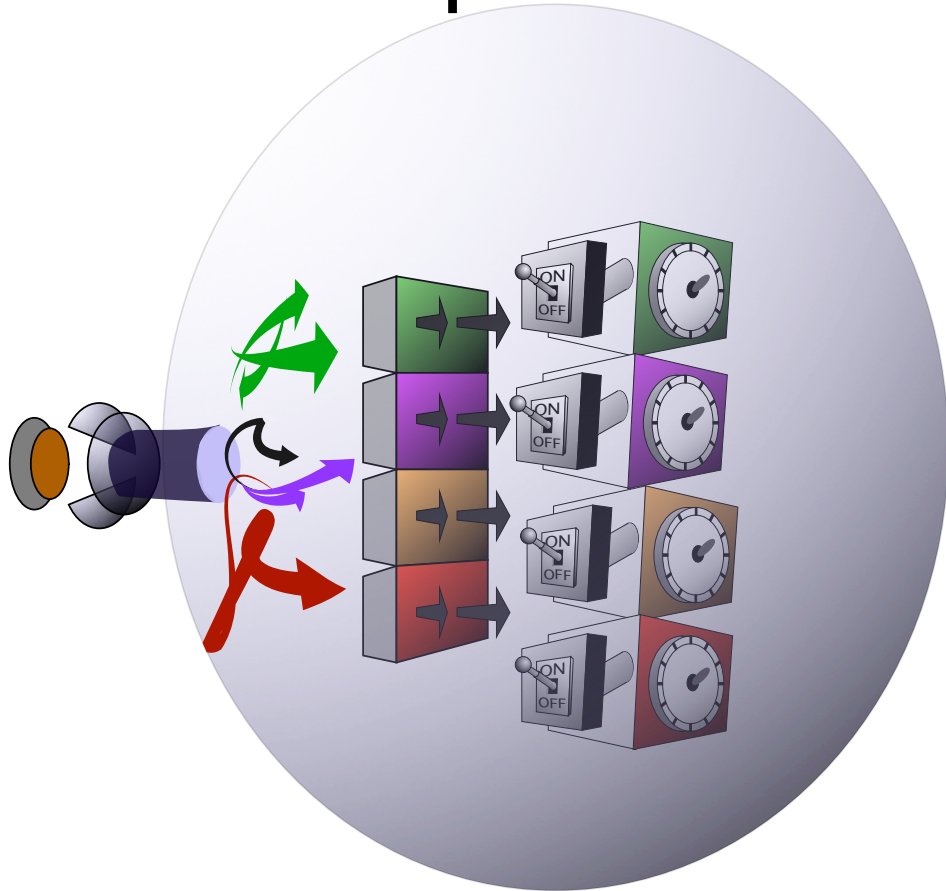


Autonomous Intracellular Competition: Hallmarks of censorship

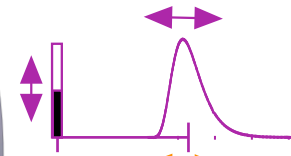


Duffy and Hodgkin, 2012 Trends in Cell Biology

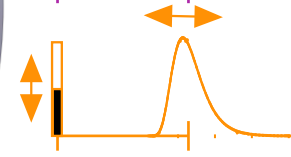
Four way Cyton - Autonomous Intracellular Competition - Rules of censorship



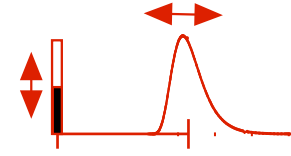
Division (restart)



Death (always wins)

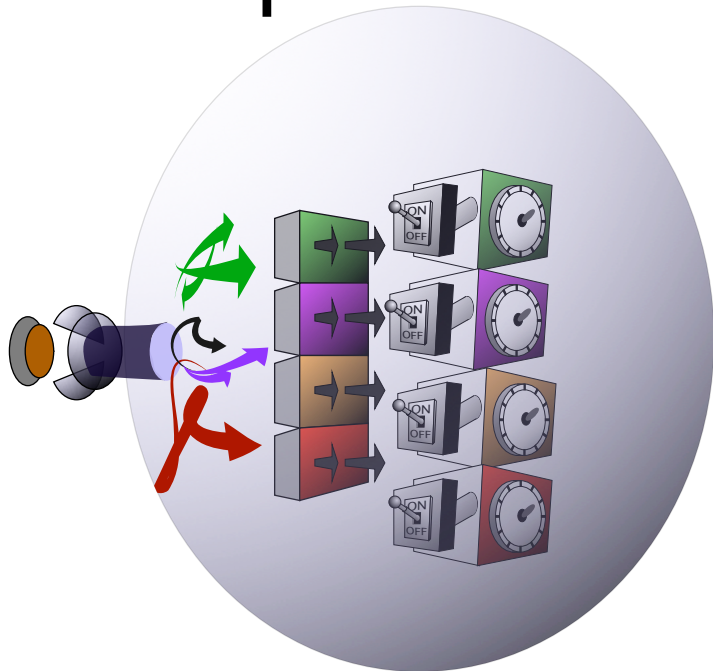


Switch (no affect)

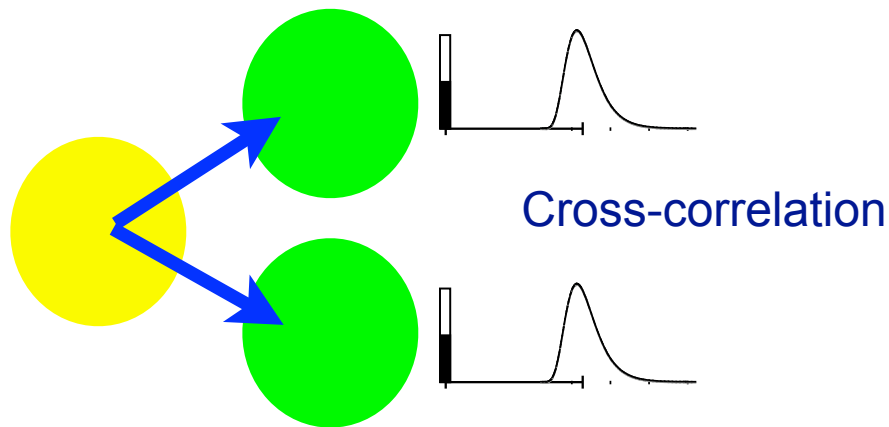


ASC (stops switch)

Autonomous Intracellular Competition - Model parameters

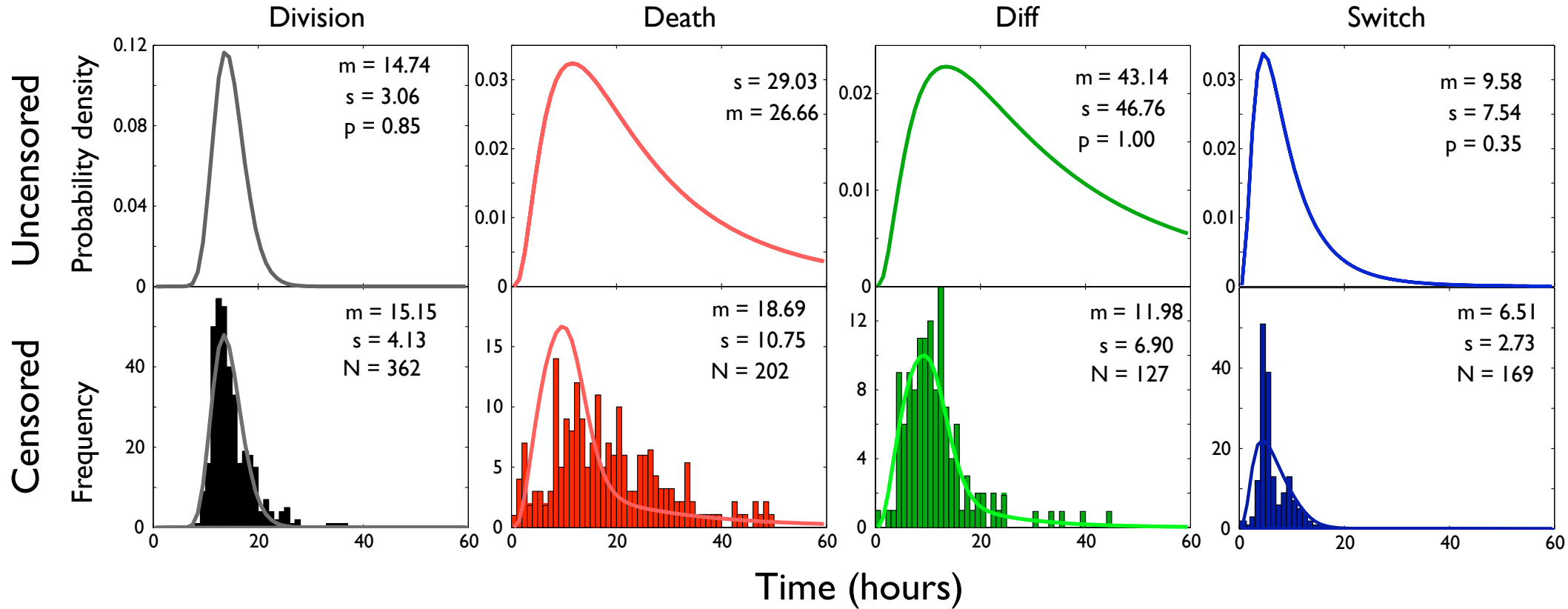


= 15 - parameters!



Statistical model (Duffy, Wellard)

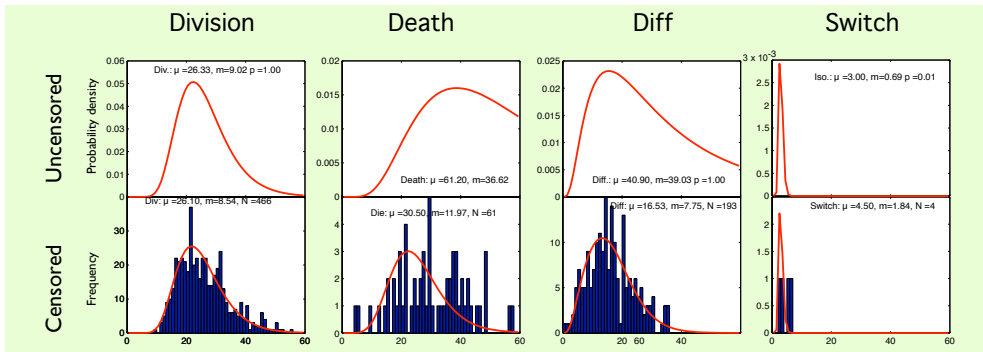
lognormal time to each event (m, s, f)
Independent with Censorship (death, div)



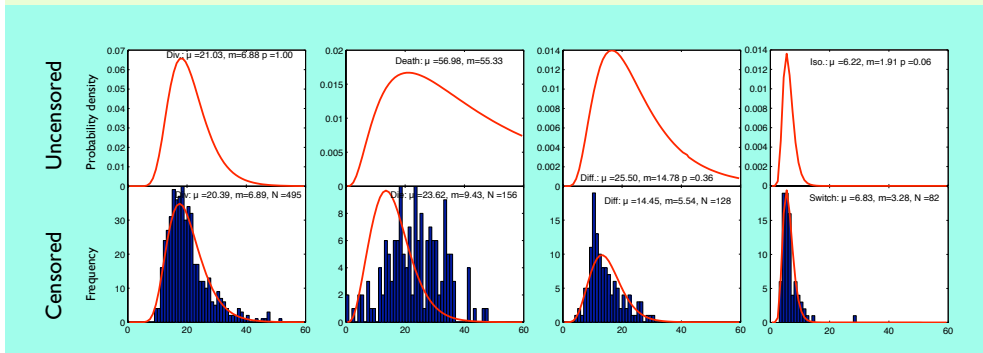
Generation 5

All generations

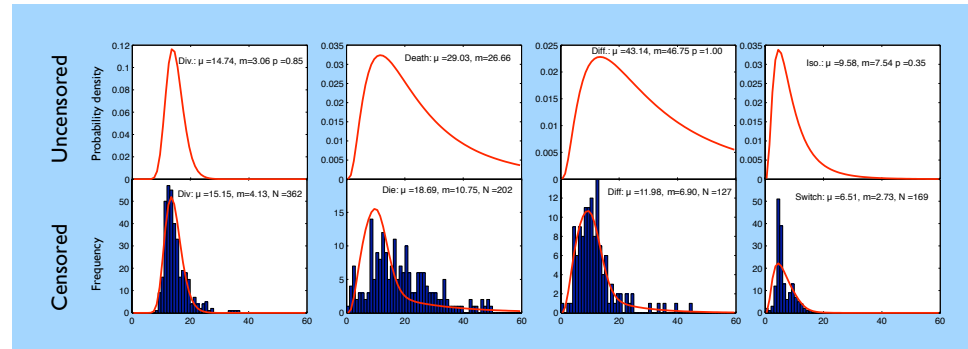
Generation 1



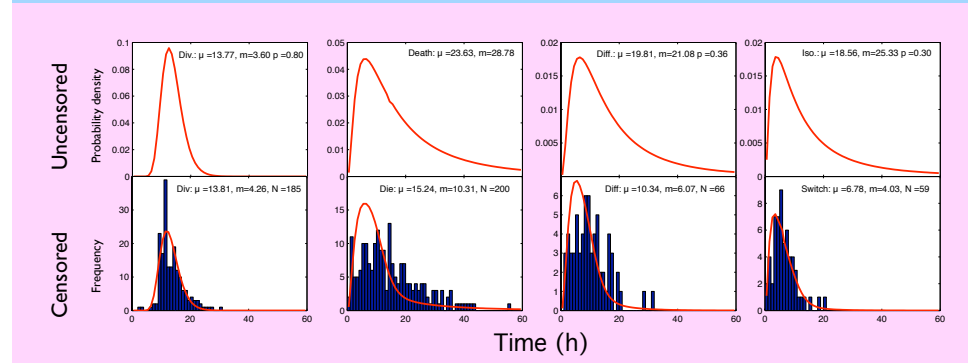
Generation 3



Generation 5

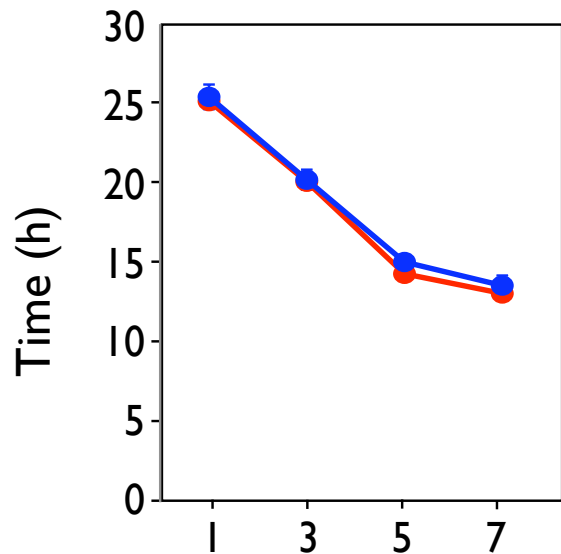


Generation 7

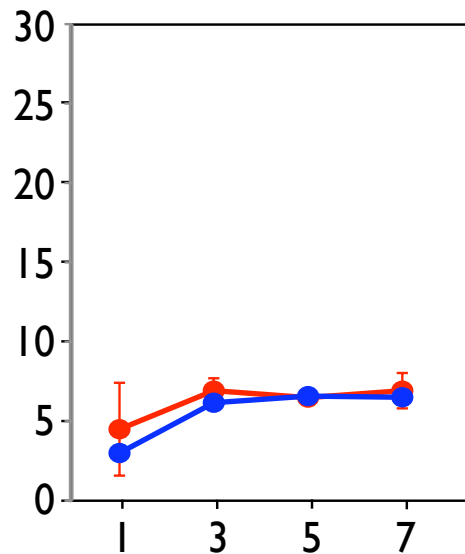


Time (h)

Division



Switch

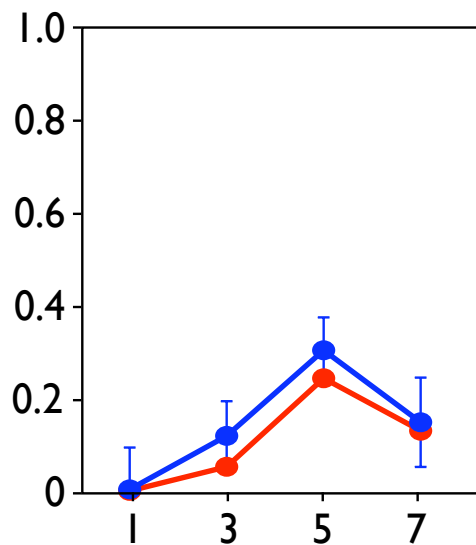
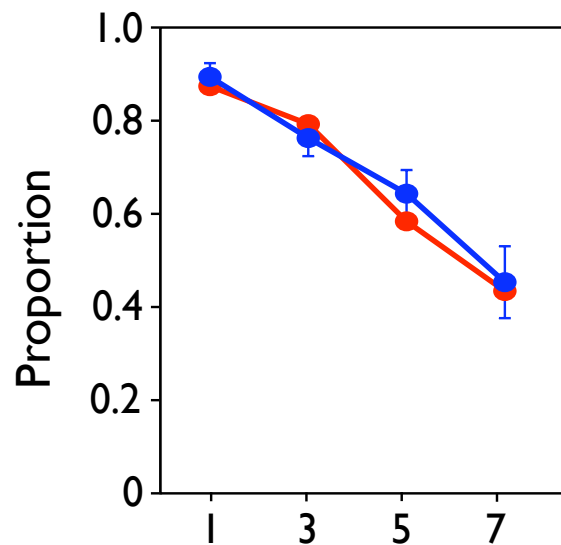


Fitted features

Time/Division

and

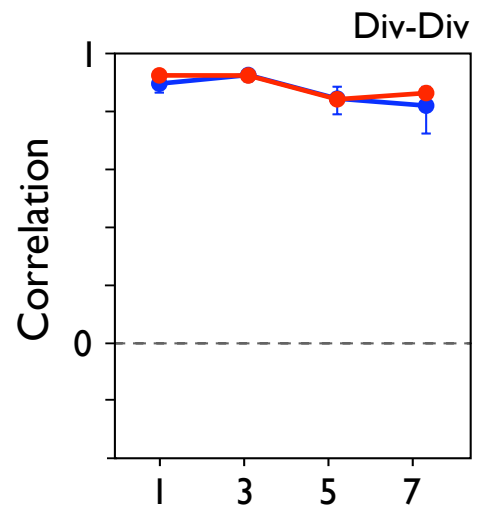
Proportion/division



Blue - measured

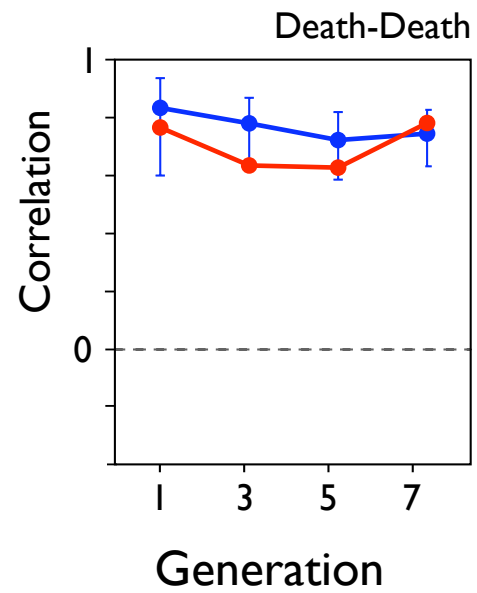
Red - model

Intercellular concordant



Fitted features

Inter-cellular Correlations



Blue - measured
Red - model

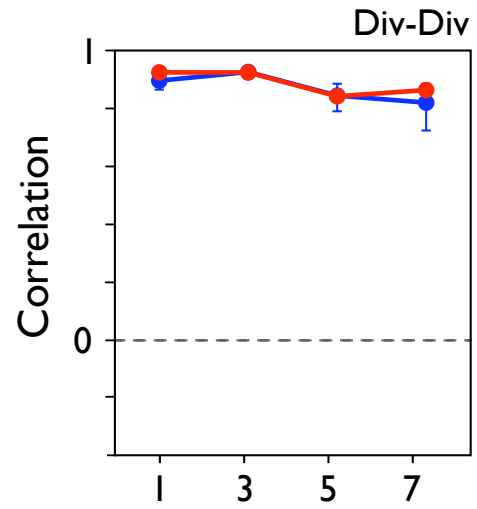
Passed test of getting back what you put in..

But can it explain the unexpected correlations?

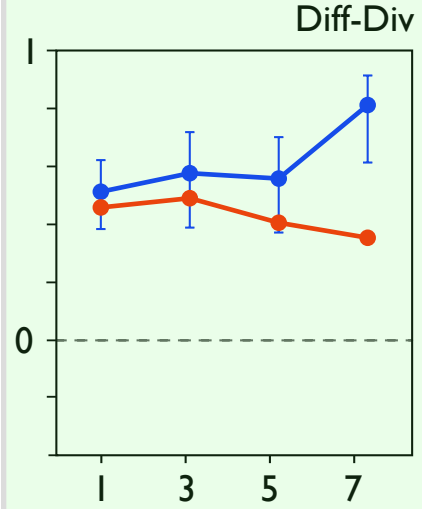
Intra cellular correlations

intercellular correlations of different fates of sibs

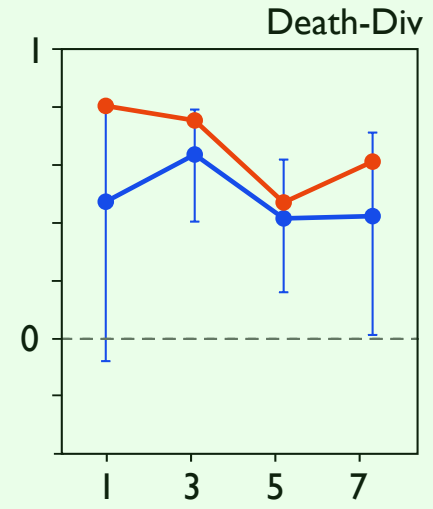
Intercellular concordant



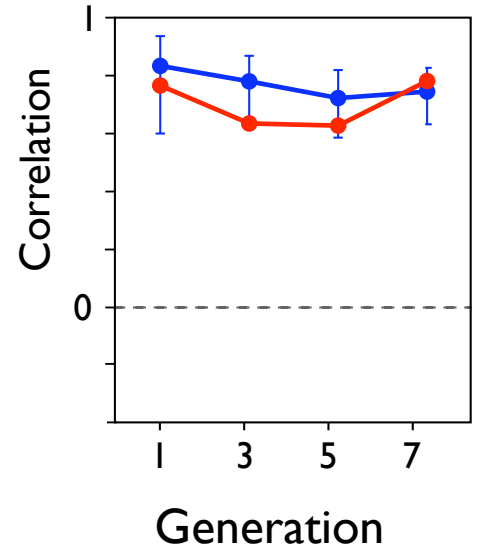
Intracellular



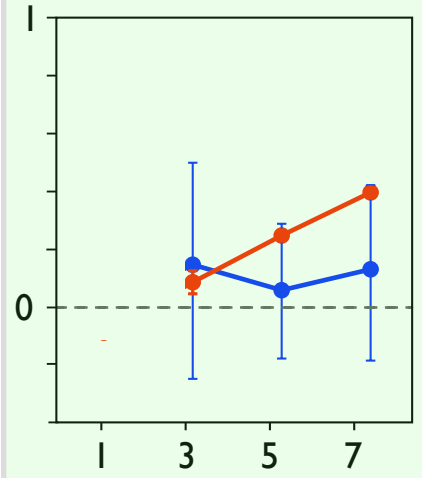
Intercellular non-concordant



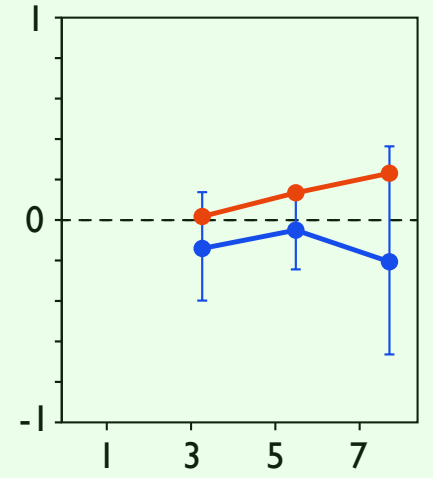
Death-Death



Death-Switch



Div-Switch

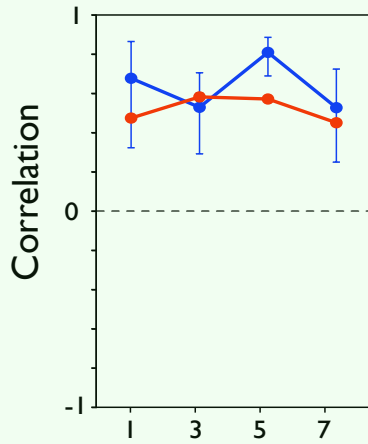


Blue - measured
Red - model

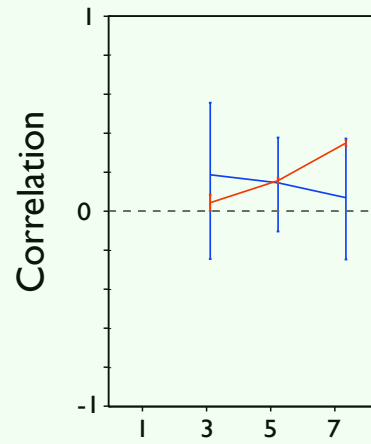
Intracellular

Intercellular non-concordant

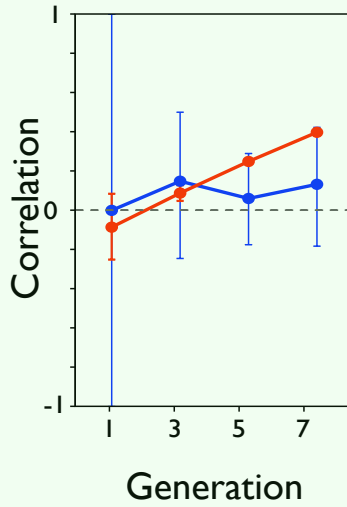
Diff-Death



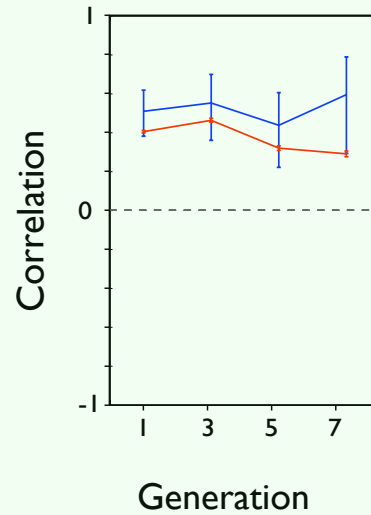
Death-Switch



Death-Switch



Div-Diff



Blue - measured
Red - model

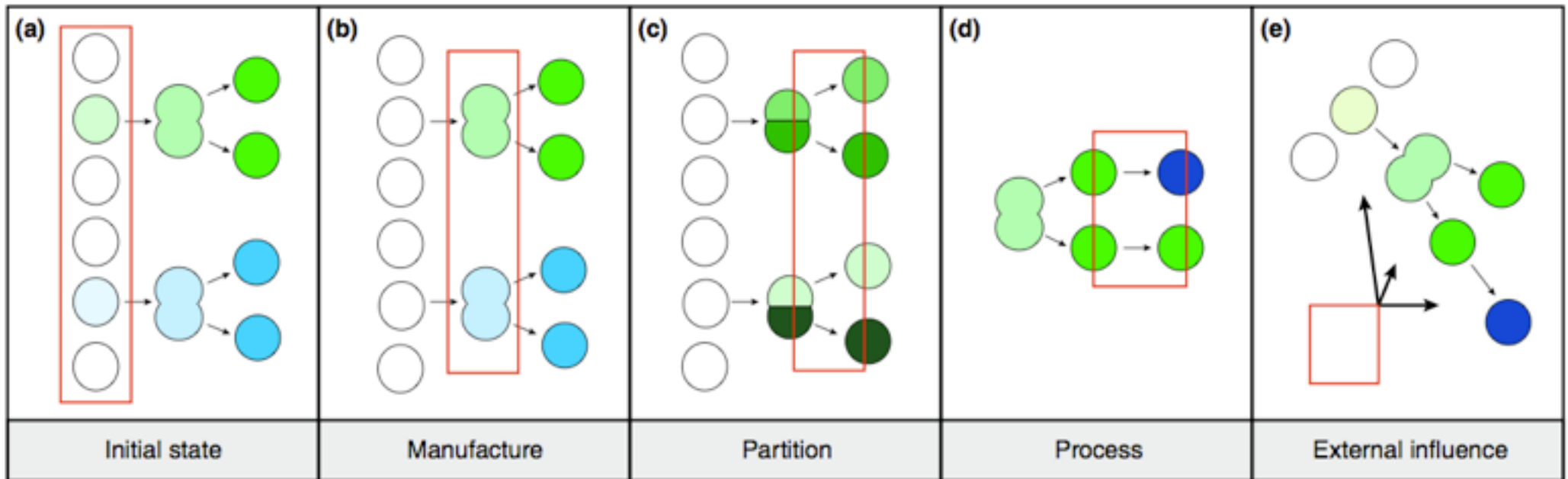
- Hallmarks of competition for fates apparent
- Asymmetric fates observed but conform to statistical likelihood

Activation-Induced B Cell Fates Are Selected by Intracellular Stochastic Competition

Ken R. Duffy,¹ Cameron J. Wellard,^{2,3} John F. Markham,⁴ Jie H. S. Zhou,^{2,3} Ross Holmberg,²
Edwin D. Hawkins,⁵ Jhagvaral Hasbold,^{2,3} Mark R. Dowling,^{2,3*} Philip D. Hodgkin^{2,3*†}

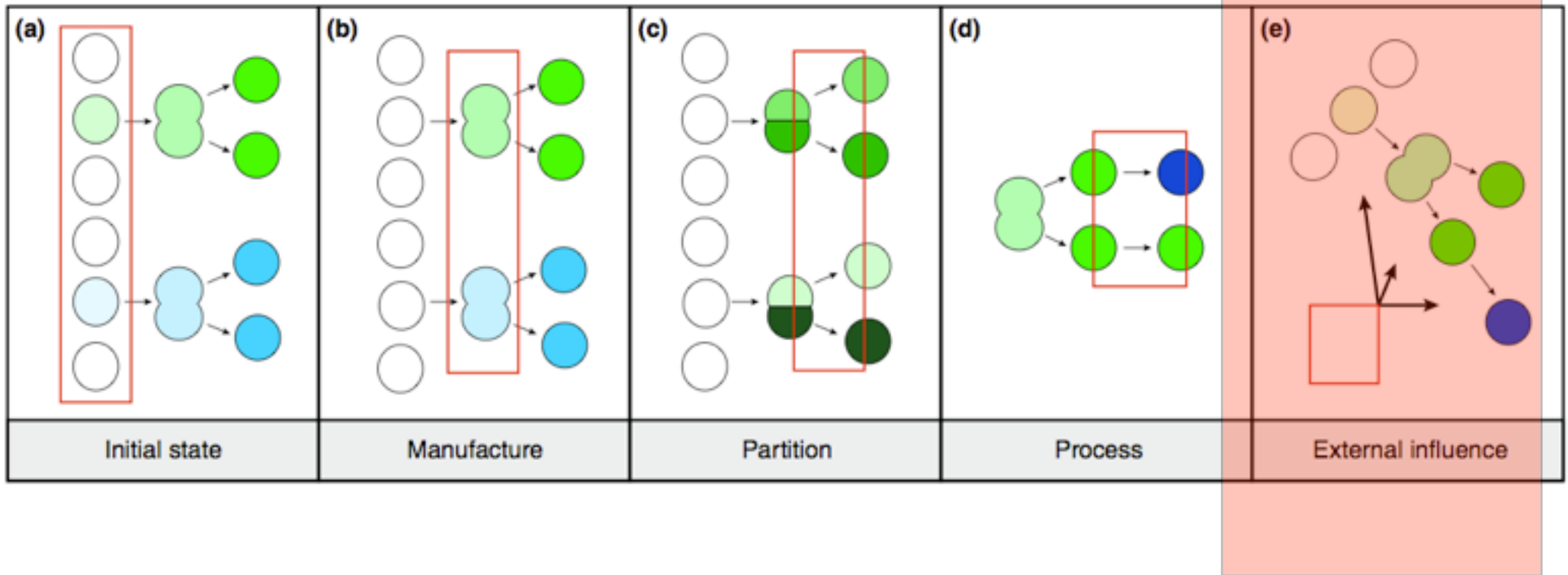
20 JANUARY 2012 VOL 335 SCIENCE

Homing in on the source of cellular heterogeneity



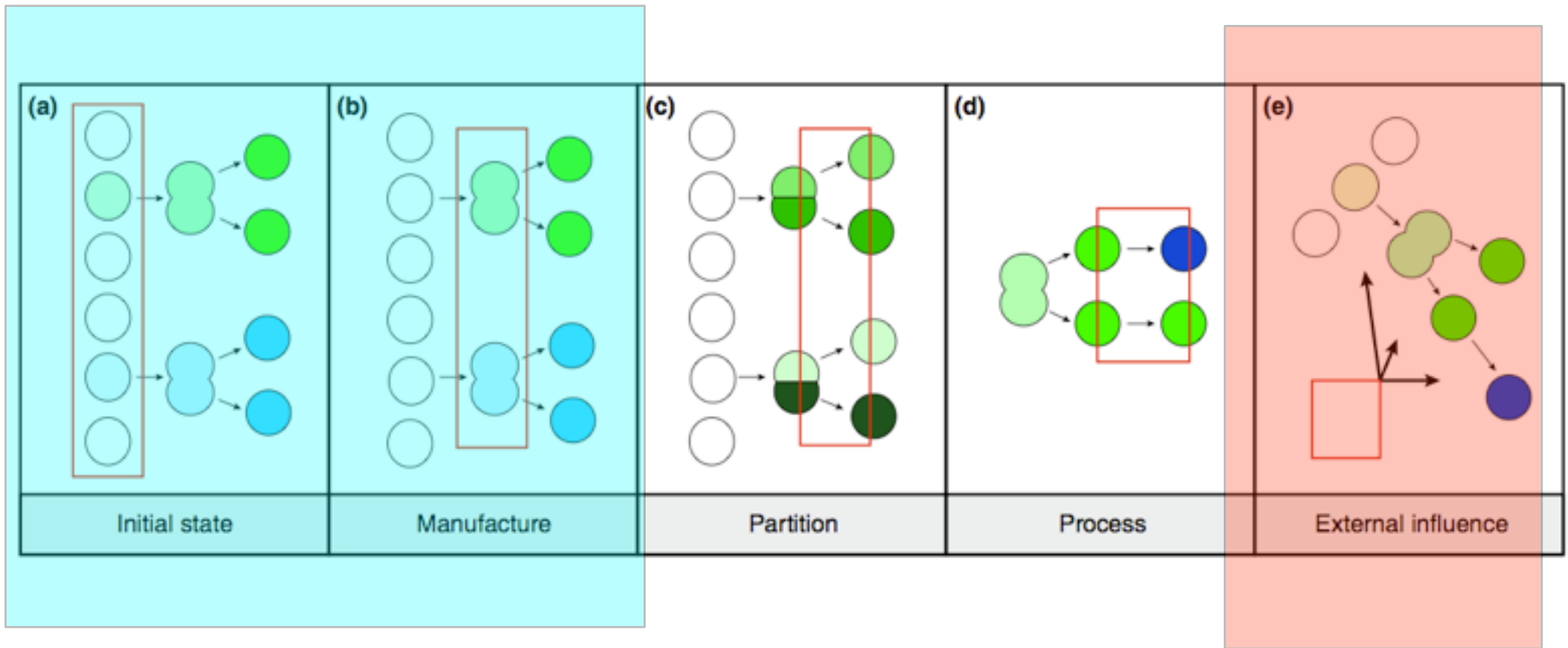
Duffy and Hodgkin, 2012 Trends in Cell Biology

Homing in on the source of cellular heterogeneity



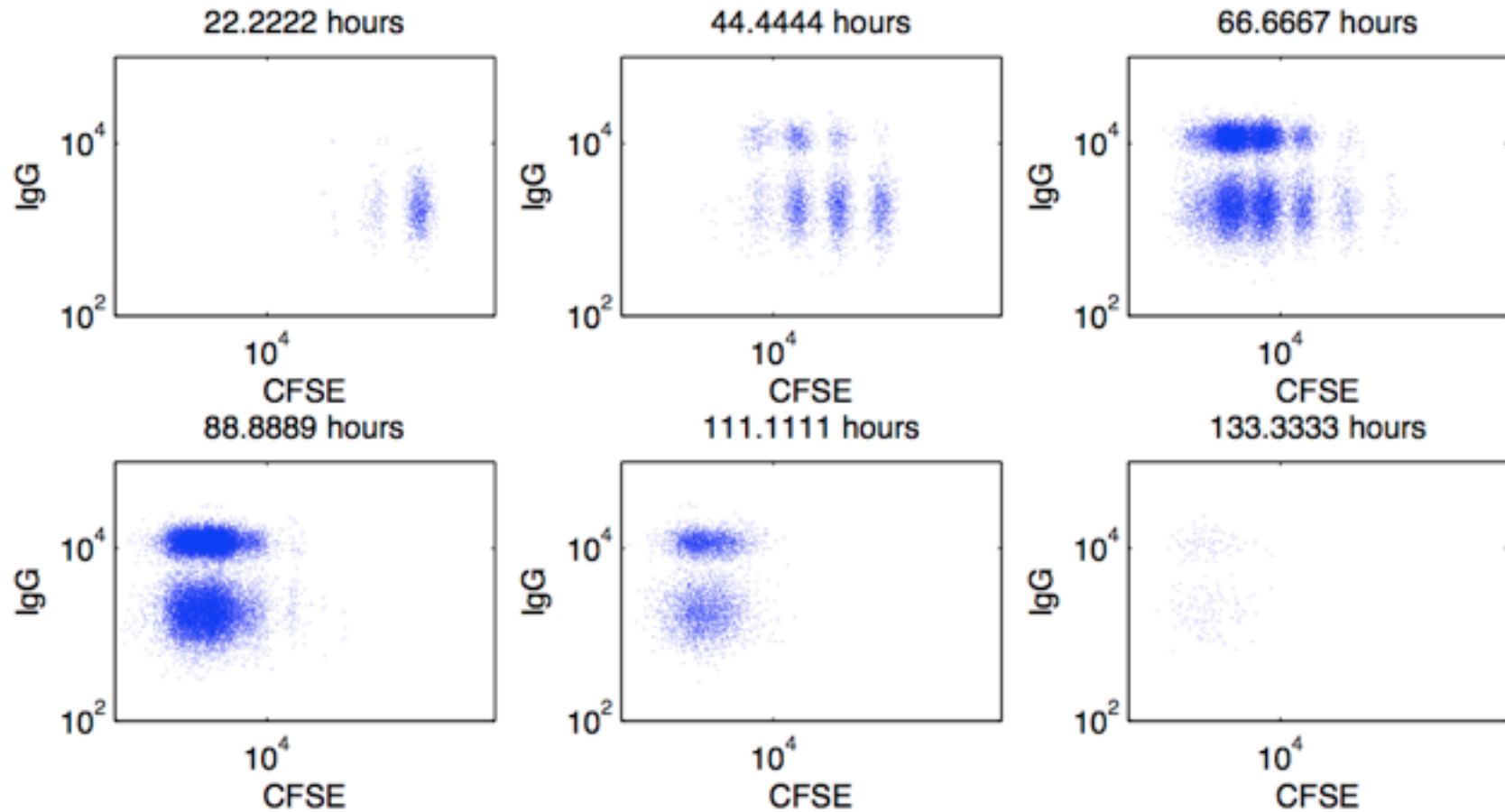
Duffy and Hodgkin, 2012 Trends in Cell Biology

Homing in on the source of cellular heterogeneity



Duffy and Hodgkin, 2012 Trends in Cell Biology

Computer flow cytometry!

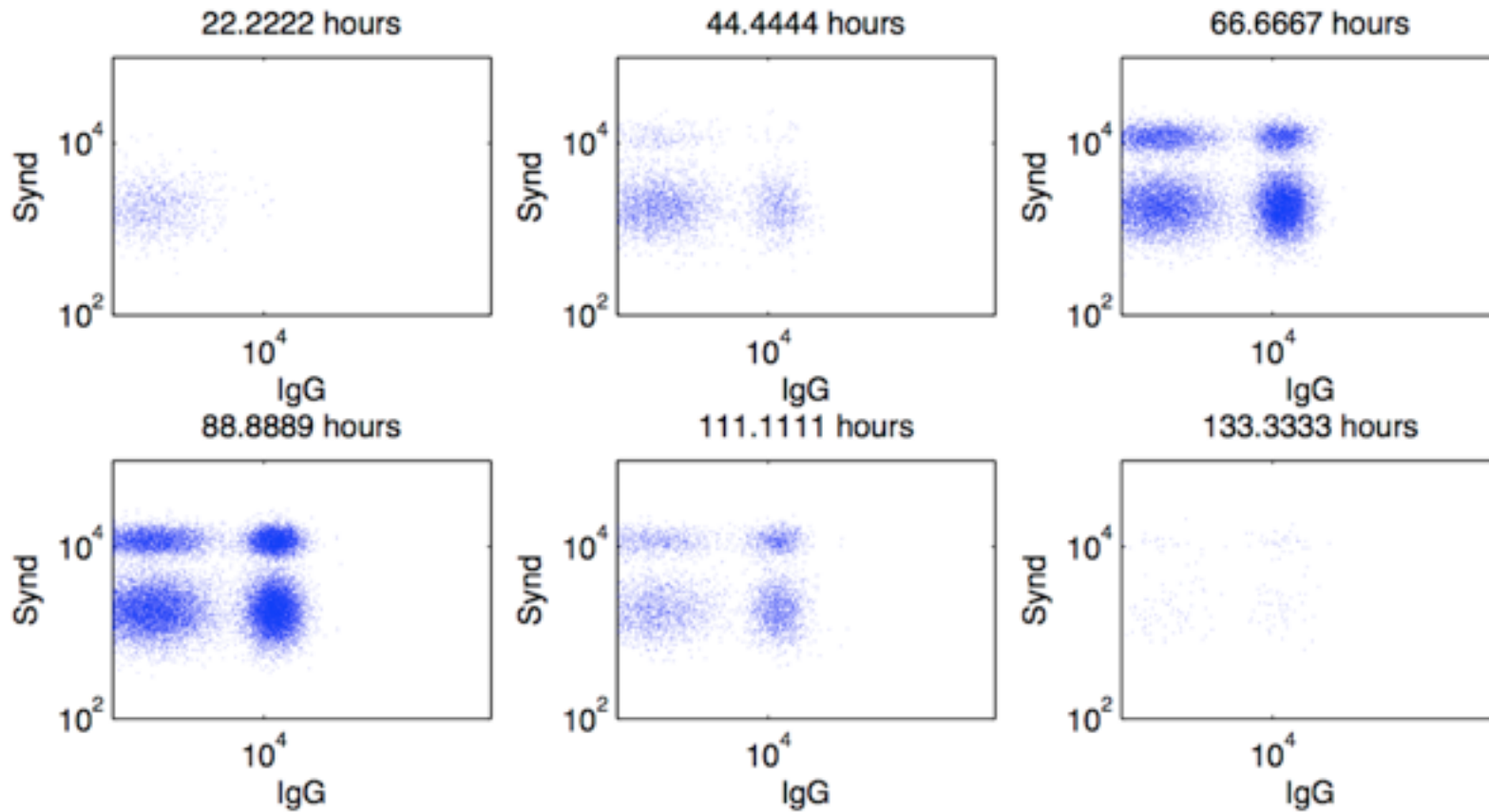


Run simulator for thousands of cells

Record features at every time for all cells

'Plot as if real cells'

Computer flow cytometry!



Run simulator for thousands of cells

Record features at every time for all cells

'Plot as if real cells'

Conclusions

Complex features of lymphocyte control result from a form of modulated 'randomness' of times to different fates set in competition in each cell

By manipulating the frequencies and times to change, by signals and cell division, a robust system for allocating different cells to large number of different fates is created

Combinatorially for example - just 20 independent surface marker 'machines' - gives one million possible 'phenotypes'

Thanks to..

B cell proliferation/ diffn/death

*Mark Dowling
*Has Hasbold
*Edwin Hawkins
*Jie Zhou
Marian Turner
Nadine Taubenheim

Differentiation simulator Image analysis/model development

*Cam Wellard

*Ken Duffy [Hamilton - Maynooth]
Vijay Subramanian [Hamilton - Maynooth]



Walter+Eliza Hall
Institute of Medical Research

Systems B cell Program WEHI

Dave Tarlinton
Lynn Corcoran
*Steve Nutt

Axel Kallies

Imaging & microwells

*John Markham [NICTA]
*Mark Dowling

*Ed Hawkins
Liam McGuinness
Ross Holmberg
**Jie Zhou

Daniel Day (Swin)
Sarah Russell (PMac)

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