

Limit theorems for Bifurcating Auto-Regressive processes with missing data

Benoîte de Saporta

Anne Gégout-Petit

Laurence Marsalle

Université de Bordeaux and INRIA Bordeaux Sud-Ouest

Université de Lille 1

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Outline

- 1 Motivation
- 2 Model
- 3 Limit theorems
- 4 Symmetry tests
- 5 Application

Modelisation of cell lineage data



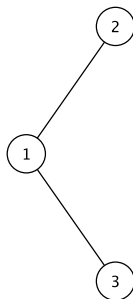
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Escherichia coli

Modelisation of cell lineage data



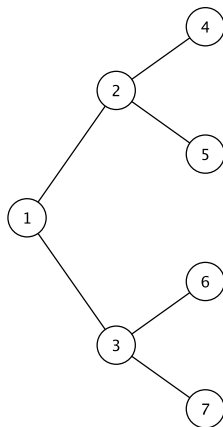
Escherichia coli



Modelisation of cell lineage data



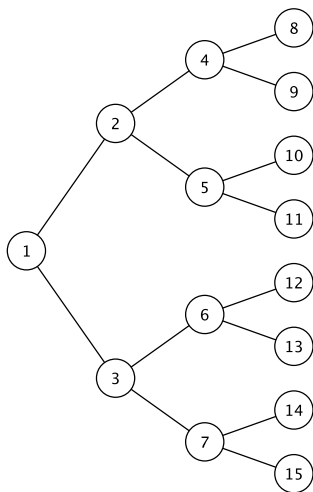
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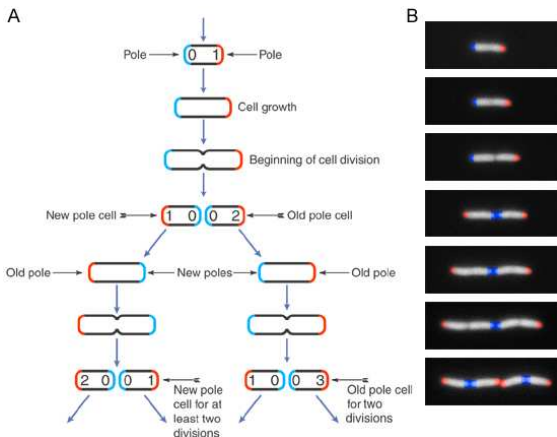
Modelisation of cell lineage data



Escherichia coli

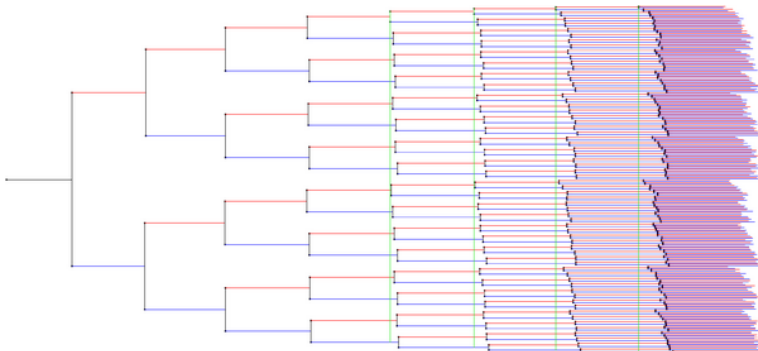


Division of Escherichia coli



From Stewart et al. *PLoS Biol.* 2005.

Do single cell organisms age?



Mean growth rate of *E. coli* for 94 genealogies up to 9 generations
From Stewart et al. *PLoS Biol.* 2005.

Aim of the talk

Aims

- propose a **bifurcating auto-regressive** model to study lineages one by one
- take possibly **missing data** into account
- study the **asymptotic properties** of the estimators of the parameters
- present **symmetry tests**
- investigate **simulated** and **real** data

Asymmetric BAR process

Bifurcating autoregressive process **BAR** = autoregressive process indexed by a **binary tree**.

$$\begin{cases} X_{2n} &= a + b X_n + \varepsilon_{2n} \\ X_{2n+1} &= c + d X_n + \varepsilon_{2n+1} \end{cases}$$

①

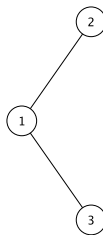
- X_1 ancestor
- $(\varepsilon_{2n}, \varepsilon_{2n+1})$ noise
- $0 < \max(|b|, |d|) < 1$

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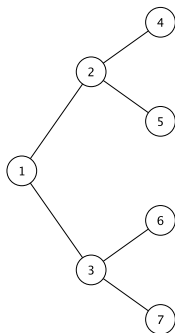


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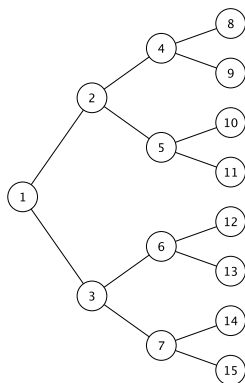


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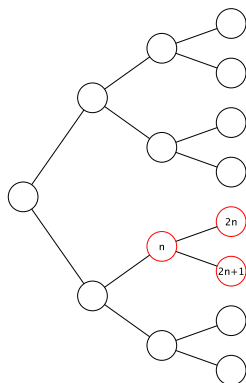


Asymmetric BAR process

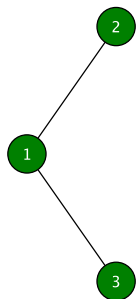
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Missing data

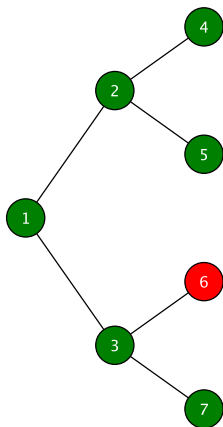


Observation process: $(\delta_n)_{n>0}$, taking values in $\{0, 1\}$

X_n observed if $\delta_n = 1$, else $\delta_n = 0$

If a cell is not observed, all its descendants are not observed.

Missing data

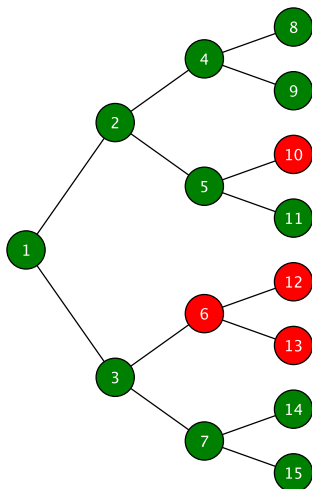


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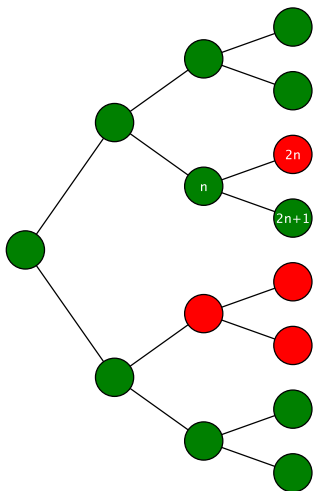


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Two-type Galton-Watson process

Reproduction laws

- all individuals reproduce **independently**
- the reproduction laws depend on the **type** of the **mother** and **daughter**
- possible **asymmetry** in the reproduction laws

$p^0(0, 0)$: proba that an **even** mother has **0** even and **0** odd daughter

$p^0(1, 0)$: proba that an **even** mother has **1** even and **0** odd daughter

$p^0(0, 1)$: proba that an **even** mother has **0** even and **1** odd daughter

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Two-type Galton-Watson process

Reproduction laws

- all individuals reproduce **independently**
- the reproduction laws depend on the **type** of the **mother** and **daughter**
- possible **asymmetry** in the reproduction laws

$p^1(0, 0)$: proba that an **odd** mother has **0** even and **0** odd daughter

$p^1(1, 0)$: proba that an **odd** mother has **1** even and **0** odd daughter

$p^1(0, 1)$: proba that an **odd** mother has **0** even and **1** odd daughter

$p^1(1, 1)$: proba that an **odd** mother has **1** even and **1** odd daughter

Extinction

Descendance matrix

$$P = \begin{pmatrix} p_{00} & p_{01} \\ p_{10} & p_{11} \end{pmatrix}$$

$p_{i0} = p^i(1, 0) + p^i(1, 1)$: expected number of **even** daughters of an individual of type i

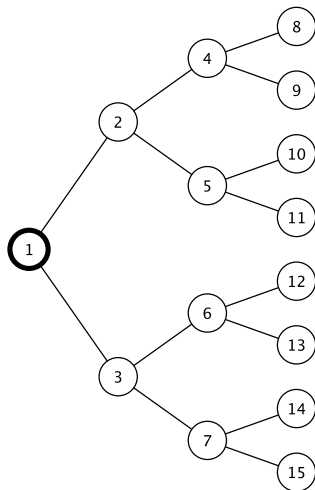
$p_{i1} = p^i(0, 1) + p^i(1, 1)$: expected number of **odd** daughters of an individual of type i

Criteria for extinction

π spectral radius of P

- if $\pi < 1$, extinction is almost sure
- if $\pi \geq 1$, extinction has probability < 1

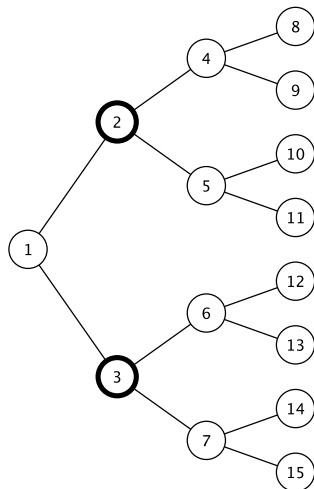
Generations



Fully observed generation 0:

$$\mathbb{G}_0 = \{1\}$$

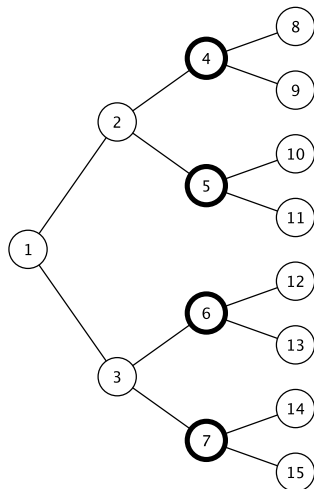
Generations



Fully observed generation 1:

$$\mathbb{G}_1 = \{2, 3\}$$

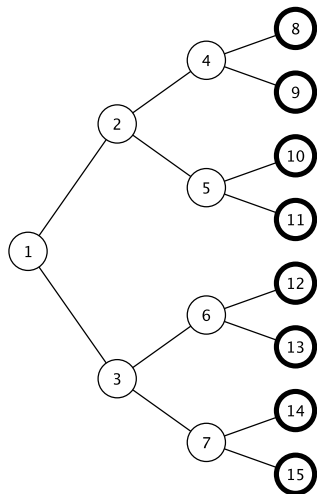
Generations



Fully observed generation 2:

$$\mathbb{G}_2 = \{4, 5, 6, 7\}$$

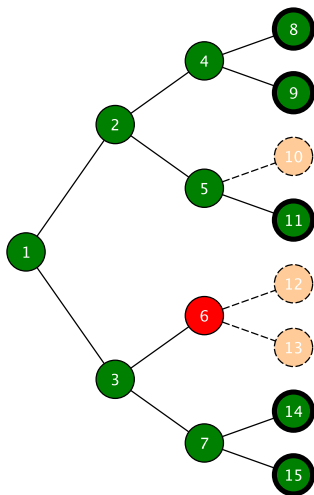
Generations



Fully observed generation n :

$$\mathbb{G}_n = \{2^n, 2^n + 1, \dots, 2^{n+1} - 1\}$$

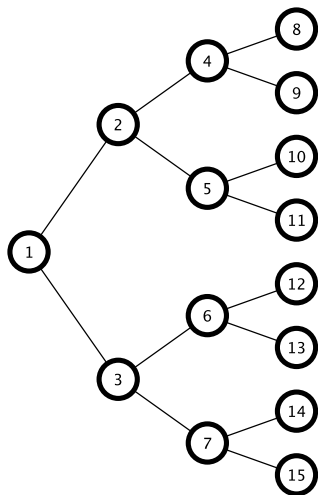
Generations



Partially observed generation n :

$$\mathbb{G}_n^* = \{k \in \mathbb{G}_n ; \delta_k = 1\}$$

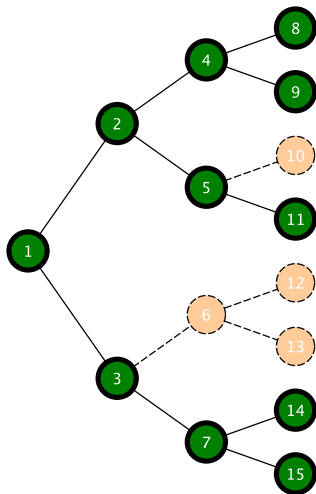
Generations



Fully observed tree up to generation n :

$$\mathbb{T}_n = \cup_{\ell=0}^n \mathbb{G}_\ell$$

Generations



Partially observed tree up to generation n :

$$\mathbb{T}_n^* = \{k \in \mathbb{T}_n ; \delta_k = 1\} = \cup_{\ell=0}^n \mathbb{G}_\ell^*$$

Assumptions

BAR model

$$\begin{cases} X_{2n} &= a + b X_n + \varepsilon_{2n} \\ X_{2n+1} &= c + d X_n + \varepsilon_{2n+1} \end{cases}$$

Assumption

- **independence** between (δ_k) and (X_k) and $(\varepsilon_{2k}, \varepsilon_{2k+1})$

Estimation of $\theta = (a, b, c, d)^t$

Least squares estimator minimizes

$$\Delta_n(\theta) = \frac{1}{2} \sum_{k \in \mathbb{T}_{n-1}} \delta_{2k} (X_{2k} - a - bX_k)^2 + \delta_{2k+1} (X_{2k+1} - c - dX_k)^2.$$

Least squares estimator

LS Estimator for θ

$$\hat{\theta}_n = \begin{pmatrix} \hat{a}_n \\ \hat{b}_n \\ \hat{c}_n \\ \hat{d}_n \end{pmatrix} = \Sigma_{n-1}^{-1} \sum_{k \in \mathbb{T}_{n-1}} \begin{pmatrix} \delta_{2k} X_{2k} \\ \delta_{2k} X_k X_{2k} \\ \delta_{2k+1} X_{2k+1} \\ \delta_{2k+1} X_k X_{2k+1} \end{pmatrix}$$

where

$$\Sigma_n = \begin{pmatrix} S_n^0 & 0 \\ 0 & S_n^1 \end{pmatrix}$$

$$S_n^0 = \sum_{k \in \mathbb{T}_n} \delta_{2k} \begin{pmatrix} 1 & X_k \\ X_k & X_k^2 \end{pmatrix} \quad S_n^1 = \sum_{k \in \mathbb{T}_n} \delta_{2k+1} \begin{pmatrix} 1 & X_k \\ X_k & X_k^2 \end{pmatrix}$$

Strong consistency with convergence rate

Theorem

Under moment assumptions on the noise sequence

$$\mathbb{1}_{\{|G_n^*|>0\}} \|\hat{\theta}_n - \theta\|^2 = \mathbb{1}_{\{|G_n^*|>0\}} \mathcal{O}\left(\frac{\log |\mathbb{T}_{n-1}^*|}{|\mathbb{T}_{n-1}^*|}\right)$$

Proof: [martingales](#)

Main martingale

$\widehat{\theta}_n - \theta = \Sigma_{n-1}^{-1} M_n$, where (M_n) is a **martingale** for the filtration of generations and the whole observation process

$$M_n = \sum_{k \in \mathbb{T}_{n-1}} \begin{pmatrix} \delta_{2k} \varepsilon_{2k} \\ \delta_{2k} X_k \varepsilon_{2k} \\ \delta_{2k+1} \varepsilon_{2k+1} \\ \delta_{2k+1} X_k \varepsilon_{2k+1} \end{pmatrix}$$

$(M_n)_{n \geq 1}$ square integrable with **increasing process** $\langle M \rangle_n = \Gamma_{n-1}$

$$\Gamma_n = \begin{pmatrix} \sigma^2 \mathbf{S}_n^0 & \rho \mathbf{S}_n^{0,1} \\ \rho \mathbf{S}_n^{0,1} & \sigma^2 \mathbf{S}_n^1 \end{pmatrix} \quad \text{and} \quad \mathbf{S}_n^{0,1} = \sum_{k \in \mathbb{T}_n} \delta_{2k} \delta_{2k+1} \begin{pmatrix} 1 & X_k \\ X_k & X_k^2 \end{pmatrix}$$

Martingale convergence results

(M_n) scalar \mathcal{F} -martingale bounded in L^2

$$\Delta M_{n+1} = M_{n+1} - M_n$$

Increasing process $\langle M \rangle_n = \sum_{k=0}^n \mathbb{E}[(\Delta M_{k+1})^2 \mid \mathcal{F}_k]$

Convergence of scalar L^2 martingales

If $\lim_{n \rightarrow \infty} \langle M \rangle_n = +\infty$, then $\frac{M_n}{\langle M \rangle_n} \rightarrow 0$ a.s.

+ moment conditions then $\left(\frac{M_n}{\langle M \rangle_n}\right)^2 = \mathcal{O}\left(\frac{\log(\langle M \rangle_n)}{\langle M \rangle_n}\right)$ a.s.

Similar results for vector-valued martingales.

Here $\langle M \rangle_n$ is a 4×4 -matrix

Convergence of the increasing process

Theorem

$\mathbb{1}_{\{|\mathbb{G}_n^*| > 0\}} \frac{\langle \mathbf{M} \rangle_n}{|\mathbb{T}_n^*|}$ converges a.s. to a fixed definite positive matrix.

Sketch of the proof

Laws of large numbers for

- the observation process (δ_k)
- the observed noise $(\delta_k \varepsilon_k)$
- the observed BAR $(\delta_{2k} X_k), \delta_{2k+1} X_k, \delta_{2k} X_k^2, \dots$

via [martingale](#) methods for various filtrations and using the [auto-regressive](#) structure

Central limit theorem

$\bar{\mathcal{E}}$ non-extinction set, complementary set of

$$\mathcal{E} = \bigcup_{n \geq 1} \{|\mathbb{G}_n^*| = 0\}.$$

New probability

$$\mathbb{P}_{\bar{\mathcal{E}}}(A) = \frac{\mathbb{P}(A \cap \bar{\mathcal{E}})}{\mathbb{P}(\bar{\mathcal{E}})}$$

Theorem

$$\sqrt{|\mathbb{T}_{n-1}^*|}(\hat{\boldsymbol{\theta}}_n - \boldsymbol{\theta}) \xrightarrow{\mathcal{L}} \mathcal{N}(0, \boldsymbol{\Sigma}^{-1} \boldsymbol{\Gamma} \boldsymbol{\Sigma}^{-1}) \quad \text{on } (\bar{\mathcal{E}}, \mathbb{P}_{\bar{\mathcal{E}}}).$$

Test for the coefficients (a, b) vs (c, d)

H0: $(a, b) = (c, d)$ vs **H1**: $(a, b) \neq (c, d)$

Test statistic :

$$\mathbf{Y}_n^2 = |\mathbb{T}_{n-1}^*| (\hat{a}_n - \hat{c}_n, \hat{b}_n - \hat{d}_n) \Delta^{-1} (\hat{a}_n - \hat{c}_n, \hat{b}_n - \hat{d}_n)^t$$

where

$$\Delta = \mathbf{d} \mathbf{g}^t \Sigma^{-1} \Gamma \Sigma^{-1} \mathbf{d} \mathbf{g} \quad \text{and} \quad \mathbf{d} \mathbf{g} = \begin{pmatrix} 1 & 0 & -1 & 0 \\ 0 & 1 & 0 & -1 \end{pmatrix}^t$$

Theorem

Under the null hypothesis (**H0**)

$$\mathbf{Y}_n^2 \xrightarrow{\mathcal{L}} \chi^2(2) \quad \text{on } (\bar{\mathcal{E}}, \mathbb{P}_{\bar{\mathcal{E}}})$$

and under the alternative hypothesis (**H1**)

$$\lim_{n \rightarrow \infty} \|\mathbf{Y}_n^2\| = +\infty \quad \text{a.s. on } (\bar{\mathcal{E}}, \mathbb{P}_{\bar{\mathcal{E}}})$$

Test for the fixed points $a/(1-b)$ vs $c/(1-d)$

H0: $a/(1-b) = c/(1-d)$ vs **H1:** $a/(1-b) \neq c/(1-d)$

Test statistic

$$Y_n^2 = |\mathbb{T}_{n-1}^*| \Delta^{-1} (\hat{a}_n/(1-\hat{b}_n) - \hat{c}_n/(1-\hat{d}_n))^2$$

where

$$\Delta = \mathbf{d}\mathbf{g}^t \boldsymbol{\Sigma}^{-1} \boldsymbol{\Gamma} \boldsymbol{\Sigma}^{-1} \mathbf{d}\mathbf{g} \quad \text{and} \quad \mathbf{d}\mathbf{g} = (1/(1-b), a/(1-b)^2, -1/(1-d), -c/(1-d)^2)^t$$

Theorem

Under the null hypothesis (**H0**)

$$Y_n^2 \xrightarrow{\mathcal{L}} \chi^2(1) \quad \text{on } (\bar{\mathcal{E}}, \mathbb{P}_{\bar{\mathcal{E}}})$$

and under the alternative hypothesis (**H1**), one has

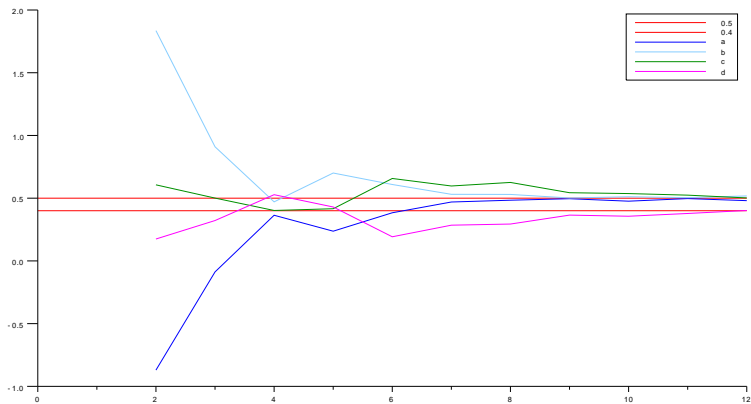
$$\lim_{n \rightarrow \infty} Y_n^2 = +\infty \quad \text{a.s. on } (\bar{\mathcal{E}}, \mathbb{P}_{\bar{\mathcal{E}}})$$

Simulated data - parameters

Galton-Watson	$p^0(0, 0)$	$p^0(0, 1)$	$p^0(1, 0)$	$p^0(1, 1)$
symmetric case	0.02	0.04	0.04	0.90
non symmetric case	0.02	0.04	0.04	0.90
	$p^1(0, 0)$	$p^1(0, 1)$	$p^1(1, 0)$	$p^1(1, 1)$
symmetric case	0.02	0.04	0.04	0.90
non symmetric case	0.015	0.075	0.075	0.0835

BAR	a	b	c	d
symmetric case	0.5	0.5	0.5	0.5
non symmetric case	0.5	0.5	0.5	0.4

Simulated data - estimation (non symmetric case)



Simulated data - Test (a, b) vs (c, d) -1000 simulations

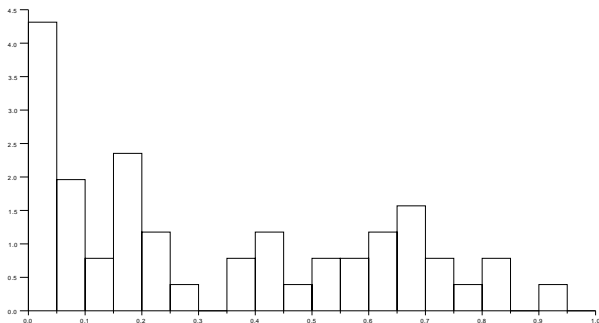
number n of generations	H0 % p-value<0.05	H1 % pvalue<0.05
$n = 7$	37.4	6.6
$n = 8$	53.6	5.5
$n = 9$	71.1	5.5
$n = 10$	86.8	6.3
$n = 11$	95.7	5.9

Simulated data - Test $a/(1 - b)$ vs $c/(1 - d)$ - 1000 simulations

number n of generations	H0	H1
	% p-value < 0.05	% p-value < 0.05
$n = 7$	23.1	2.2
$n = 8$	41.3	3.3
$n = 9$	64.6	3.8
$n = 10$	82.9	4.7
$n = 11$	94.5	5.5

Real data - growth rate of Escherichia coli

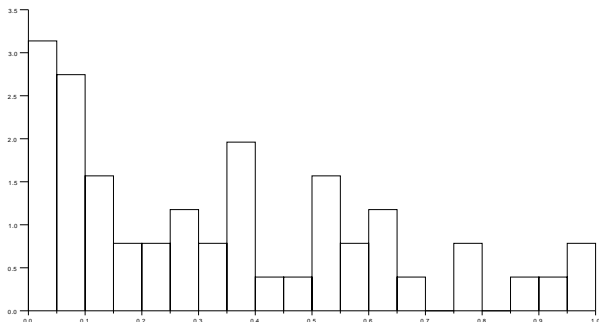
51 genealogies of cells dividing between 8 and 9 times



p-values First test (a, b) vs (c, d)

Real data - growth rate of Escherichia coli

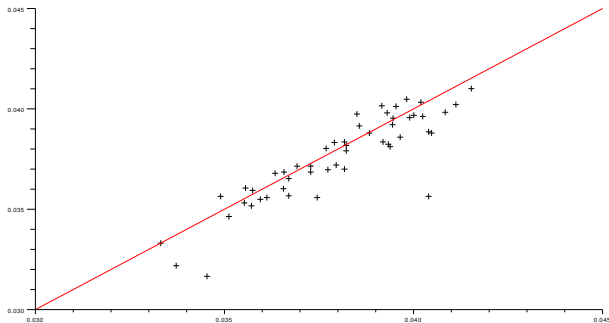
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p-values Second test $a/(1-b)$ vs $c/(1-d)$

Real data - growth rate of Escherichia coli

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Fixed points

References

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