

Exponential Growth or Death?

Peter Jagers

First Franco-Nordic Mathematics Congress

Reykjavik, 6-9 January 2005.



Malthus's Law:

“A population, when unchecked, increases in a geometrical ratio. Subsistence increases only in an arithmetical ratio. A slight acquaintance with numbers will shew the immensity of the first power in comparison of the second.”

(An Essay on the Principle of Population as it Affects the Future Improvement of Society..., 1798)



Growth or Extinction:

- “Quam ob causam maxime ridiculæ sunt eorum incredulorum hominum obiectones, qui negant tam brevi tempore spatio ab uno homine universam terram incolis impleri potuisse.” (Leonhard Euler, *Introductio in analysin infinitorum*, 1748)

Growth or Extinction:

- “Quam ob causam maxime ridiculæ sunt eorum incredulorum hominum obiectones, qui negant tam brevi tempore spatio ab uno homine universam terram incolis impleri potuisse.” (Leonhard Euler, *Introductio in analysin infinitorum*, 1748)
- In 200 years (from 1583 to 1783) 379 out of 487 Berne families died out. (Malthus)



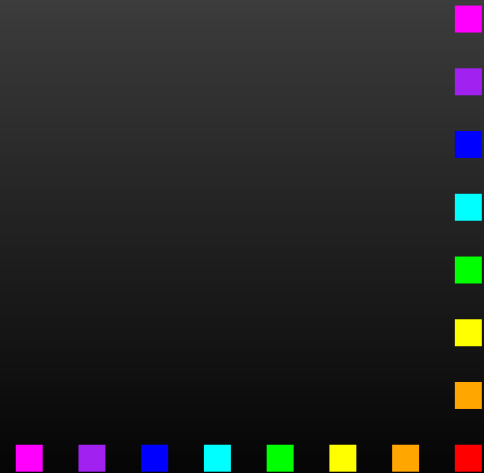
Growth or Extinction:

- “Quam ob causam maxime ridiculæ sunt eorum incredulorum hominum obiectones, qui negant tam brevi tempore spatio ab uno homine universam terram incolis impleri potuisse.” (Leonhard Euler, *Introductio in analysin infinitorum*, 1748)
- In 200 years (from 1583 to 1783) 379 out of 487 Berne families died out. (Malthus)
- I. J. Bienaymé (1845) and F. Galton (1873).



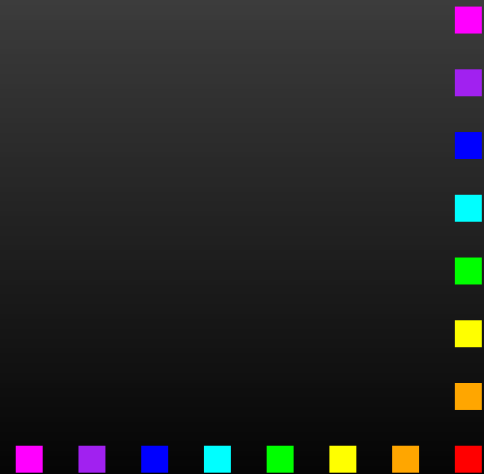
The Malthus – Bienaymé – Galton Dichotomy

- All (decent) *branching* processes either grow exponentially or die out.



The Malthus – Bienaymé – Galton Dichotomy

- All (decent) *branching* processes either grow exponentially or die out.
- Branching populations are sets of *independently* reproducing individuals.



The Malthus – Bienaymé – Galton Dichotomy

- All (decent) *branching* processes either grow exponentially or die out.
- Branching populations are sets of *independently* reproducing individuals.
- These can be of various *types*. Individuals of the same type have the same *reproduction distribution*.

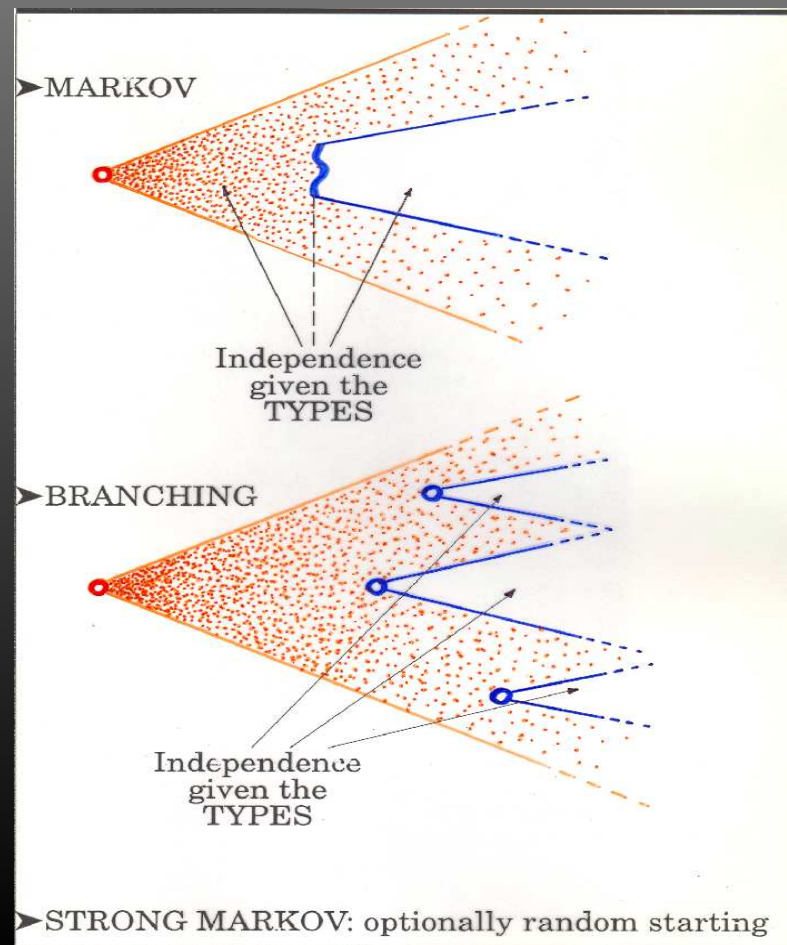
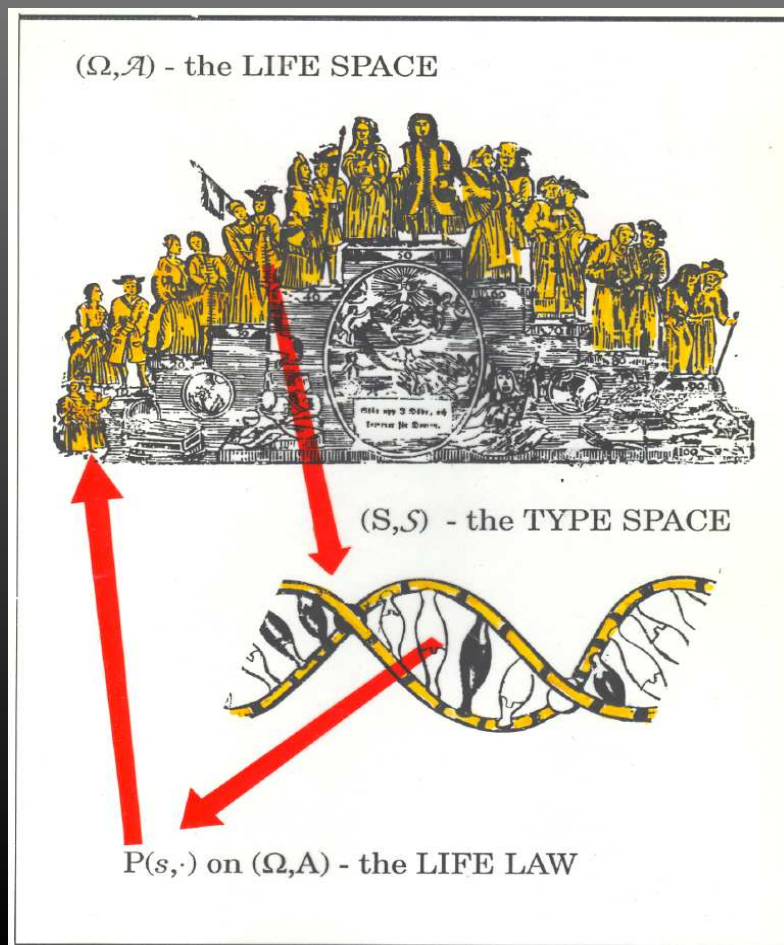


The Malthus – Bienaymé – Galton Dichotomy

- All (decent) *branching* processes either grow exponentially or die out.
- Branching populations are sets of *independently* reproducing individuals.
- These can be of various *types*. Individuals of the same type have the same *reproduction distribution*.
- Individual reproduction is any point process over (mother's) age \times (child's) type.



The Markov and Branching Properties



What if a population is “checked”?

Consider non-negative X_1, X_2, \dots such that $X_n = 0 \Rightarrow X_{n+1} = 0$. Suppose that there is a history-independent risk of extinction:
For any x there is a $\delta > 0$ such that

$$\mathbf{P}(\exists n; X_n = 0 \mid X_1, \dots, X_k) \geq \delta,$$

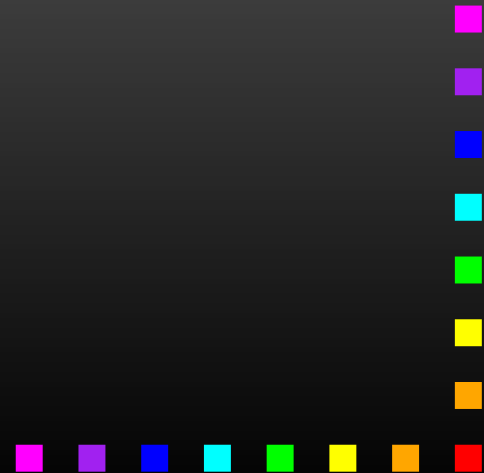
if only $X_k \leq x$. Then, with probability one either

- there is an n such that all $X_k = 0$ for $k \geq n$ or
- $X_k \rightarrow \infty$ as $k \rightarrow \infty$.



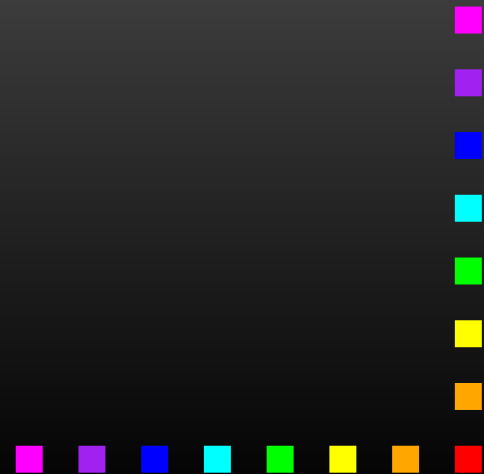
Branching with Dependence

- Local interaction – between relatives



Branching with Dependence

- Local interaction – between relatives
- Stationary or i.i.d random environments – independent branching | environment



Branching with Dependence

- Local interaction – between relatives
- Stationary or i.i.d random environments – independent branching | environment
- Global feedback: population size dependence
Let $m_z =$ expected offspring number | population of size z . Assume $\sum (m_z - m)/z < \infty$.



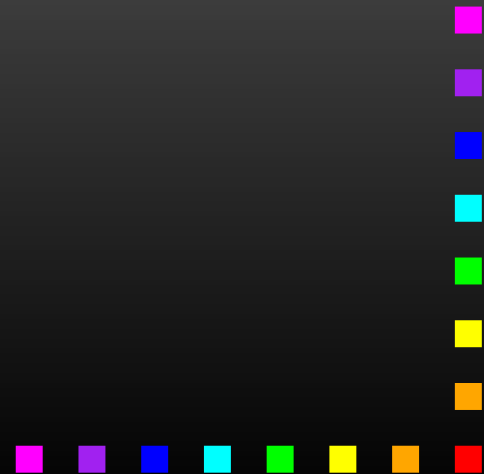
Branching with Dependence

- Local interaction – between relatives
- Stationary or i.i.d random environments – independent branching | environment
- Global feedback: population size dependence
Let $m_z =$ expected offspring number | population of size z . Assume $\sum (m_z - m)/z < \infty$.
- In all these cases the dichotomy holds.



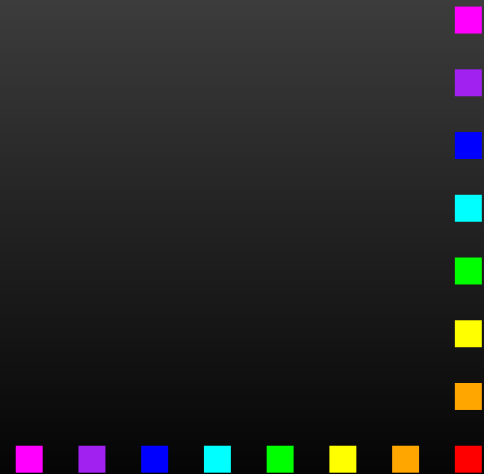
Population Size Dependence

- Galton-Watson (Klebaner 80's),



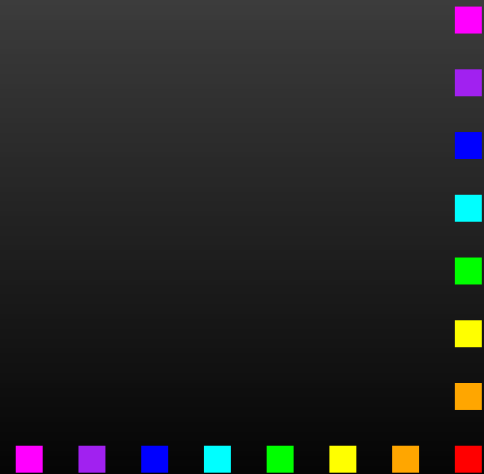
Population Size Dependence

- Galton-Watson (Klebaner 80's),
- multi-type GW (Klebaner 90's),



Population Size Dependence

- Galton-Watson (Klebaner 80's),
- multi-type GW (Klebaner 90's),
- discrete time general process (PJ & Sagitov, 2000),



Population Size Dependence

- Galton-Watson (Klebaner 80's),
- multi-type GW (Klebaner 90's),
- discrete time general process (PJ & Sagitov, 2000),
- continuous time age- and population-size-dependent reproduction (PJ & Klebaner 2000).



Population Size Dependence

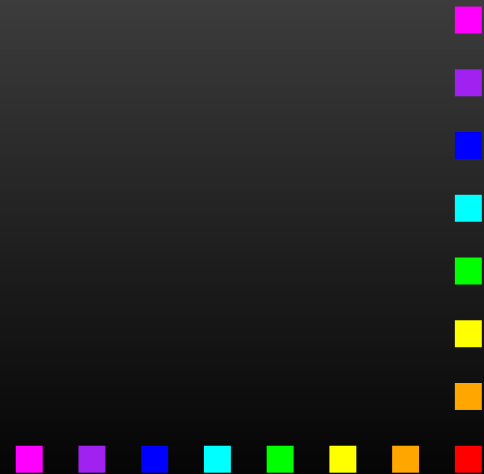
- Galton-Watson (Klebaner 80's),
- multi-type GW (Klebaner 90's),
- discrete time general process (PJ & Sagitov, 2000),
- continuous time age- and population-size-dependent reproduction (PJ & Klebaner 2000).
- The general case remains.



Simple Processes in General Environments

\mathcal{E}_n is the environment up to and including season n , Z_n is the population size then.

- Population size is part of the environment,
 $Z_n \in \mathcal{E}_n$



Simple Processes in General Environments

\mathcal{E}_n is the environment up to and including season n , Z_n is the population size then.

- Population size is part of the environment,
 $Z_n \in \mathcal{E}_n$
- Reproduction probabilities are determined by \mathcal{E}_n .



Simple Processes in General Environments

\mathcal{E}_n is the environment up to and including season n , Z_n is the population size then.

- Population size is part of the environment,
 $Z_n \in \mathcal{E}_n$
- Reproduction probabilities are determined by \mathcal{E}_n .
- Given \mathcal{E}_n , individuals reproduce independently (and according to the same distribution – single type).



Near Critical Reproduction

Write μ_n for the expected offspring per individual, given \mathcal{E}_n , Assume $\mu_n = 1 + C_n/Z_n + R_n$, if $Z_n > 0$. C_n independent of Z_n , $R_n = o(1/Z_n)$.

- Then, with $R_n = 0$,

$$\mathbf{E}[Z_{n+1}] = \mathbf{E}[Z_n \mu_n] = \mathbf{E}[Z_n] + \mathbf{E}[C_n; Z_n > 0] .$$



Near Critical Reproduction

Write μ_n for the expected offspring per individual, given \mathcal{E}_n , Assume $\mu_n = 1 + C_n/Z_n + R_n$, if $Z_n > 0$. C_n independent of Z_n , $R_n = o(1/Z_n)$.

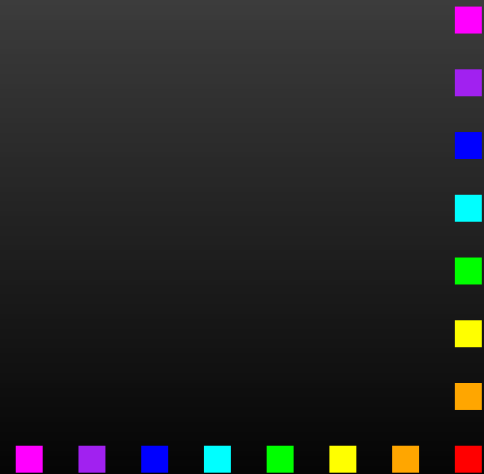
- Then, with $R_n = 0$,
$$\mathbf{E}[Z_{n+1}] = \mathbf{E}[Z_n \mu_n] = \mathbf{E}[Z_n] + \mathbf{E}[C_n; Z_n > 0] .$$
- With $c_n = \mathbf{E}[C_n]$, independence yields
$$\mathbf{E}[Z_N] = \mathbf{E}[Z_0] + \sum_{n=1}^N c_n \mathbf{P}(Z_n > 0).$$



Asymptotics

- With $\mathbf{P}(Z_n > 0) \rightarrow \mathbf{P}(Z_n \rightarrow \infty) = 1 - q > 0$,

$$\mathbf{E}[Z_N] - \mathbf{E}[Z_0] \sim (1 - q) \sum_{n=1}^N c_n. \quad (1)$$



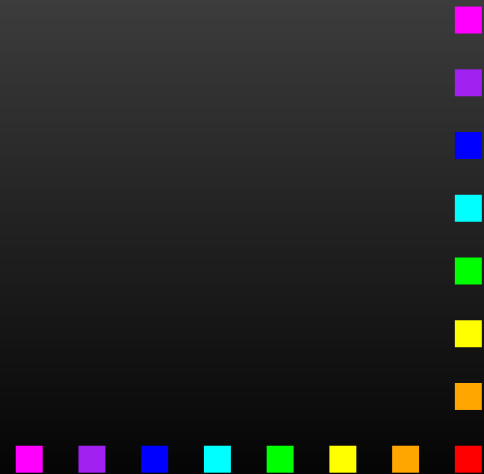
Asymptotics

- With $\mathbf{P}(Z_n > 0) \rightarrow \mathbf{P}(Z_n \rightarrow \infty) = 1 - q > 0$,

$$\mathbf{E}[Z_N] - \mathbf{E}[Z_0] \sim (1 - q) \sum_{n=1}^N c_n. \quad (1)$$

- What about the process itself,

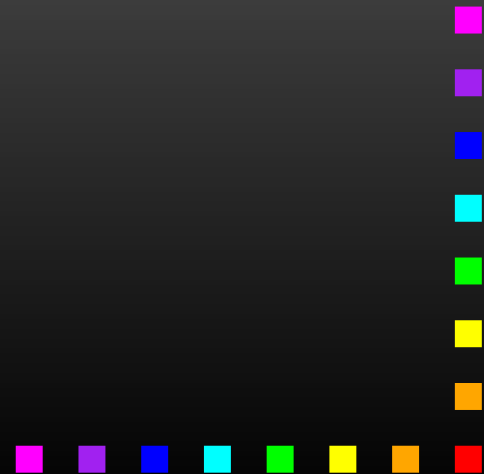
$$Z_N / \sum_1^N c_n \sim ?$$



Interesting cases:

- Stationary exterior conditions (like the C_n i.i.d. with $c_n = c$)

$$Z_N \sim cN.$$



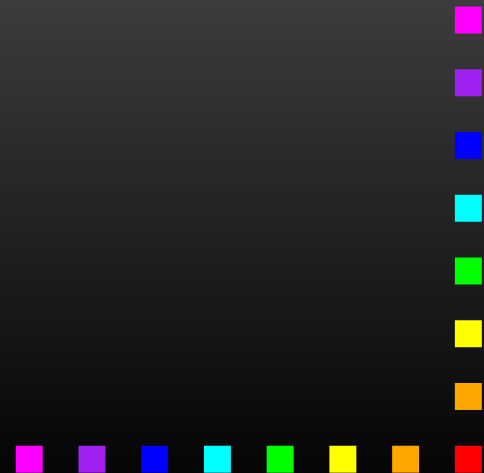
Interesting cases:

- Stationary exterior conditions (like the C_n i.i.d. with $c_n = c$)

$$Z_N \sim cN.$$

- Deteriorating exterior conditions (like C_n independent with $c_n = c/n$)

$$Z_N \sim c \log N.$$



Interesting cases:

- Stationary exterior conditions (like the C_n i.i.d. with $c_n = c$)

$$Z_N \sim cN.$$

- Deteriorating exterior conditions (like C_n independent with $c_n = c/n$)

$$Z_N \sim c \log N.$$

- Improving outer environment (like $c_n \sim cn$)

$$Z_N \sim cN^2/2.$$



Process Convergence – Assumptions:

(1) The conditional offspring distributions at times $n = 0, 1, 2, \dots$, possess moments of all orders.

For each order, these are bounded above;

(2) $\sum_{j=1}^n c_j \rightarrow \infty$ and $c_n / \sum_{j=1}^n c_j \rightarrow 0$;

(3) Reproduction variances σ_n^2 stabilise as the population grows: $\sigma_n^2 \rightarrow V_n$, $\mathbf{E}[V_n] = v_n$, V_n (like C_n) independent of Z_n .

(4) Reproduction variances do not swamp means:

$$v_n \sim ac_n, a \geq 0.$$



Process Convergence —

Conclusion:

Then, either the population goes extinct or

$$\frac{Z_n}{\sum_{j=1}^n c_j} \xrightarrow{d} \Gamma\left(\frac{2}{a}, \frac{2}{a}\right),$$

where $\Gamma(\infty, \infty) = 1$.

Proof by the method of moments and induction:

$$\mathbf{E}[Z_{n+1}^k | \mathcal{E}_n] = Z_n^k + \left(kC_n + \binom{k}{2} V_n \right) Z_n^{k-1} + o(Z_n^{k-1}).$$



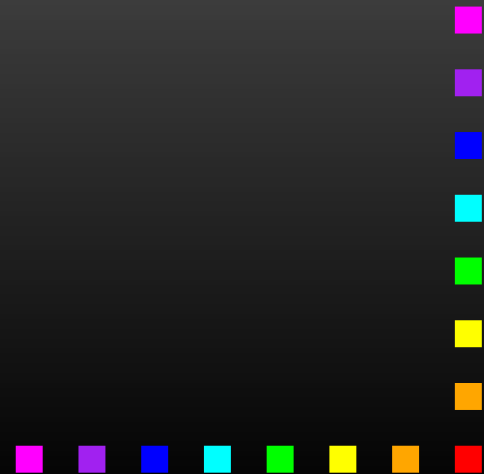
PCR

The *Polymerase Chain Reaction*, generates a test sample out of a minuscule amount of DNA. In *quantitative* PCR the successive growth of the amount of DNA is followed from a threshold of observation up to large molecule numbers. In ordinary PCR focus is on the end product, which is sufficiently big to detect viruses or mutations, and for paternity testing or forensic matters.



PCR Steps

- The first PCR step is *denaturation*: heating up to 90°C so that DNA strands separate.



PCR Steps

- The first PCR step is *denaturation*: heating up to 90°C so that DNA strands separate.
- In the subsequent *annealing* phase, at around 50°C, short synthetic so called *primers* bind to the target sequences.

PCR Steps

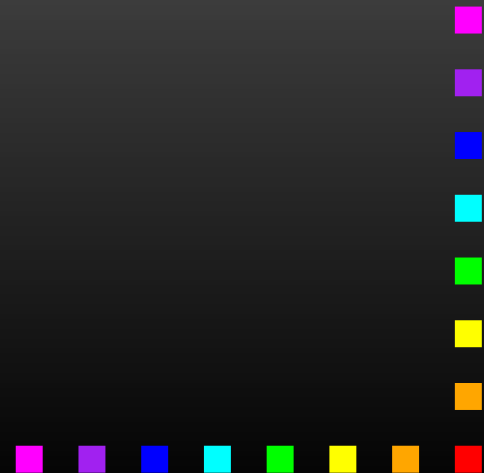
- The first PCR step is *denaturation*: heating up to 90°C so that DNA strands separate.
- In the subsequent *annealing* phase, at around 50°C, short synthetic so called *primers* bind to the target sequences.
- Temperature is raised and an enzyme, the *polymerase* promotes the synthesis process at the region marked by the primers, the primer extending into a complementary DNA string.



Quantitative PCR

In Q-PCR three phases are conventionally discerned:

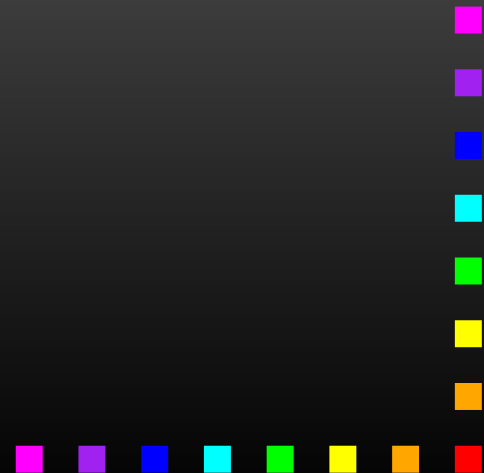
- After an unclear beginning, *exponential* growth,



Quantitative PCR

In Q-PCR three phases are conventionally discerned:

- After an unclear beginning, *exponential* growth,
- followed by *linear* increase,



Quantitative PCR

In Q-PCR three phases are conventionally discerned:

- After an unclear beginning, *exponential* growth,
- followed by *linear* increase,
- gradually changing over into *saturation*.



Model

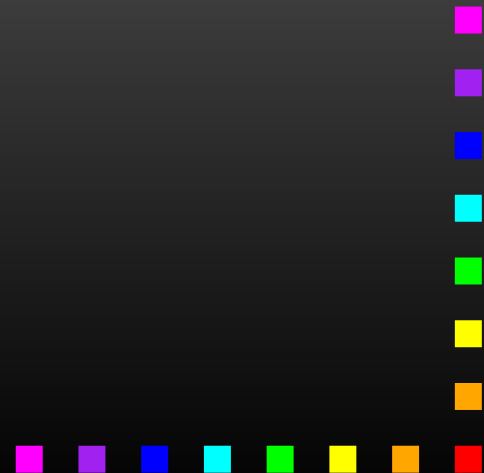
$$\# \text{ "offspring" } = \begin{cases} 2 & \text{with probability } p \\ 1 & \text{with probability } 1 - p \end{cases}$$

In most literature, $p = 1$ – or constant. But $p = p(\text{target DNA, primers, Taq polymerase, deoxynucleic triphosphate, MgCl}_2 \dots)$.



Michaelis-Menten kinetics

- The critical ingredient is the amount of target DNA, z .



Michaelis-Menten kinetics

- The critical ingredient is the amount of target DNA, z .
- By classical enzyme kinetics, the rate of the reaction is $z/(z + K)$, where K (very large) summarises all other influences.

Michaelis-Menten kinetics

- The critical ingredient is the amount of target DNA, z .
- By classical enzyme kinetics, the rate of the reaction is $z/(z + K)$, where K (very large) summarises all other influences.
- Thus, in one cycle starting from z , $zp(z) \propto z/(z + K)$ new molecules are synthesized.

PCR is near-critical

Since $p(1) \approx 1$, $p(z) = \frac{K}{K+z}$. Let K_n be the Michaelis Menten constant in the n :th cycle. Then,

$$\mu_n = 1 + \frac{K_n}{K_n + Z_n} = 1 + K_n/Z_n + o(1/Z_n)$$

and

$$\sigma_n^2 = \frac{K_n Z_n}{(K_n + Z_n)^2}.$$



and grows linearly:

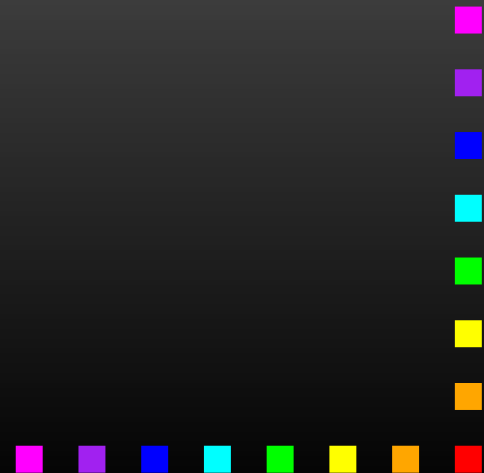
If the M-M constants have the same mean,

$$c_n = \mathbf{E}[K_n] = c, \text{ then } \sum_1^n c_j = cn,$$

and $v_n = o(1)$, so that

$$Z_n/n \rightarrow c$$

in probability.



The Martingale

Doob: $Z_{j+1} = \mu_j Z_j + Z_{j+1} - \mathbf{E}[Z_{j+1} | \mathcal{E}_j] =$
 $Z_j + \frac{K_j Z_j}{K_j + Z_j} + Z_{j+1} - \mathbf{E}[Z_{j+1} | \mathcal{E}_j]$. Repeat to obtain
 $Z_n = Z_0 + A_n + M_n$, where

$$A_n = \sum_{j=0}^{n-1} \frac{K_j Z_j}{K_j + Z_j}$$

is increasing and in \mathcal{E}_{n-1} . Since $Z_j \rightarrow \infty$,
 $A_n/n \rightarrow c$ a. s., by LLN, eg. if the K_j are i.i.d.

Almost sure convergence

$M_n = \sum_{j=1}^n Z_j - \mathbf{E}[Z_j | \mathcal{E}_{j-1}]$ – a martingale with respect to $\{\mathcal{E}_n\}$.

$$\begin{aligned} \mathbf{E}[(M_n - M_{n-1})^2 | \mathcal{E}_{n-1}] &= \mathbf{Var}[Z_n | \mathcal{E}_{n-1}] = \\ &= \frac{K_{n-1} Z_{n-1}^2}{(K_{n-1} + Z_{n-1})^2} \leq K_{n-1}. \end{aligned}$$

Hence, $\sum_{n=1}^{\infty} \mathbf{E}[(M_n - M_{n-1})^2] / n^2 < \infty$.

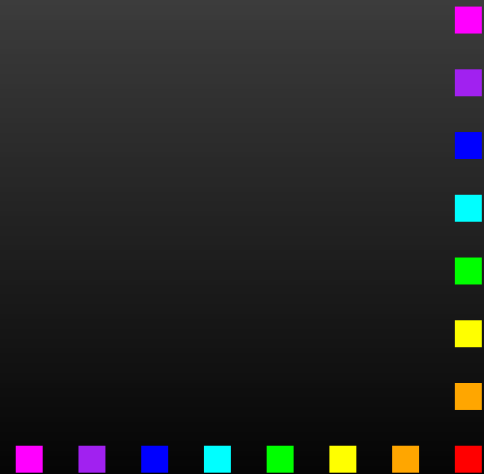


Almost sure convergence

Since $\sum_{n=1}^{\infty} \mathbf{E}[(M_n - M_{n-1})^2]/n^2 < \infty$, the Martingale Law of Large Numbers yields $M_n/n \rightarrow 0$ a. s. Hence,

$$Z_n/n = Z_0/n + A_n/n + M_n/n \rightarrow 0 + c + 0,$$

with probability one.



Linear growth

This recovers the linear phase of quantitative PCR. Saturation requires a more subtle discussion of K_n when Z_n or $n \rightarrow \infty$.

Joint work with F. C. Klebaner. Partly published in J. Appl. Prob 41 (A), 2004.