

The dynamics of fitness under selection, and context-dependent mutation

Guillaume Martin & Lionel Roques

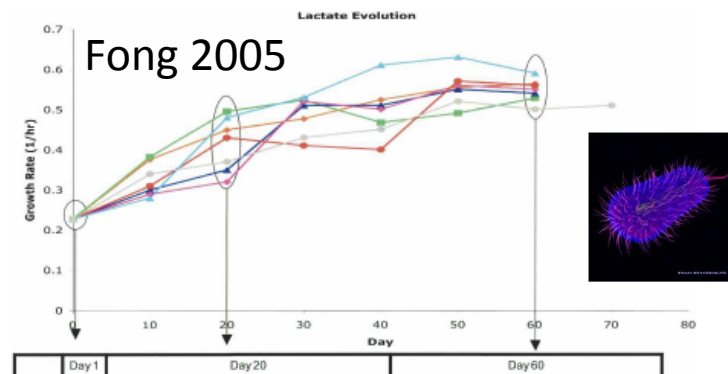
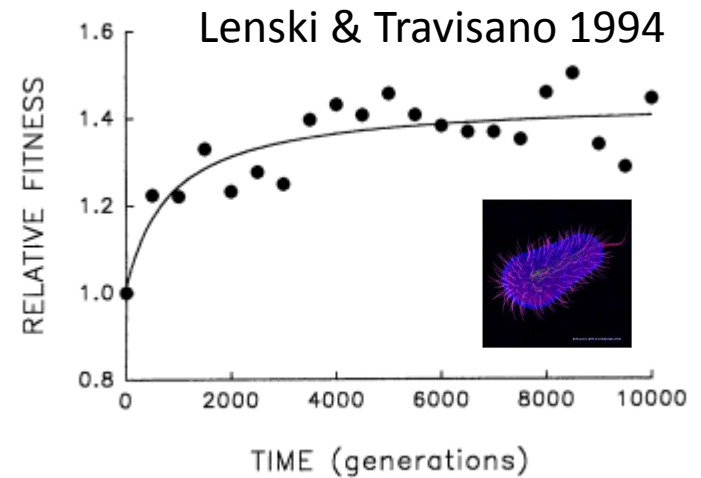
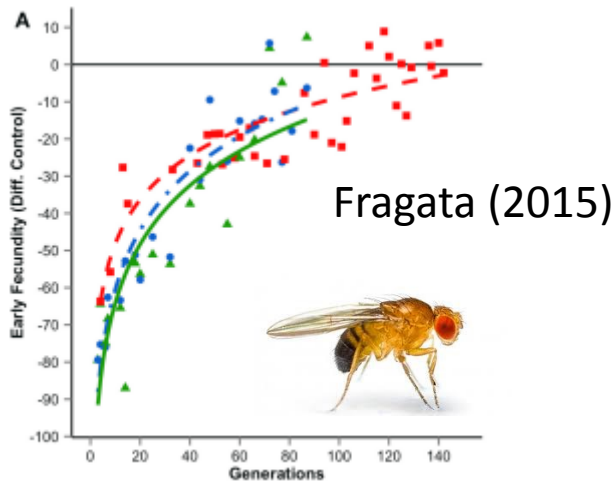
Models describing the rate of adaptation

Typical adaptation models: **stable regime assumptions** $\Rightarrow \partial_t E(\bar{m}) = \text{constant}$

fitness: substitutions (*Gerrish & Lenski 1998*), Travelling waves (*Desai & Fisher 2007*)

traits: Quantitative genetics (*Lande 1979*) etc.

Typical adaptation data: **saturation**



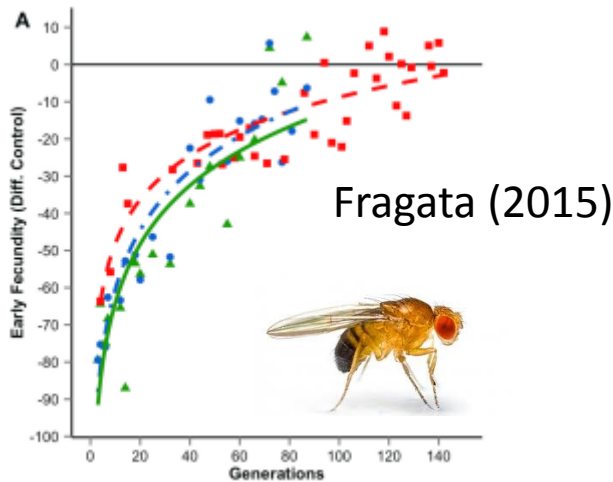
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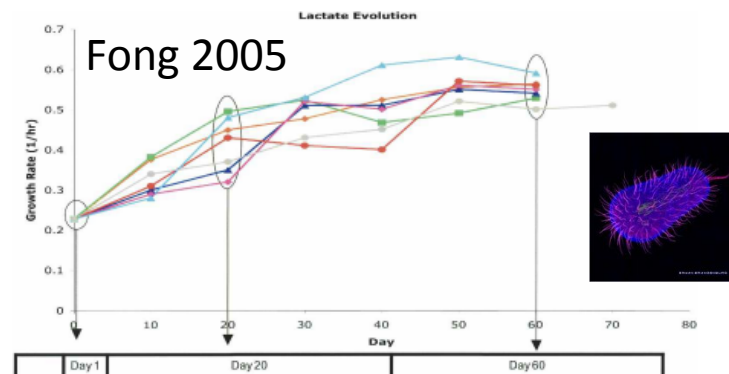
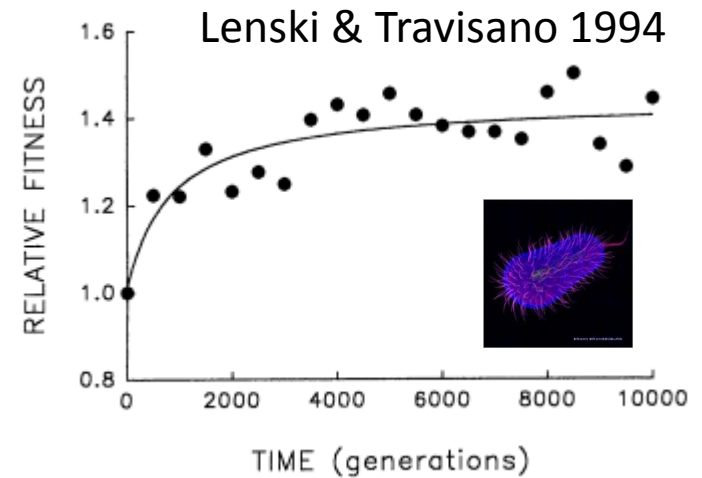
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traits: Quantitative genetics (*Lande 1979*) etc.

Typical adaptation data: **saturation**



- Erosion of initial variance
- Epistasis



Fitness distribution dynamics with epistasis + standing variance

generating function of the fitness distribution

Generating functions can prove handy:

R. Bürger (1991) : trait mutation

Johnson (1999), Desai & Fisher (2011) : deleterious mutation

Gerrish & Sniegowski (2012): mutation + short term

Ratray & Shapiro (2001): biallelic locus + mutation + drift

Good & Desai (2013): fitness-based , deleterious mutation + drift

Often: Infinite set of moment equations (but see Johnson 1999)

No epistasis

Fitness trajectories with epistasis:

Kryazhimskiy (2009) Dwyer (2012), Good & Desai (2015)

Analytic progress = away from clonal interference regime

Definitions / Assumptions

Asexuals, multitype wright fisher diffusion (continuous time approx)

m_i : malthusian fitness of genotype i (no frequency/density dependence)

Mutation : poisson process at rate U per unit time per capita

Distribution of Fitness Effects (DFE): $f(s|m_i)$ in background m_i

Cumulant Generating Function (CGF) :

$$C_t(z) = \log(\sum p_i(t) e^{z m_i}) \quad \text{at time } t \quad (z \in \mathbb{R}^+ \text{ if } m < 0)$$

Derivatives in $z \Rightarrow$ cumulants moments: $C'_t(0) = \bar{m}_t$: mean fitness

Dynamics of C_t ?

Selection + drift

$\mathcal{C}_t(z) = \langle C_t(z) \rangle$: expected CGF over stochastic process:

$\mathbf{p}(t) = \{p_i(t)\}_{i \in [1, K]} \in [0, 1]^K \sim K$ –type Wright-Fisher diffusion

e.g. apply Feynman-Kac Theorem:

$$\partial_t \mathcal{C}_t(z) = \boxed{C'_t(z) - C'_t(0)} - \boxed{\frac{1 - \langle e^{C_t(2z) - 2C_t(z)} \rangle}{2N_e}}$$

Selection (clonal interference)

drift

When can we neglect drift here ?

$$\partial_t C_t(z) = \boxed{C'_t(z) - C'_t(0)} - \boxed{\frac{1 - \langle e^{C_t(2z) - 2C_t(z)} \rangle}{2N_e}}$$

Selection
Drift effect δ

- $\text{Var}(\text{within}) \gg \text{Var}(\text{between})$

p_{max} : frequency of the fittest class

$$0 \leq |\delta| \leq \frac{1}{2N_e} |1 - \langle p_{max}^{-1} \rangle|$$

OK in models where the fittest class remains or quickly becomes substantial ($p_{max} \gg 1/N$)
 = fitness upper bound (optimum, purely deleterious before ratchet)

Not OK on unbounded fitness sets (travelling waves) or with Muller's ratchet (late effect)

Background-dependent Mutation

Mutation effects CGF: $C_S(z, m) = \log(\int e^{s z} f(s|m) ds)$

Assume: linear context-dependence: $C_S(z, m) \approx \omega(z)m + C_*(z)$

$C_*(z)$: CGF of DFE in background $m = 0$ (e.g. optimum)

$\omega(z) = \partial_m C_S(z, m)|_{m=0}$: « context-dependence function »

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Small m approx when nearing a maximum fitness set at $m = 0$:

=> mutation – selection balance with diminishing returns epistasis

Exact at all times in some particular models:

- $\omega(z) = 0$: any non-epistatic model (finite moments)
- $\omega(z) = -z$: House Of Cards model (absolute effect independent of m)
- Fisher's geometrical model (quadratic trait- fitness function), see next

Closed approximate dynamics

$$\partial_t \mathcal{C}_t(z) \approx \boxed{c'_t(z) - c'_t(0)} + \boxed{U(\exp(c_*(z) + c_t(z + \omega(z))) - c_t(z)) - 1}$$

selection

Background-dependent mutation

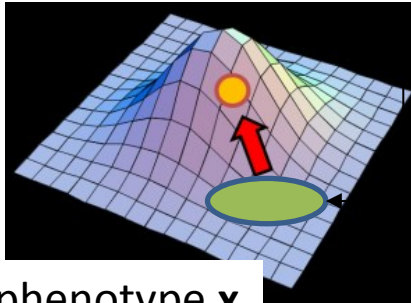
$\mathcal{C}_0(z)$: initial condition (standing fitness variance)

Nonlinear nonlocal PDE

can be solved (at least numerically) for given $c_*(z)$, $\omega(z)$, $\mathcal{C}_0(z)$

Example : Fisher's model

fitness $e^{m(\mathbf{x})}$



phenotype \mathbf{x}

Mutant cloud around background \mathbf{x}

- variance λ per trait (isotropy)
- dimension n
- Normally distributed $\mathbf{dx} \sim N(\mathbf{0}, \lambda \mathbf{I}_n)$ (relaxed in some cases)

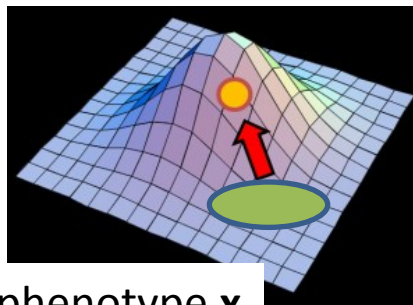
$$\Rightarrow C_s(z, m) = \omega(z) m + C_*(z)$$

Martin (2014)

$C_*(z) = -n/2 \log(1 + \lambda z)$: neg. gamma DFE at optimum

$\omega(z) = -\lambda z^2 / (1 + \lambda z)$: epistasis

fitness $e^{m(x)}$



phenotype x

Example : Fisher's model

Example: at $t = 0$: a clone with fitness m_0
 \Rightarrow **expected mean fitness trajectory**: $\mathcal{C}'_t(0) = \langle \bar{m}_t \rangle$

- Strong Selection Weak Mutation SSWM ($U \ll s$)

$$\begin{cases} \partial_t \langle \bar{m}_t \rangle = U e^{\langle \bar{m}_t \rangle \omega(t)} (\langle \bar{m}_t \rangle M_*(t) \omega'(t) + M'_*(t)) \\ \langle \bar{m}_0 \rangle = m_0 \end{cases} \quad : \text{ODE easy to solve, fit etc.}$$

- Weak selection strong mutation WSSM ($U \gg s$)

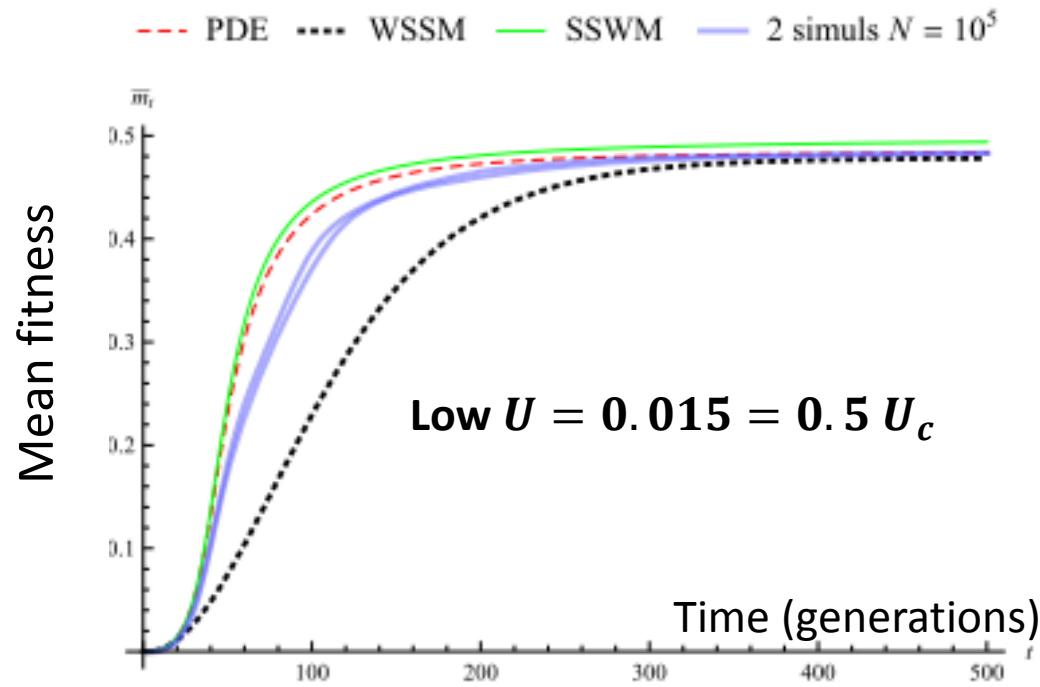
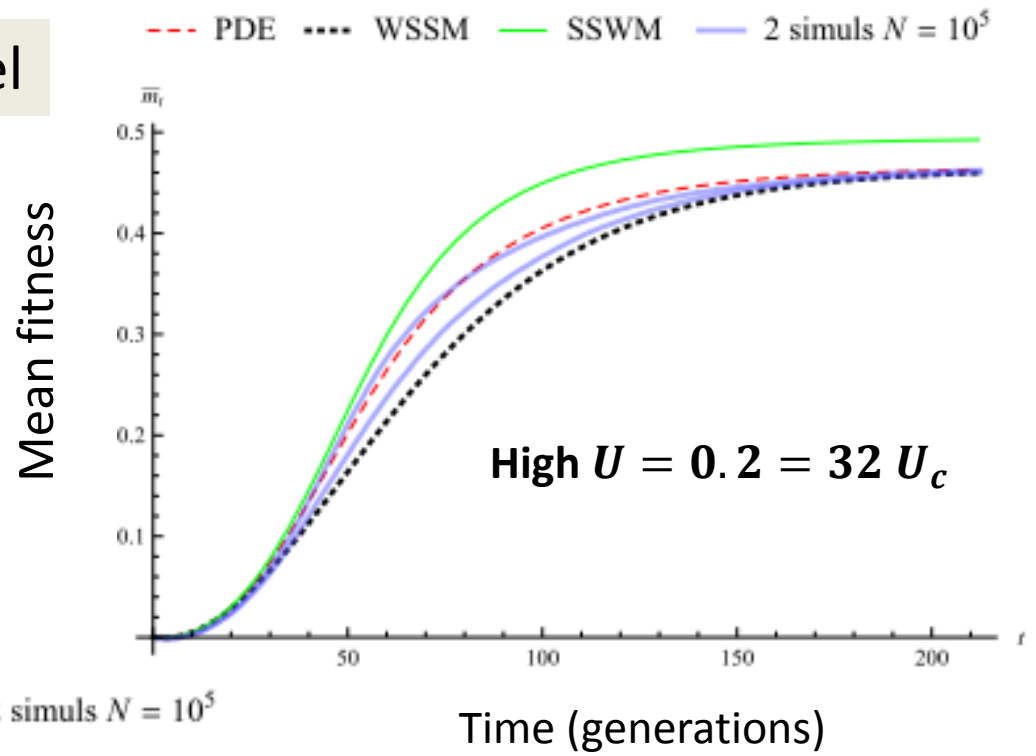
Leading order in $\lambda \Rightarrow$ analytic solution to linearized PDE $\Rightarrow \langle \bar{m}_t \rangle = \mathcal{C}'_t(0)$

$$\begin{aligned} \langle \bar{m}_t \rangle - m_0 &= -m_0 \tanh(\mu t)^2 - n/2 \mu \tanh(\mu t) \\ \mu &= \sqrt{U \lambda} \end{aligned}$$

Valid when $U \gg U_c = n^2 \lambda / 4$

Fisher's Geometrical Model

$$n = 5, \lambda = 0.001, N = 10^5$$



Strategies for empirical testing

Obvious quantitative test: high power to reject/compare models

$$\langle \bar{m}_t \rangle = f(U, n, \lambda, m_0, t) \quad \left\{ \begin{array}{l} \text{PDE + analytical approximations} \\ \text{Fisher's model and House of cards} \end{array} \right.$$

1. Fit trajectories of \bar{m}_t
2. Compare fitted parameters to direct estimates (U, n, λ)

challenge: precision/stability of estimates across environments/labs etc.

Heuristic test: Use WSSM approx trajectory

$$\langle \bar{m}_t \rangle - m_0 = -m_0 \tanh(\mu t)^2 - n/2 \mu \tanh(\mu t)$$

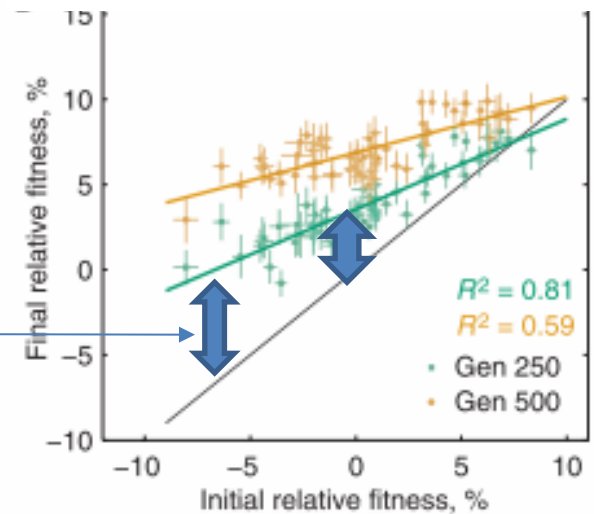
Expected cumulated improvement \propto initial fitness

- 1/ check linearity
- 2/ predict regression coefficients (depend on time + mutational parameters)

NB: Linearity with m_0 seems to hold outside the WSSM approx (formal proof ?)

Yeast in YPD 30°C:
 Founders with various m_0
 Mean fitness after $t = 250$ or 500 generation

m_{ref} : fitness of reference strain for competitions



1/ Good linearity ($R^2 \geq 60\%$):

2/ Observed regression coefs at $t = 250$

$\hat{\mu} = 0.0034, \hat{m}_{ref} = 0.09$

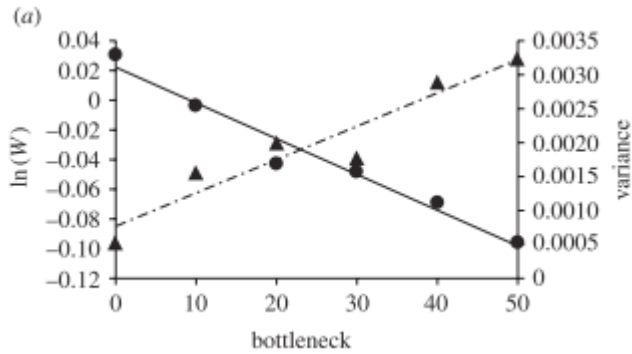
~ predicts regression coefs at $t = 500$

coef.	predicted	observed
β_{500}	-0.87	-0.67
α_{500}	7.86	7.88

NB: rejects context indep. model (slope = 0) and unique optimum reached (slope = -1)

Mutation Accumulation data

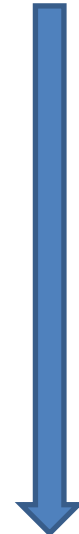
E. coli in LB, fitness in competition



Trindade et al. 2010

Perfeito et al. *Evolution* (2013)

→ $\mu_{pred} \sim 0.007$

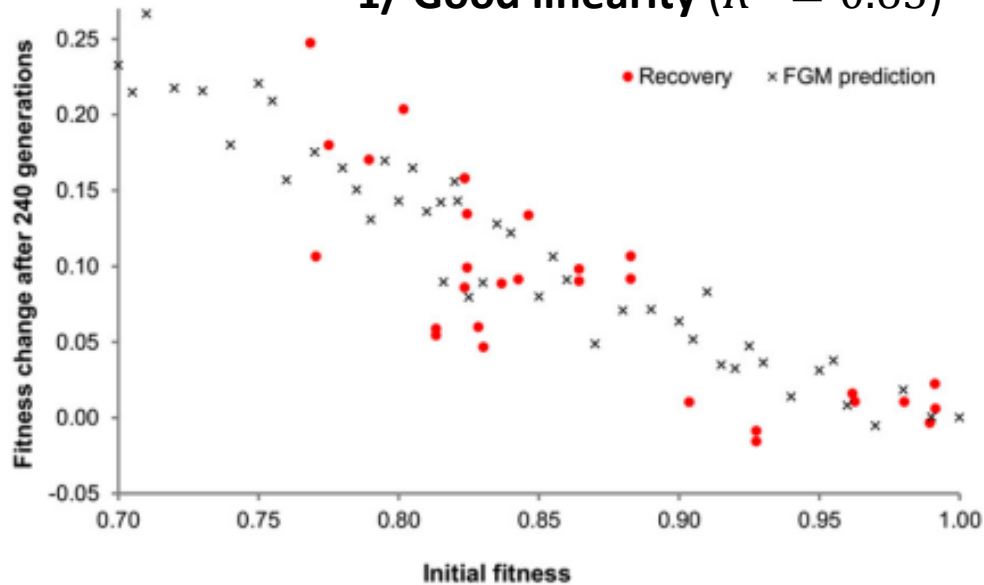


Perfeito et al. *Evolution* (2013): already tested FGM by ABC

Dynamics of fitness recovery in LB after 240 generations, same conditions

Against ancestor ($m_{ref} = 0$)

1/ Good linearity ($R^2 = 0.63$)



2/ ~ Predicts regression coefficients

coef.	predicted	observed
β_{240}	- 0.87	-0.77
α_{240}	- 0.019	- 0.02

Conclusions

Null models for experimental evolution

+

Classic methods to parameterize them from independent data

} quantitatively test
population genetics

Drift need not be modelled sometimes, even in asexuals and with clonal interf.

=> Prediction robust to drift parameters (birth death rates, in each geno etc.)

- Heuristic test uses WSSM which is not optimal here a priori (just orders)

- Accounting for drift sensitive regimes (bias + envelope around expectations)

some predictable patterns overlooked in experimental evolution (e.g. standing variance)

Extensions: e.g. coupling with rescue dynamics, Yoann Anciaux, poster 1



sequences data / coalescent: fitness distribution with predictable

shape/scale/variance over time. Implement into MMC Coalescents ?

Thanks

The organizers and you

Ongoing tests / extensions :

Thomas Lenormand (CEFE Montpellier)

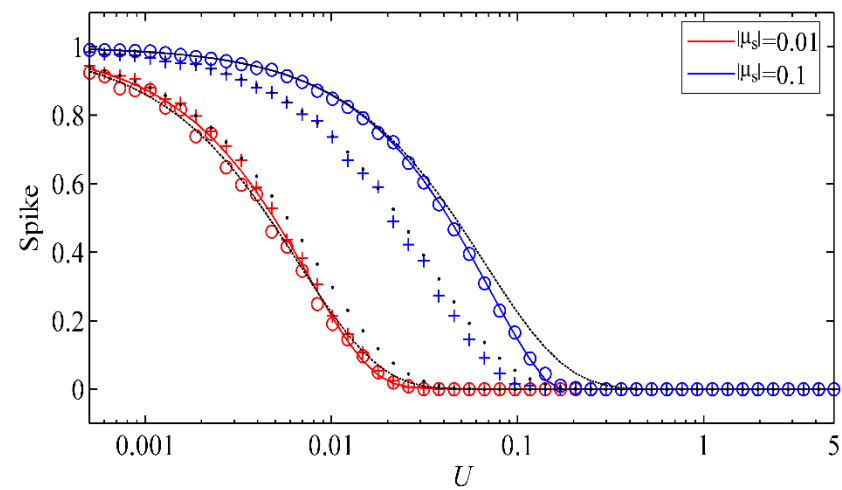
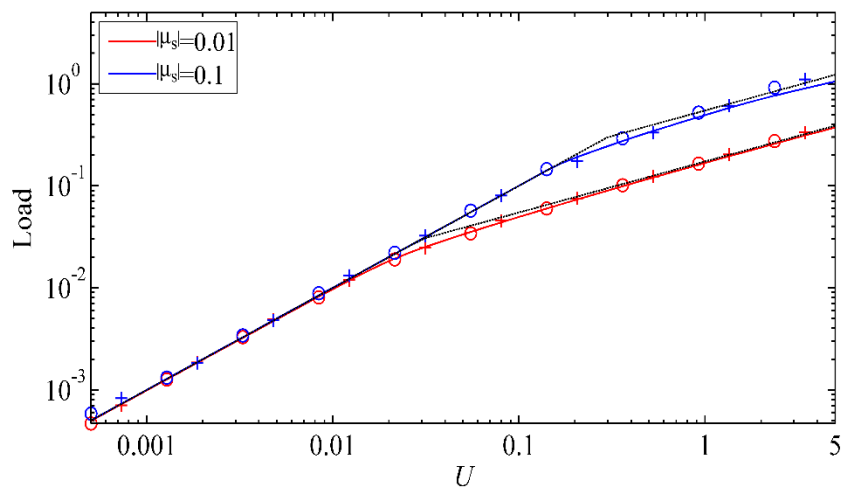
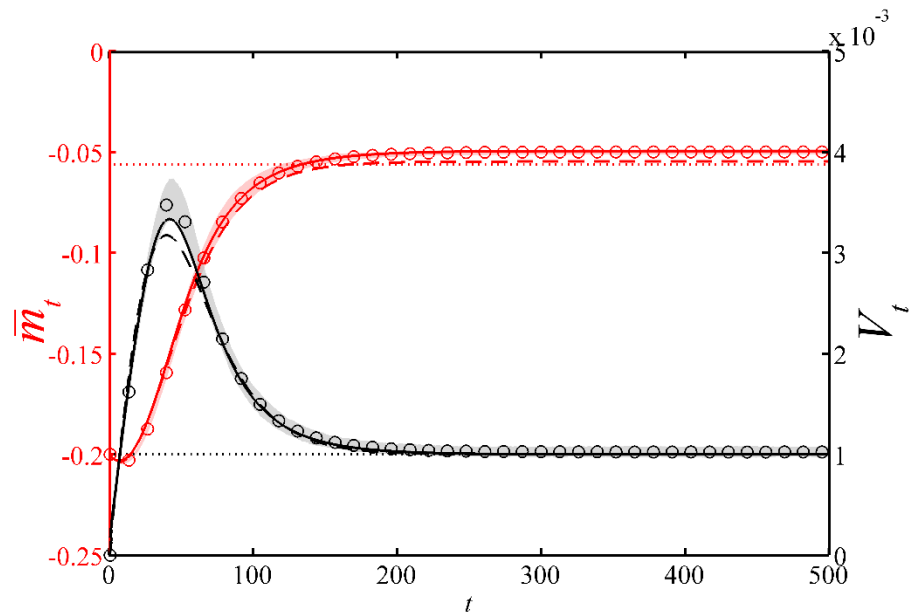
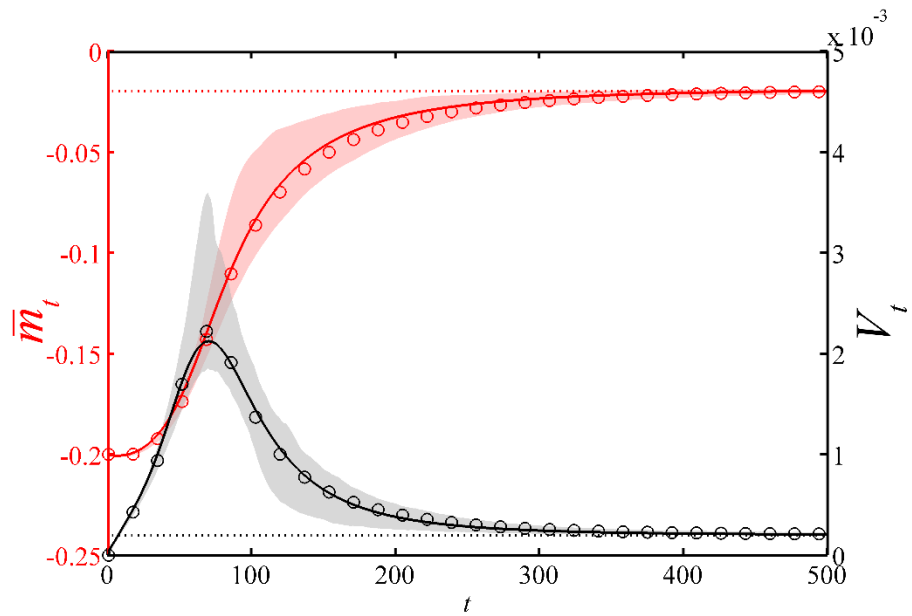
Yoann Anciaux (ISEM Montpellier)

Ophelie Ronce (ISEM Montpellier)

Amaury Lambert (UPMC, Collège de France)

Noemie Harmand (CEFE, Montpellier)

Luis Miguel Chevin (CEFE, Montpellier)

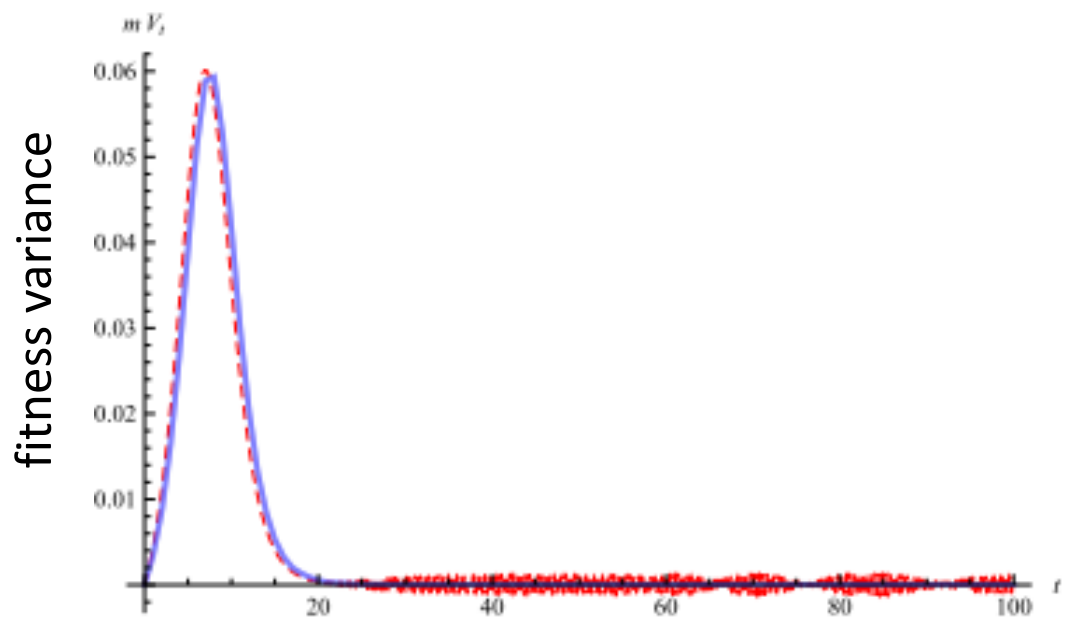
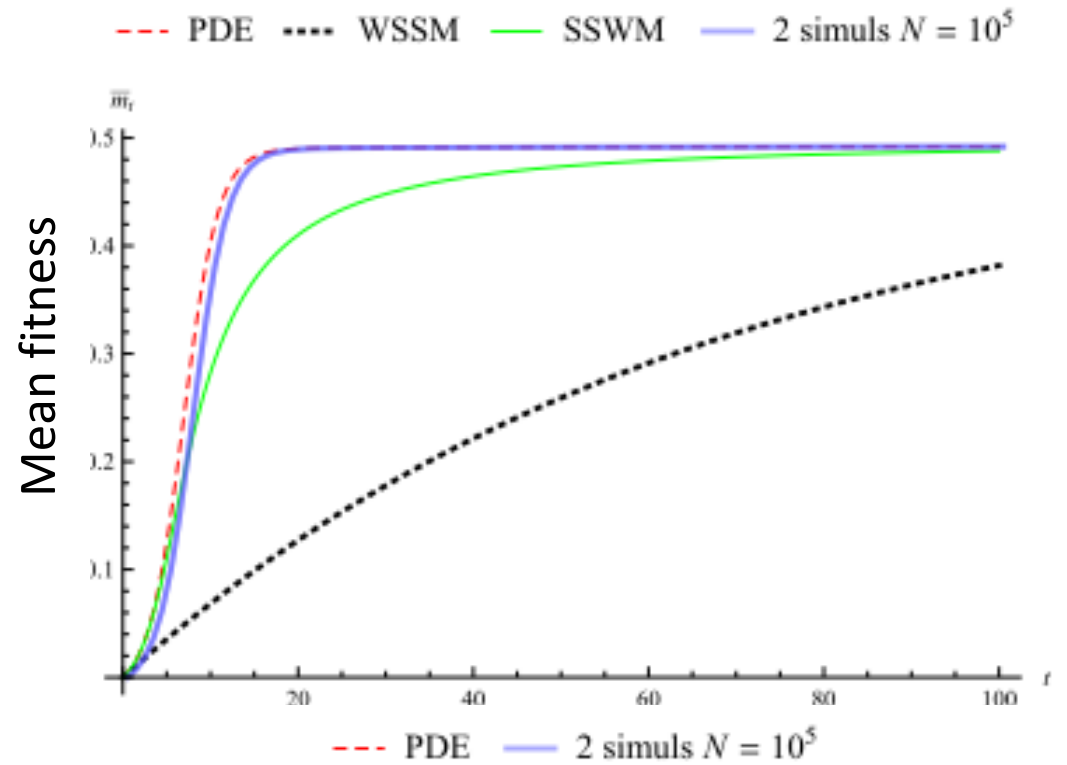


House of Cards Model

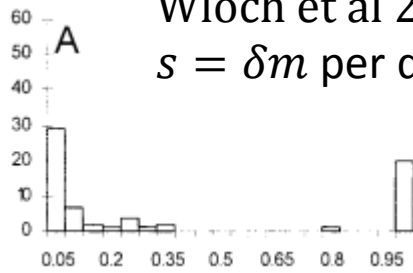
$C_*(z)$: arbitrary

(here gamma DFE)

$$\omega(z) = -z$$



Wloch et al 2001, random single mutations in yeast
 $s = \delta m$ per division in (haploid effect in YPD)

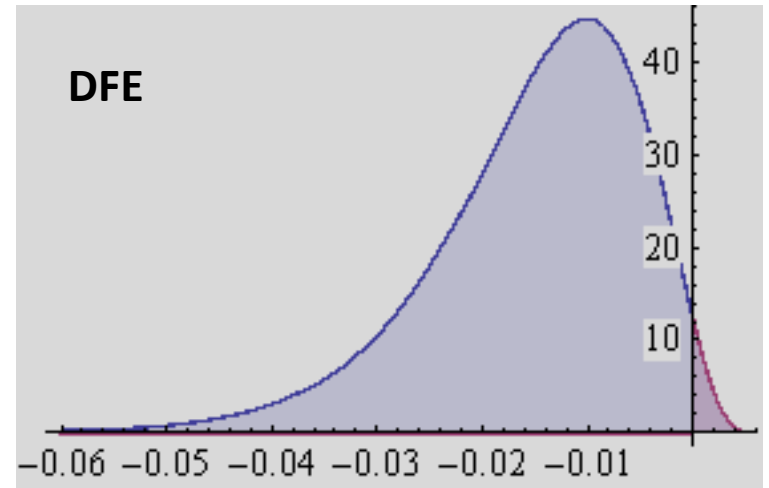
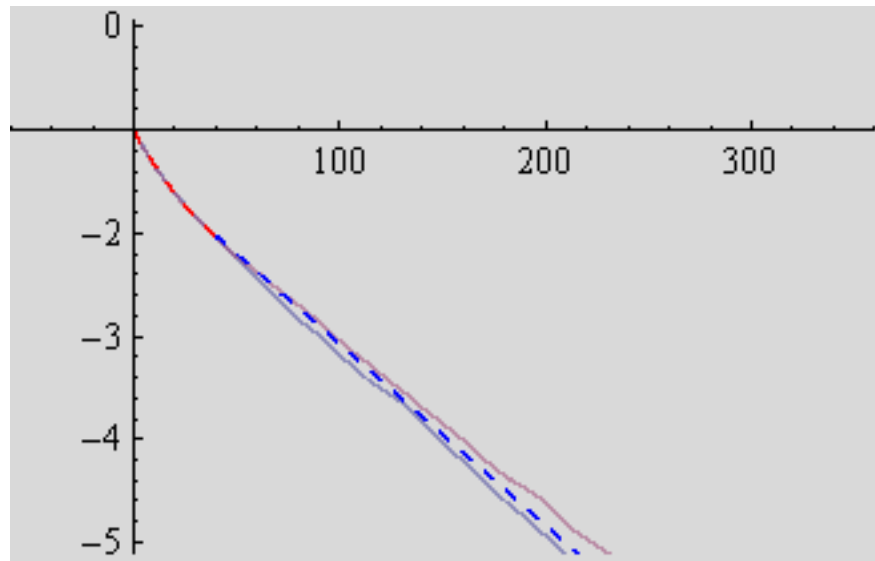
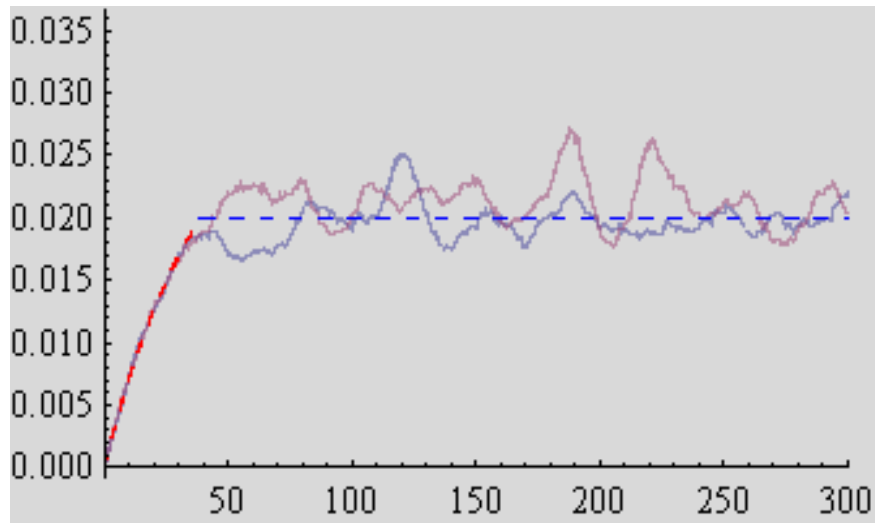


$$U = 0.66 \times 10^{-3} (\text{div}^{-1})$$

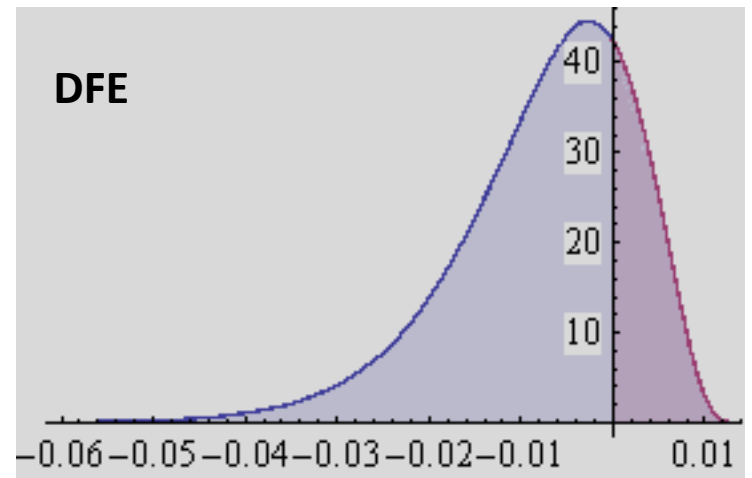
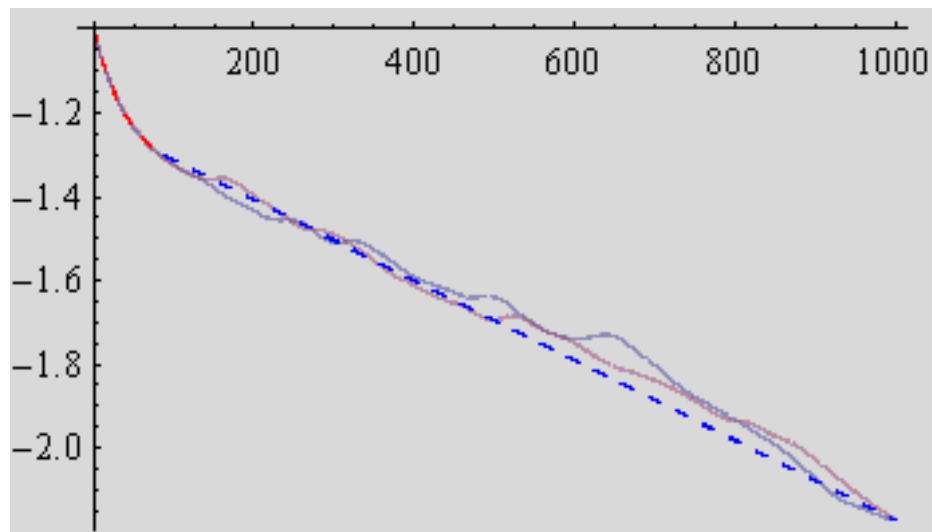
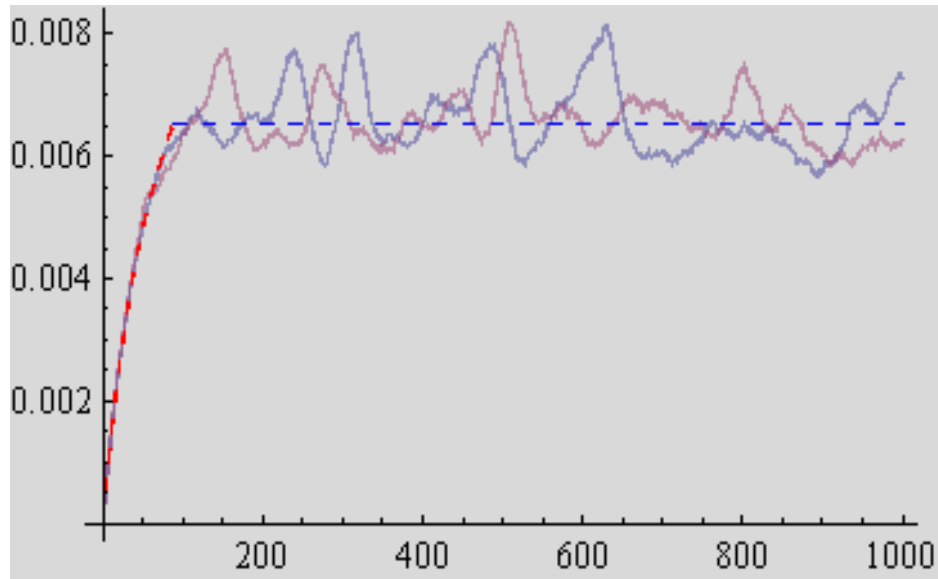
$$\bar{s} = 0.086 (\text{div}^{-1})$$

$$n = 1/CV^2$$

$$N = 10^4, U = 500\lambda, \theta = 4, s_0 = \lambda, \lambda = 0.005$$



$$N = 10^5, U = 200\lambda, \theta = 4, s_0 = 2.5\lambda, \lambda = 0.005$$



$$N = 10^5, U = 200\lambda, \theta = 4, s_0 = 3\lambda, \lambda = 0.01$$

